

Developing Costimulatory Molecules for Immunotherapy of Diseases

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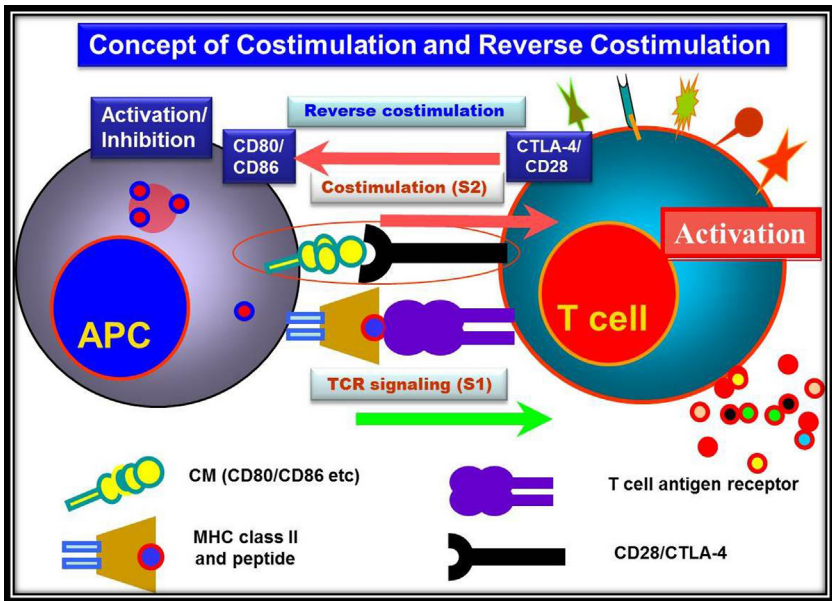
**Dedicated to
My
Beloved Family
And
Teachers**

About the Editor

Dr. Manzoor completed his masters in life sciences with a gold medal and after qualifying with the prestigious JRF-NET-CSIR examination completed his PhD in immunopathology from IMTECH-JNU. His PhD work includes understanding the role of reverse costimulation in the survival of intracellular pathogens and cancer. His area of research includes costimulation biology and immunology of stroke. He is Assistant Professor at the Department of Bioresources, University of Kashmir, Srinagar, India, and presently working in the Stroke Research



Chair on novel recombinational therapies for stroke as Research Scientist at the College of Applied Medical Sciences, Majmaah University, Kingdom of Saudi Arabia. He has several international research papers, review articles, and books to his credit. He is a member of many scientific societies across the globe. He has many courses to his credit besides a doctorate degree in immunology which includes PGDHE and PGDEE. Dr. Manzoor has already authored many books on costimulation immunotherapy and stroke biology.



Preface

Considering the importance of complex roles played by CD28 and B7 costimulatory families in regulating the immune system, we predict that novel approaches targeting these families will yield new therapies for the treatment of inflammation, autoimmunity, transplantation, cancer, allergies, asthma, and other diseases. This book highlights the novel concept of reverse costimulation, which can be effectively exploited to develop immunotherapy either using humanized antibodies against CD80, CD86, and other costimulatory molecules or CD28 fusinogenic proteins for the treatment of cancer, especially relapse and refractory follicular lymphomas; infectious diseases like tuberculosis, typhoid, etc.; and transplantation therapeutics, inflammation, etc.

Therapeutic modalities targeting the B7 and CD28 costimulatory families of ligands and receptors are showing promise in the clinic. The encouraging results in targeting the B7/CD28/CTLA-4 pathway in the autoimmune diseases like psoriasis, multiple sclerosis, and rheumatoid arthritis underscore the importance of this costimulatory pathway, and validate the pathogenic contributions of T cells in the etiology of these diseases. This novel strategy of costimulation activation/inhibition can be effectively exploited to develop immunotherapy for the treatment of intracellular pathogens like *M. tuberculosis*, HIV, *L. donovani*, *T. cruzi*, etc. This strategy can also be used as an alternative strategy or in combination with the drugs. Because this approach is based on modulating the immune system of the hosts rather than targeting the pathogen, it significantly diminishes the chance of emergence of drug resistant strains of pathogens and if applied properly may overcome the rising menace of infectious diseases. To develop alternative or adjunct (with drugs) therapies using costimulatory molecules, an intensive effort has been undertaken in the last decade to understand how intracellular pathogens exploit costimulatory molecules, which are the tour de force of the immune system. The potent role of costimulatory molecules is aptly established in the optimum activation of T cells and APCs: cells that play a cardinal role in curbing the infections. Hence, immunotherapy involving costimulatory molecules can be a breakthrough strategy to treat various diseases, minimize side effects inflicted by drug therapies, and restrict the emergence of drug resistance.

This book highlights the novel concept of reverse costimulation, which can be effectively exploited to develop immunotherapy using either humanized antibodies against CD80, CD86, and other costimulatory molecules or CD28 fusinogenic proteins for the treatment of diseases like allergies, asthma, rheumatoid arthritis, multiple sclerosis, lupus nephritis and severe psoriasis vulgaris, tuberculosis, typhoid, therapeutic transplantation, cancer, and inflammation. Moreover the development of second and third generation CARs for the treatment of various cancers and viral infections is also discussed in Chapter 6 entitled “T cell Costimulation and Its Role in Diseases.”

This book highlights the importance of complex roles played by CD28 and B7 costimulatory families in regulating the immune system. Despite the complex roles and interactions within the CD28 and B7 costimulatory families, the novel approaches targeting these families will yield new therapies for the treatment of inflammation, autoimmunity, transplantation, cancer, and infectious diseases. To translate this field into the clinic, it is urgent to develop novel methods to target the costimulatory pathway as currently understood and deeply comprehend the pathophysiology of the diseases involving costimulatory molecules.

Moreover no such book regarding developing costimulatory molecules for the treatment of diseases like allergy, asthma, lymphomas, cancer, autoimmunity, and transplantation was available to the best of my knowledge. Therefore it became imperative to pen a book with the concept of reverse costimulation immunotherapy in the treatment of diseases by developing various costimulatory molecules. Additionally up-to-date trends in the research like development of second and third generation CARs are included in this book with many tables and diagrams to make it more concise and readable. This book highlights mainly the following points:

- Describes the breakthrough strategy of immunotherapy involving costimulatory molecules to treat various diseases, minimizing side effects inflicted by drug therapies
- Contains many diagrams and tables highlighting costimulatory interactions
- Provides cutting-edge knowledge on costimulation and immunotherapy with relation to different diseases and their treatment
- Includes coverage of therapeutic and preventative methods
- Brings the basic science and clinical perspectives together in a single volume, facilitating translational possibilities
- Helps to integrate the value of costimulation immunotherapy outside of a cancer setting
- Brings out the development of second and third generation CARs for treatment of cancer and viral infections

The target audience will get up-to-date cutting-edge knowledge of costimulation and immunotherapy with relation to different diseases and their treatment. Moreover information is given on how the immune system is regulated by the costimulatory molecules and which costimulatory/coinhibitory pathways are important in disease control. This book highlights the regulation of the immune functioning by different costimulatory molecules and the treatment of lymphomas using reverse costimulation immunotherapy. Further, information is provided on how pathogens cause dreadful diseases and modulate the expression of various costimulatory molecules.

Acknowledgments

First I would like to thank Almighty for giving me strength, belief, and good health. It is under his grace that we all live, learn, and flourish.

Although it is difficult to thank everybody and interweave in words the genuine efforts made by the people directly or indirectly to make this book entitled *Developing Costimulatory Molecules for Immunotherapy of Diseases*, I would like to take this opportunity to pen down my appreciation for a number of people whose contribution in numerous ways immensely helped me to move toward my destination.

It gives me immense pleasure to express my sense of gratitude and respect to my mentors Dr. Javed N Agrewala, Prof Talat Ahmad, Prof Mohammad Afzal Zargar, Prof Inayetullah Tahir, Prof Raid Saleem Albaradie, and Prof Farooq Ahmad Malik whose constant support, utmost patience, invaluable advice, and guidance rekindled my interest for science from time to time. I will benefit for a long time from their sincerity, originality, and truthfulness which has nourished my intellectual maturity. I admire and respect them all for their sincerity, dedication, devotion, amazing memory, thorough knowledge, and constructive criticism.

The congenial company of Qazi Imtiyaz Sb, Raid, Fuzail, Faizan, and Aiman at Majmaah University and Tajamal, Reiaz, Altaf and Rffat, showkat at Kashmir University deserve a special mention. I am short of words to justify the care, affection, and untiring support showered by these very good friends.

I owe deepest gratitude to my parents Mohammad Abdullah Mir and Zoona Begum. A seed of loyalty, hard work, dedication, and sincerity sown in me by my parents played a crucial role in completion of this work. They have been a source of great strength throughout my life. I would like to thank other family members for continuous and unconditional support of all my undertakings, scholastic and otherwise. Without their affection, support, best wishes, love, and care this work would not have been completed.

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Manzoor Ahmad Mir

Chapter 1

Introduction to Costimulation and Costimulatory Molecules

INTRODUCTION

We live in an environment that contains a huge range of pathogenic microbes and toxic substances that threaten normal homeostasis and challenge the host by a very broad selection of pathogenic mechanisms. Pathogenic microbes possess a diverse collection of mechanisms by which they replicate, spread, and threaten normal host functions. The immune system must eliminate pathological microbes and toxic or allergenic proteins, and at the same time, it must avoid responses that produce excessive damage of self-tissues or that might eliminate beneficial, commensal microbes. It is not surprising, therefore, that the immune system uses a complex array of protective mechanisms to control and usually eliminate these organisms and toxins. A general feature of the immune system is that these mechanisms rely on detecting structural features of the pathogen or toxin that mark it as distinct from host cells. Such host–pathogen or host–toxin discrimination is essential to permit the host to eliminate the threat without damaging its own tissues.

Our immune system uses many mechanisms to combat infection by microbes. These mechanisms work together, and the fully integrated immune response draws elements from many effector systems in order to tailor a response to the specific invading pathogen. Activation of the immune system is dependent upon both innate and adaptive responses to environmental challenges such as infection. Activation of adaptive immune responses requires signals through antigen-specific T- and B-cell receptors (TCRs and BCRs, respectively). But this antigen-specific signal alone was not sufficient to drive the activation of naive T cells and led to the concept of two-signal model of T-cell activation, according to which productive T-cell activation requires a first signal provided by the interaction of antigenic peptide/major histocompatibility complex (MHC) with the TCR and a second, antigen-independent cosignal, the “costimulatory signal.” The “first signal” delivered by the TCR mediates the specificity of a T-cell response via the recognition of specific epitopes of a given antigen presented in combination with

the MHC on antigen-presenting cells (APCs). However, activation of T cells by this receptor interaction alone fails to induce cytokine production and sustained proliferation, but rather results in T-cell apoptosis, or the induction of specific nonresponsiveness (anergy) to subsequent stimulation with the same antigen.¹ This gives rise to the concept of the two-signal model of T-cell activation (Figure 1.1). According to the two-signal model, T cells, to become fully activated, require additional signals delivered by the so-called costimulatory molecules. These molecules are transmembrane proteins that induce an intracellular signaling cascade via their cytoplasmic tail that modifies the TCR-mediated signal. Costimulatory molecules cannot activate T cells without concomitant TCR cross-linking. According to this definition, adhesion molecules such as intercellular adhesion molecule-1 are not costimulatory because they enhance T-cell activation merely by facilitating the contact between T cell and APC. Furthermore,

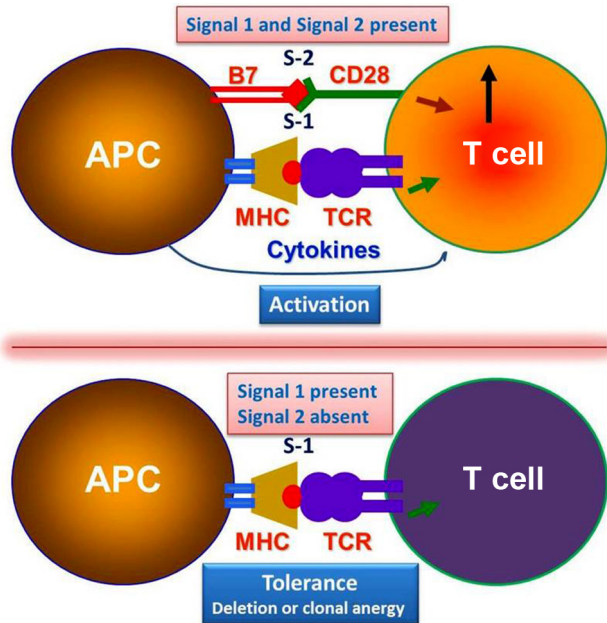


FIGURE 1.1 Two-signal model of T-cell activation. Two signals are needed for optimal activation of T cells. Signal 1, which is specific, is delivered through TCR on T cells and Signal 2, which is antigen independent and nonspecific, is delivered through costimulatory molecules present on APCs. Signal 2 is required for sustained T-cell proliferation, effector/memory cell generation, and prevention of anergy or apoptosis. If costimulatory signal is absent, the cell will undergo clonal anergy (unresponsiveness) or deletion. Expression of costimulatory molecules on professional APCs ensures that only pathogen-specific T cells are activated and minimizes the chances of acquired immune responses being mounted against self.

T-cell costimulation refers to a signal that is delivered to the T cell exclusively. In this respect, the CD40–CD40L interaction is also not considered as costimulatory, although this is an important receptor–ligand pair in T/B cooperation. The dependence of T-cell activation on this “second signal” delivered by costimulators adds a second line of regulation to antigen-specific T-cell responses that reaches far beyond a mere “on–off” command. With a growing number of both stimulatory and inhibitory costimulatory molecules being identified, the classic concept of costimulation as a two-signal event has changed. T cells simultaneously express an adjustable spectrum of costimulatory molecules. Today, T-cell costimulation is recognized as an integrating process of various positive and negative signals that determine the mode of T-cell activation.² The T-cell response to any given antigen involves the clonal expansion of a small number of T cells that share specificity but express unique TCRs. The integration of signals generated by these interactions determines the final outcome of encounters between T cells and APCs. The primary source of this costimulatory signal is interactions between the T-cell integral membrane protein CD28 and its ligands, B7-1 and B7-2, on the APC. Currently, the B7 family has seven members (B7-1, B7-2, inducible costimulator ligand (ICOS-L), programmed death ligand-1 (PD-L1), PD-L2, B7-H3, and B7-H4), whereas tumor necrosis factor (TNF) receptor (TNFR)/TNF family has six members (herpes virus entry mediator (HVEM), B- and T-lymphocyte attenuator (BTLA), CD70, CD30, 4-1BB-L, and OX40L) (Table 1.1 and Figure 1.2). The B7 family includes transmembrane or glycosylphosphatidylinositol-linked proteins characterized by extracellular IgV and IgC domains. B7-1 and B7-2 can engage either CD28 or cytotoxic T-lymphocyte antigen-4 (CTLA-4), resulting in complex positive and negative regulation of antigen receptor signaling.³ ICOS-L is the only identified ligand for inducible costimulator (ICOS). Both PD-L1 and PD-L2 engage programmed death-1 (PD-1), although data suggest the existence of additional receptors for PD-L1 and PD-L2. Receptors for B7-H3 and B7-H4 have not been identified, although expression of these ligands modulates immune responses. The cell surface expression of B7 ligands is highly regulated, although, in general, ligands for costimulatory receptors are constitutively expressed on APCs, whereas ligands for coinhibitory receptors are expressed after induction by various inflammatory stimuli. The activation of a T cell is thus dictated by the integration of signals derived from the TCR, CD28, and CTLA-4. By regulating the expansion of individual T cells, CD28 and CTLA-4 could modulate the composition of a polyclonal antigen-specific T-cell response. The balance between activating signals generated by CD28 and inhibitory signals delivered by CTLA-4 determine the outcome of a T cell’s interaction with an APC.

COSTIMULATORY MOLECULES

It is now well established that the optimum activation of T cells requires two distinct signals from APCs. The first signal is triggered by interaction

TABLE 1.1 Currently Known Costimulatory Molecules and Ligands

Expression	Superfamily	Costimulatory Molecule	Ligand	Signal Type
Constitutive	CD28/B7	CD28	B7-1, B7-2 (CD80, CD86)	Positive
	TNF/TNFR	CD27	CD70	Positive
	TNF/TNFR, CD28/B7	HVEM	LIGHT, BTLA	Positive
	TNF/TNFR, CD28/B7	BTLA	HVEM	Negative
Inducible	CD28/B7	ICOS	ICOSL	Positive
	TNF/TNFR	OX40 (CD134)	OX40L	Positive
	TNF/TNFR	4-1BB (CD137)	4-1BB-L	Positive
	TNF/TNFR	CD30	CD30L (CD153)	Positive
	CD2	SLAM (CD150)	SLAM (CD150)	Positive
	CD28/B7	CTLA-4 (CD152)	B7-1, B7-2 (CD80, CD86)	Negative
	CD28/B7	PD-1	PD-L1, PDL-2	Negative
	CD28/B7	Unknown	B7-H4	Negative
	CD28/B7	Unknown	B7-H3	Obscure

Note: TNF: tumor necrosis factor; R: receptor; HVEM: herpes virus entry mediator; LIGHT: homologous to lymphotoxin, inducible expression, competing for GpD of herpes virus, expressed on activated T lymphocytes; BTLA: B- and T-lymphocyte attenuator; ICOS: inducible costimulator; L: ligand; SLAM: signaling lymphocyte activation molecule; CTLA: cytotoxic T-lymphocyte antigen; PD: programmed death; B7-H4 and -H3: B7 homologues 4 and 3.

of antigen-specific TCR with the peptide MHC complex,⁴ whereas the second nonspecific signal is delivered by the interaction of costimulatory molecules present on the surface of the APCs and their corresponding receptors on T cells.^{5–8} The costimulatory signals are neither antigen specific nor MHC restricted, yet they are essential and critical for the development of effector functions. T-cell activation requires engagement of the TCR with the peptide–MHC complex presented on the cell surface of APCs.⁹ In addition to this antigen-specific interaction, a second interaction involving costimulatory receptors (CD28, ICOS) on T cells and their respective ligands (B7-1/B7-2, ICOSL) on APCs is required for optimal T-cell activation.^{10,11} Although these receptor–ligand pairs augment T-cell activation, interactions between T-cell costimulatory/inhibitory receptors CTLA-4 and PD-1 with their ligands, B7-1/B7-2 and PD-L1/PD-L2,

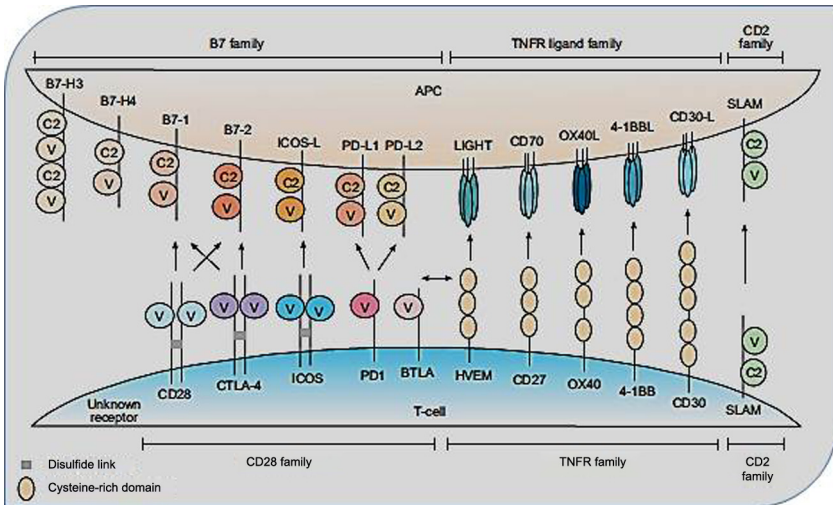


FIGURE 1.2 Currently known costimulatory molecules and their ligands. TNFR: tumor necrosis factor receptor; APC: antigen-presenting cell; B7-H3 and -H4: B7 homologues 3 and 4; ICOS: inducible costimulator; L: ligand; PD: programmed death; LIGHT: homologous to lymphotoxin, inducible expression, competing for GpD of herpes virus, expressed on activated T lymphocytes; V: Ig-like variable domain; C2: constant type-2 Ig-like domain; SLAM: signaling lymphocyte activation molecule; CTLA: cytotoxic T-lymphocyte antigen; BTLA: B- and T-lymphocyte attenuator; HVEM: herpes virus entry mediator.

respectively, lead to attenuation of the T-cell response.^{12,13} The integration of signals delivered as a consequence of these interactions is essential for the initiation, modulation, and regulation of an effective T-cell response. The immunoregulatory potential of costimulation became clear when it was shown that TCR ligation in the absence of costimulatory signal can lead to anergy, deletion, or unresponsiveness in T cells.^{14–17} This gave rise to two-signal model of T-cell costimulation.

Two-Signal Model of T-cell Activation

The basic concept that optimal and sustained activation of T cells, leading to proliferation, cytokine production, and effector functions, requires two signals has been known as “the two-signal model.”² The first signal is delivered through the TCR after interaction with antigenic peptides presented on MHC molecules of APCs. This MHC–peptide–TCR signal is specific and determines which CD4⁺ T cells will respond to a particular antigen. The second signal is also accessory cell derived, but not antigen specific and has been termed as a costimulatory signal because, although essential, it does not by itself induce any response in T cells.^{1,18,19} However, when a T cell has its receptor ligated and receives a costimulatory signal, the T cell proliferates

and differentiates into an armed effector cell. These secondary, the so-called costimulatory, signals are critically required for the process of T-cell activation. Because landmark studies defined that T cells receiving a TCR signal without a costimulatory signal are tolerized *in vitro*, the investigation of T-cell costimulation has attracted intense interest. Early studies demonstrated that interrupting T-cell costimulation allows attenuation of the alloresponse, which is particularly difficult to modulate due to the clone size of alloreactive T cells.

The understanding of costimulation has since evolved substantially and now encompasses not only positive signals involved in T-cell activation but also negative signals that inhibit T-cell activation and promote T-cell tolerance. Moreover, the significance of costimulatory molecules in T-cell activation has gained considerable impetus following the observation that T cells that bind to antigens but do not receive costimulatory signals are thought to die or become anergic, a state of unresponsiveness, in which the cell cannot be activated even if it receives both the signals vital for its activation.^{14,19,20} Thus, an encounter with the antigen can lead to two quite distinct outcomes, proliferation and differentiation into effector cells, or inactivation or death. Which outcome occurs is determined by the appropriate delivery of costimulatory signals through costimulatory molecules.¹⁵ At present, several reports are available in the literature highlighting the potential role of a number of cell surface molecules necessary for the initiation of the differentiation events occurring during T cell–APC interaction. The various known costimulatory molecules expressed on the APC have both adhesive and costimulatory functions.

The known costimulatory molecules are CD80, CD86, CD40, B7-DC (PDL-2), B7H-3/B7H-4, CD70, OX40L, B7-H2/LICOS, B7-H1, and 4-1BBL. The recently discovered costimulatory molecules include the B7-DC (PDL-2), B7H-3, and B7H-4.²¹ The exact mechanism of sequence of signals provided by these molecules in the activation and differentiation of cells on which they are present is not very well known. The best-characterized costimulatory pathway involves the CD28/CTLA-4 receptors present on T cells that bind to costimulatory molecules, CD80 and CD86.¹² CD28 is constitutively expressed on all T cells in mice and on 95% of CD4+ T cells and 50% of CD8+ T cells in humans. CD80 and CD86 are expressed mainly on APCs, including dendritic cells (DCs), macrophages, and B cells. The expression of CD80 and CD86 on APCs is enhanced by the presence of microbes and by cytokines that are produced in response to microbes. This regulated expression of B7 costimulators ensures that T cells respond best only when necessary, that is, when faced with pathogens. The interaction of CD80 and CD86 with CD28, in concert with T-cell–receptor signaling, promotes the expansion of antigen-stimulated T cells and their differentiation into effector and memory cells. CD28 is the major costimulatory receptor for naive T cells and is therefore important for initiating T-cell responses. CD28 signals enhance the production of interleukin-2 (IL-2) and other cytokines; upregulate cell survival genes (such

as Bcl-xL); promote energy metabolism (glucose uptake and rate of glycolysis); and facilitate cell cycle progression.²² One of the best-characterized costimulatory interactions is the binding of CD28/CTLA-4 on T cells with B7-1/B7-2 on APCs. CD28 and CTLA-4 are homologous receptors with approximately 30% sequence identity that belong to the immunoglobulin (Ig) superfamily and contain one membrane proximal IgV-like domain.^{23,24} Furthermore, they are clustered in close proximity on chromosome 1C1 in mice and 2q33 in humans.^{25,26} They are expressed as disulfide-linked homodimers and bind to the same ligands, B7-1 and B7-2. Although CTLA-4 is bivalent in nature, recent structural studies suggest that CD28 is monovalent.²⁷ Moreover, the expression pattern and the immunological roles of CD28 and CTLA-4 are vastly different. Although CD28 is constitutively expressed on both naive and activated T cells, CTLA-4 follows a more complex pattern of expression. In resting T cells, CTLA-4 is associated with the clathrin adaptor complex AP-1 and is present predominantly in intracellular vesicles. Upon T-cell activation, CTLA-4 is transported to the cell surface where it is again rapidly internalized by another clathrin adaptor complex AP-2.^{28,29} The directional recruitment of CTLA-4 from intracellular vesicles to the immunological synapse is regulated by the TCR signal strength. Furthermore, CD28 has a slow turnover rate, whereas CTLA-4 has a rapid turnover with a half-life of 2 h.³⁰ As a consequence of these dynamic processes, only a small fraction of CTLA-4 is present on the cell surface, which seems to be sufficient to deliver the requisite signal for termination of T-cell activation. Functionally, the engagement of CD28 with B7 ligands delivers a positive signal to T cells that culminates in T-cell proliferation and cytokine production, and prevents induction of T-cell tolerance.^{31,32} In contrast, the interaction of CTLA-4 with B7-1/B7-2 results in negative signaling, which leads to attenuation of T-cell activation and induction of T-cell anergy. Such processes are essential for regulating a T-cell response and maintaining T-cell homeostasis.^{33–35} The engagement of CD28 and CTLA-4 also indirectly delivers distinct signals to T cells through their different effects on DCs. The expression pattern of various known B7-CD28 family of costimulatory molecules along with their corresponding ligands is listed in [Table 1.2](#) and described below in detail.

THE B7 FAMILY OF COSTIMULATORY LIGANDS

The B7:CD28 family of costimulatory molecules includes the costimulatory ligands present on the APCs and their corresponding receptors on the T cells. The B7 costimulatory ligands involve the CD80, CD86, ICOS-L, PDL-1 (B7-H1), PDL-2 (B7-DC), B7-H3, and B7-H4 (B7x/B7-S1) ([Figure 1.3](#)).^{10,36,37} The receptors for the B7 family members present on T cells include the members of CD28 family such as CD28, CD152 (CTLA-4), ICOS, PD-1, and BTLA.^{11,12} The functional characterization of all the costimulatory molecules

TABLE 1.2 Protein Expression of CD28 and B7 Superfamily Members

Costimulatory Ligands/Receptors	Type	Hematopoietic Cells	Nonhematopoietic Cells	
Ligands	B7-1 (CD80)	Induced on T cells, B cells, myeloid cells, and macrophages		
	B7-2 (CD86)	Constitutive on B cells, DCs, monocytes/macrophages, and upregulated with activation		
	ICOSL	Induced on T cells, B cells, DCs, and monocytes/macrophages	Intestinal epithelial cells and muscle cells	
	PDL-1	Constitutive on B cells, DCs, monocytes/macrophages, and upregulated with activation	Induced on T cells	Endothelial cells, Kupfer cells, thymic, intestinal, and renal tubular epithelial cells, syncytiotrophoblasts, pancreatic islets, keratinocytes, and tumor cells (carcinomas, melanoma)
		PDL-2		Induced on monocytes/macrophages, and DCs
	B7-H3	Induced on T cells, B cells, NK cells, DCs, and monocytes/macrophages	Osteoblasts	
	B7-H4	Induced on T cells, B cells, DCs, and monocytes/macrophages	Tumor (lung, ovarian)	
Receptors	CD28	Constitutive on T cells		
	CTLA-4	Induced on T cells		
	ICOS	Induced on T cells, B cells, and NK cells		
	PD-1	Induced on T cells, B cells, myeloid cells, and macrophages		

Note: ICOS: inducible costimulator; CTLA: cytotoxic T-lymphocyte antigen; PD: programmed death; B7-H3 and B7-H4: B7 homologues 3 and 4; NK: nuclear killer.

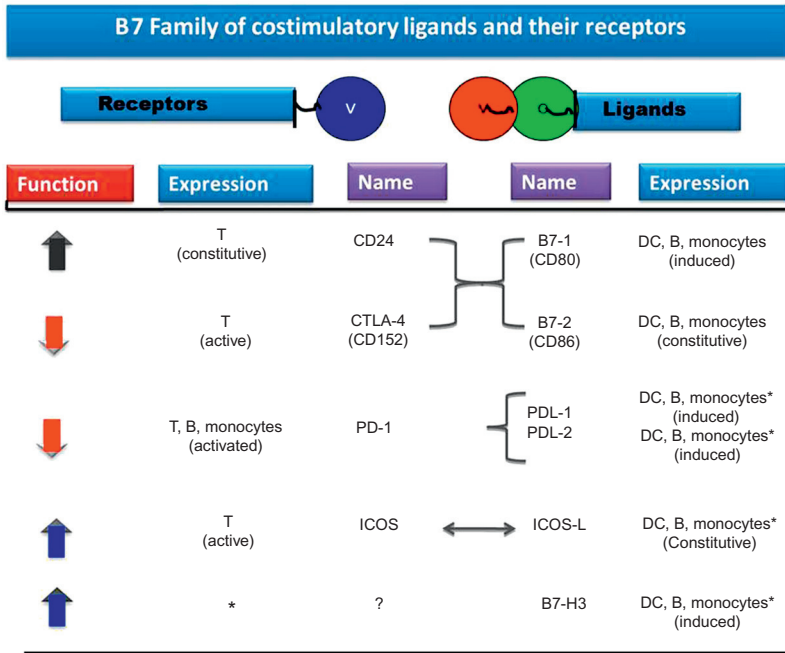


FIGURE 1.3 Summary of B7 family ligands and their receptors. The names of receptors and ligands are indicated, as well as a brief summary of predominant expression patterns for each. The conserved structure of a single IgV extracellular domain for receptors and IgV and IgC extracellular domains for ligands is depicted at the top. Function arrows summarize whether the pathway is thought predominantly to costimulate or inhibit the response of the receptor-bearing cell. Integration of signals through this family of costimulatory and inhibitory receptors and their ligands is critical for activation of immune responses and tolerance.

and their ligands is listed in [Table 1.3](#). The discovery of new functions for the original B7 family members together with the identification of additional B7 and CD28 family members has revealed new ways in which the B7:CD28 family regulates T-cell activation and tolerance. The interaction between these molecules does not regulate T cells but also the cells on which these molecules are present, that is, APCs.¹²

Pathways in the B7:CD28 family have key roles in regulating T-cell activation and tolerance, and are thus promising therapeutic targets. In the B7 family, CD80 and CD86 are the best-defined and best-characterized costimulatory molecules present on APCs. T-cell activation is dependent upon signals delivered through the antigen-specific TCR and accessory receptors on the T cell. A primary costimulatory signal is delivered through the CD28 receptor after engagement with its ligands, CD80 or CD86, resulting in T-cell activation, whereas the engagement of CTLA-4 (CD152) with CD80 or CD86 ligands results in attenuation of T-cell responses. CD80/CD86:CD28/CD152

TABLE 1.3 Functional Characterization of Costimulatory Molecules

Functional Aspect	Characterization	Costimulatory Molecule
Expression pattern	Constitutive	CD28, CD27, HVEM, BTLA
	Inducible	ICOS, CTLA-4, PD-1, OX40, 4-1BB, CD80, SLAM, putative receptor for B7-H3 and B7-H4
T-cell modulation	Positive/enhancement	CD28, ICOS, OX40, CD27, 4-1BB, CD30, HVEM, SLAM
	Negative/Inhibition	CTLA-4, PD-1, BTLA-4, putative receptor for B7-H4
T-cell differentiation	Th1	4-1BB, SLAM, BTLA
	Th2	ICOS, OX40, CD30
	Treg	ICOS, CTLA-4, PD-1
T-cell function	T effector/helper	ICOS, CTLA-4, PD-1, OX40
	T memory	OX40, CD30, CD27
Location of action	Central	CD28, HVEM, CTLA-4
	Central or peripheral	ICOS, PD-1, OX40, 4-1BB

Note: HVEM: herpes virus entry mediator; BTLA: B- and T-lymphocyte attenuator; ICOS: inducible costimulator; CTLA: cytotoxic T-lymphocyte antigen; PD: programmed death; SLAM: signaling lymphocyte activation molecule; B7-H3 and B7-H4: B7 homologues 3 and 4; Th: T-helper cell; Treg: regulatory T cell.

interactions not only promote initial T-cell activation but also regulate self-tolerance by supporting CD4⁺ CD25⁺ regulatory T cell (Treg) homeostasis.¹²

The well-defined and best-characterized costimulatory ligands in T-cell activation are the two structurally related type I transmembrane glycoproteins CD80 and CD86 belonging to Ig superfamily. They share 25% sequence homology and have two Ig-like extracellular domains (one variable and one constant), in addition to transmembrane and short cytoplasmic domains. These proteins are expressed as monomers/dimers on professional APCs, namely, DC, activated B cell, and macrophage. CD80 expression is found on activated B and T cells and macrophages. CD86 is constitutively expressed on interdigitating DCs, Langerhans cells, peripheral blood DCs, memory B cells, and germinal center B cells. Both CD80 and CD86 are capable of binding the receptors, CD28 and CTLA-4. Both human and mouse CD80 and CD86 can bind to either human or mouse CD28 and CTLA-4. This suggests that there are conserved amino acids that form the critical binding sites. CD28 plays a role in activation, whereas CTLA-4 functions as an inhibitory receptor important for down-modulating the immune response.^{3,13,38,39} These costimulatory

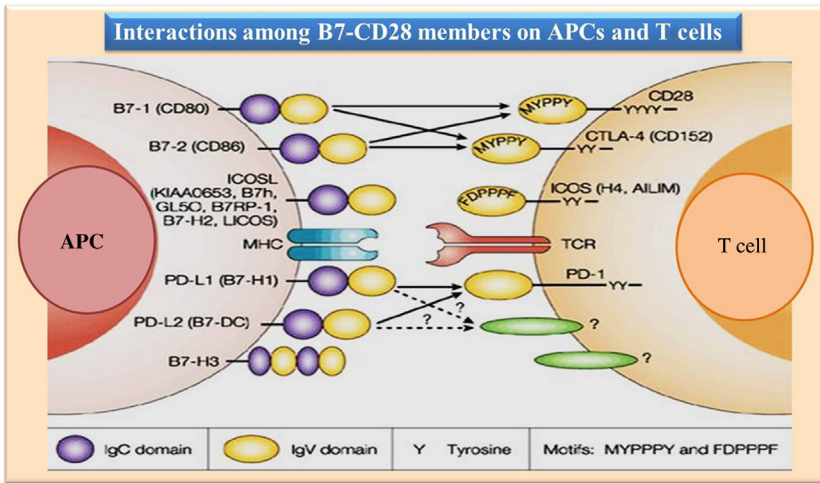


FIGURE 1.4 Schematic representation of the interactions among CD28 and B7 family members. Alternate names for receptors and ligands are shown in parentheses. IDO, indoleamine 2,3-dioxygenase.

molecules on APCs interact with their corresponding ligands on T cells and cause the inhibition/activation of the cell function depending upon the molecules involved and the intracellular domain of the molecules involved. For interactions of these molecules on APC and T cell, see [Figure 1.4](#). Although CD86 is generally the first B7 molecule encountered, due to its constitutive expression on numerous APCs, there does not appear to be significant differences in the functions of CD80 and CD86. Rather, it depends on the type of APC encountered and its activation state.

The B7 and CD28 gene families have grown substantially with mining of the human genome sequence, and the pathways defined by these new members regulate the activation, inhibition, and fine-tuning of T-cell responses. A great surprise has been the abundance of inhibitory signals delivered by these new B7/CD28 family members. The newer CD28 family members, ICOS, PD-1, and BTLA are inducibly expressed on T cells and have important roles in controlling the balance between effector T cells and Tregs. PD-1 and BTLA are also expressed on B cells and APCs, and appear to have broader immunoregulatory functions, which necessitate cell-type specific analyses of their functions. The new B7 family members, ICOS ligand (B7h, GL50, B7RP-1, ligand of ICOS (LICOS), and B7-H2); PD-L1 (B7-H1); PD-L2 (B7-DC); B7-H3 and B7-H4 (B7x/B7-S1); B7S3; and BTNL2, can be expressed not only on professional APCs but also on nonhematopoietic cells. The nonhematopoietic cell expression of these B7 family members suggests that they may function to regulate T-cell activation and/or tolerance in nonlymphoid organs.

CD80 Costimulatory Molecule

CD80, a type-1 transmembrane glycoprotein, was first identified on Epstein–Barr virus-activated B-cell blasts, Burkitt’s lymphomas, and B lymphoblastoid cell lines.⁴⁰ It was the first ligand to be identified for CD28 and later for CTLA-4.^{41,42} Both the human and murine CD80 genes were cloned and shown to be the members of the Ig superfamily. Human and mouse CD80 shares approximately 44% identity.^{43,44} CD80 has 10-fold higher affinities for both CD28 and CTLA-4 than for CD86. CD80 expression is found on activated B cells, activated T cells, and macrophages. The functional importance of CD80 molecule has been demonstrated in a number of studies of T-cell activation. The potent costimulatory role of CD80 has been demonstrated *in vivo* in transgenic mice in which CD80 was ectopically expressed on the cells of the islets of Langerhans.⁴⁵

The role of CD80 as a costimulatory molecule was established by the ability of CD80-transfected cells to provide the costimulation to T cells. CD80 transfected Chinese hamster ovary cells, synergies with anti-CD3 antibody (Ab)-induced T-cell activation, ultimately leading to T-cell proliferation and the production of IL-2. The proliferation was enhanced in a CD28-dependent fashion because T-cell activation was blocked by anti-CD28 Ab.⁴⁶ In subsequent studies, CD80 was also shown to promote the breakdown of tolerance. Despite the apparent ability of CD80 to provide sufficient costimulation, it has been very difficult to demonstrate its function on normal APCs in mice. Anti-CD80 Ab minimally blocks a primary Mixed Lymphocyte Reaction (MLR), whereas human Cytotoxic T Lymphocyte Antigen 4 Immunoglobulin (hCTLA-4Ig) inhibits the response by 80%.⁴⁷ Additionally, the staining of either lipopolysaccharide (LPS)-activated B cells or whole spleen with hCTLA-4Ig was not inhibited by anti-CD80 Ab, which suggests the existence of an additional CTLA-4 ligand.^{48,49} CD80 $-/-$ mice are capable of mounting an immune response to nominal antigens and APCs isolated from these mice could be stained with labeled hCTLA-4Ig.⁵⁰ These observations led to the identification and eventual cloning of a second B7 family member, CD86.^{50,51}

CD86 Costimulatory Molecule

CD86 is a 70-kDa glycoprotein made up of 329 amino acids, a transmembrane region, and a longer cytoplasmic domain than CD80.⁵⁰ CD86 is constitutively expressed on interdigitating DCs, Langerhans cells, peripheral blood DCs, memory B cells and germinal center B cells, and macrophages. CD86 is rapidly upregulated on B cells following activation by cross-linking of the Ig receptor or the addition of a variety of cytokines.⁴⁷ Additionally, CD86 is expressed at low levels on monocytes and is upregulated through interferon- γ (IFN- γ) stimulation. CD86 shows 25% identity with CD80 and are both coded on human chromosome 3q13.33q21. Similar to CD80 transfectant, CD86 transfectants augment T-cell proliferation and IL-2 production to suboptimal

stimulation with anti-CD3 Ab or phorbol myristate acetate (PMA). This costimulation was inhibited by either hCTLA-4Ig or anti-CD28 Ab, but not by anti-CD80 Abs, demonstrating therefore that CD86 binds to both CD28 and CTLA-4.^{50,52} A combination of both anti-CD80 and anti-CD86 Abs were the most effective at inhibiting the MLR. The role for both molecules in primary responses is further supported by the findings that a combination of anti-CD80 and anti-CD86 Abs can induce anergy.^{53–55}

CD80 and CD86 Expression Kinetics

CD28 shares the ability to bind B7 family members with its homologue CTLA-4. However, B7 binds to CTLA-4 with up to 2500-fold higher avidity than to CD28.²³ Upon ligation to B7, CD28 and CTLA-4 deliver opposing signals for T-cell activation, with CD28 generating an activating costimulatory signal and CTLA-4 functioning to inhibit the response.³⁹ CD28 ligation enhances T-cell proliferation through multiple mechanisms, including increased production and stability of IL-2 mRNA and upregulation of antiapoptotic genes, such as Bcl-xL. In contrast, CTLA-4 signaling reduces the production of the growth factor IL-2³⁴ and cyclin D3, cyclin-dependent kinase (cdk)4, and cdk6, thus restricting T-cell expansion.^{37,56} The expression of CD80 and CD86 is primarily limited to APCs; however, recently they have been shown to be expressed on T cells, but their functional significance is not very well understood. CD86 is constitutively expressed on some resting T cells, whereas CD80 is not present on resting T cells; however, both can be upregulated on T cells by different stimuli.⁵⁷ Resting B cells express no detectable CD80 and very low levels of CD86. Both CD80 and CD86 can be upregulated on B cells by LPS or Concovalin A (ConA). CD86 is expressed more rapidly than CD80 on B cells stimulated with LPS or anti-CD40 Ab.⁵⁸ CD80 is generally absent on unstimulated cells, whereas CD86 is constitutively expressed at moderate levels on DCs, macrophages, and monocytes.⁵¹ CD80 and CD86 are rapidly upregulated by Granulocyte Maturing-Colony Stimulating Factor (GM-CSF), LPS, IFN- γ , and CD40 signaling on DCs and macrophages.^{59,60} The induction of CD86 expression occurs within 6 h of stimulation, with maximal level of expression achieved between 18 and 24 h. In contrast, CD80 expression is not detected until 24 h poststimulation and does not reach maximal levels until 48–72 h.^{47,48} Some microorganisms including *Mycobacterium leprae* inhibit B7 expression, whereas some including *Toxoplasma gondii* upregulate the CD86 expression.⁶¹ CD80 expression is induced, while that of CD86 is upregulated when human monocytes encounter the viable *T. gondii* pathogen.^{61,62} A number of cytokines have been shown to differentially regulate CD80 and CD86 expression. IL-4 is one of the most potent inducers of CD86 and, to a lesser extent, CD80 on B cells.^{63,64} Incubation of small resting B cells with IL-4 upregulates CD86 expression within 6 h with maximal induction occurring by 24 h.⁶⁴ IFN- γ increases the expression of CD86 on B cells, peritoneal macrophages, and

peripheral blood monocytes^{49,64} but downregulates the expression of CD80 on peritoneal macrophages.⁵⁹ Interestingly, the engagement of Fc receptor downregulates both CD80 and CD86 on monocytes that have been activated with either IFN- γ or GM-CSF.⁶⁵ IL-10 blocks both CD80 and CD86 upregulation on peritoneal macrophages and downregulates CD86, but not CD80 on human DCs.^{66,67} These results suggest that immunosuppressive properties of the IL-10 may, in part, be a result of its regulation of CD28/CTLA-4 ligands. Thus, the differences in the ability of the respective cytokines to regulate the level and temporal expression of CD80 and CD86 both qualitatively and quantitatively may result in the distinct effects during an immune response.

CD80 and CD86 Binding Kinetics to CD28 and CTLA-4

Recent evidence suggests that CD80 and CD86 have similar low avidities for CD28 and high avidities for CTLA-4.¹⁰ Both these proteins have very fast on-and-off rates, but their binding kinetics to CD28 and CTLA-4 are distinct. CD80 is a more potent ligand for CTLA-4 based on its higher affinity and avidity. CD80, in contrast to CD86, binds two to three times more strongly to both CD28 and CTLA-4, with faster binding kinetics and slower dissociation constants. CD80 binds CTLA-4 and CD28 with equilibrium dissociation constants (K_d) of 0.2 and 4 μM , respectively (Figure 1.5), whereas CD86 exhibits 5–10-fold lower affinities with K_d of 2.6 and 20 μM for CTLA-4 and CD28, respectively.¹⁰ Stoichiometrically, two B7 molecules, CD80 and CD86, bind to two sites on a CD28 or CTLA-4 homodimer. This stoichiometry and the different on/off rates for CD80 and CD86 suggest that the response to low concentration of B7 would not be linear. Although CD80 and CD86 bind to the same general region of CD28 and CTLA-4, the specific amino acids critical for CD80 or CD86 binding are different.⁴² Interaction of CD80 or CD86 with CTLA-4 results in an inhibitory signal, in contrast to the positive signal generated after its interaction with CD28.

Different Oligomeric States of CD80 and CD86

Costimulatory ligands CD80 and CD86 expressed on the surface of APCs interact with their receptors CTLA-4 and CD28 on T cells. Although CD80 and CTLA-4 are homologous ligands having common receptors, they exhibit distinct biochemical features and roles in immune regulation. It was recently demonstrated that CD80 and CD86 adopt different oligomeric states on the cell surface. CD86 is present as a monomer on the cell surface, whereas CD80 exists as a mixed population of monomers and dimers with predominating dimers.^{23,68,69} The point mutations made in the putative dimer interface of CD80 disrupt the dimeric state of the CD80 molecule and result in the generation of a population with enhanced monomers. This existence of the different oligomeric states of CD80 and CD86 indicates that

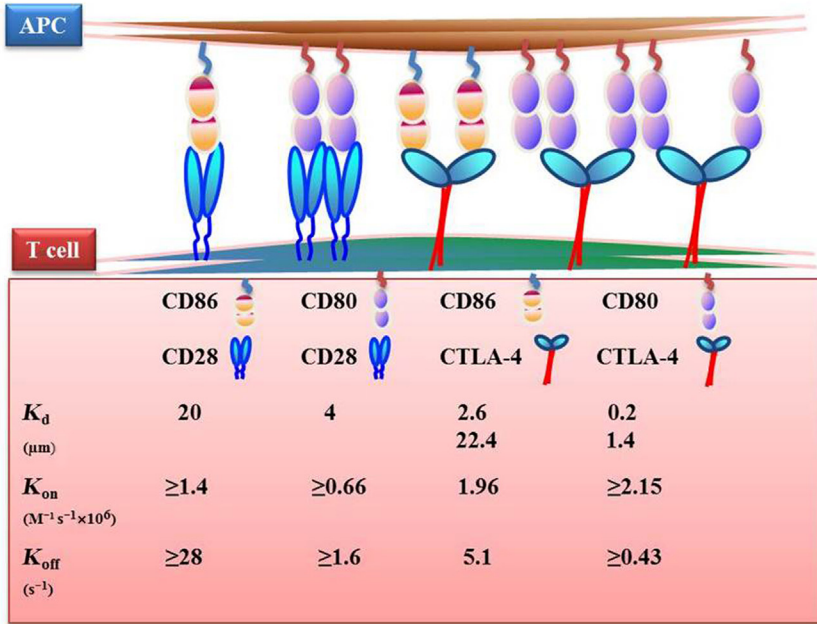
Interaction of CD86/CD80 with CD28/CTLA-4 molecules during costimulation

FIGURE 1.5 Interaction of CD80/CD86 with CD28/CTLA-4. The figure represents the interaction of CD80/CD86 with CTLA-4 and CD28. On the basis of expression, binding affinities, and crystal structure, it appears that CD80 shows better interaction with CTLA-4, and CD86 shows better interaction with CD28, thereby delivering bidirectional signaling, activating not only T cells but also APC. The association of CD80 with CTLA-4 is also considered better due to its faster association and slower dissociation (K_d) rates compared to CD28, which shows slower association and faster dissociation kinetics. The association of CD86 with CD28 is also considered better due to its faster association and slower dissociation rates compared to CTLA-4, which shows slower association and faster dissociation kinetics.

these molecules form costimulatory receptor–ligand complexes with distinct cell surface organizations, which may represent an important mechanistic determinant for the assembly and functional properties of signaling complexes at the T cell–APC interface.^{10,68,69} This presence of different oligomeric states of CD80 and CD86 along with the different dissociation constants of CD80 and CD86 for CD28 and CTLA-4 may be the reasons why CD80 and CD86 behave differently and show different functional properties. CTLA-4 is only expressed by T cells after activation.

PD-L1/B7-H1 Costimulatory Molecule

B7-H1, also called PD-L1, is a recently identified member of B7 family of costimulatory molecules. B7-H1 shares approximately 20% amino acid

identity with CD80 and 15% amino acid identity with CD86. Human and mouse B7-H1 (PD-L1) shares approximately 69% amino acid identity.^{70–72} B7-H1 is a ligand for PD-1, a member of the CD28/CTLA-4 family.⁷³ B7-H1 is widely expressed on normal tissues (e.g., liver, lung, pancreas, heart) but not on resting peripheral blood cells. Expression on monocytes, macrophages, and DCs is upregulated with IFN- γ stimulation or upon activation. Experiments using TCR transgenic CD4+ or CD8+ T cells stimulated with APCs expressing PD-L1 show that both T-cell subsets are susceptible to this inhibitory pathway. However, CD8+ T cells may be more sensitive to modulation by the PD-1:PD-L pathway because of their intrinsic inability to produce significant levels of IL-2.^{72,74} However, B7-H1 does not bind to CD28, CTLA-4, or ICOS.^{3,73} The molecule can act as a T-cell costimulatory ligand during suboptimal TCR stimulation. As with B7-H2, both proliferation and IL-10 secretion are costimulated in the presence of immobilized B7-H1 or B7-H1-transfected COS cells, whereas IL-2 production is minimally enhanced. PD-L1 appears to have a strong inhibitory influence on autoreactive T cells in the islets, possibly by preventing cytokine secretion at the tissue site.^{75,76} It has been shown to be an attractive target for the treatment of autoimmune diseases because it was found to be the main ligand restricting autoimmune diabetes in nonobese diabetic (NOD) mice.⁷⁷

Interactions between the inhibitory receptor PD-1 and its ligands, PD-L1 and PD-L2, regulate both the induction and maintenance of peripheral T-cell tolerance.⁷⁸ In addition, PD-1 and PD-L1 exert critical inhibitory functions in the setting of persistent antigen expression, as in chronic viral infections and tumors.^{79,80} Studies in the chronic lymphocytic choriomeningitis virus infection model provided the first indication that this pathway contributes to T-cell dysfunction during chronic viral infection.⁸¹ PD-1 is overexpressed on exhausted T cells, and immune function in these exhausted T cells is restored upon PD-1 or PD-L1 blockade. This observation has been extended to human immunodeficiency virus, Simian immunodeficiency virus, hepatitis B virus, and hepatitis C virus infections, and has led to new strategies for treating chronic viral infections.⁸² High PD-L1 expression on tumor cells correlates with poor prognosis in a variety of human cancers, and PD-1 or PD-L1 blockade has been shown to mediate some tumor protection in mice. A humanized anti-PD-1 antibody is in clinical trials in cancer patients. Relatively little is known about PD-1 signaling. Riley⁸⁰ reviews the current understanding of signaling pathways altered by PD-1 and discusses distinct mechanisms by which PD-1 and CTLA-4 inhibit T cells. One key difference is that PD-1 engagement blocks induction of phosphatidylinositol 3-kinase activity, whereas CTLA-4 does not. The targeting of distinct signaling molecules by PD-1 and CTLA-4 suggests that approaches that block PD-1 and CTLA-4 signaling should be synergistic.

B7-H2/B7RP-1/LICOS Costimulatory Molecule

B7-H2, also known as B7RP-1, B7-H2, LICOS, and GL50, is another member of the growing B7 family of immune costimulatory proteins.³ As with other B7 family members, B7-H2 is a type I transmembrane glycoprotein that is a member of the Ig superfamily. B7-H2 shares approximately 20–30% amino acid sequence identity with other members of the B7 family. Human and mouse B7-H2 shares approximately 49% amino acid sequence identity.^{83,84} B7-H2 is expressed constitutively on resting B cells, DCs, and at low levels on monocytes, but IFN- γ stimulation upregulates expression on these cell types. The mechanisms of upregulation of B7-H2 expression appear to be distinct from those for CD80 and CD86.⁸⁵ B7-H2 expression could be regulated by inflammatory signals present in peripheral sites. It is the ligand for ICOS and does not bind to CD28, CTLA-4, or PD-1.⁸⁶ Consistent with a role of B7-H2 as a costimulatory ligand, T-cell proliferation in response to antigen presentation by B7-H2-transfected fibroblasts is significantly higher than that with untransfected fibroblasts, similar to that observed with CD86-transfected cells. B7-H2 costimulation of T cells, however, does not depend on either CD28 or CTLA-4 molecules because antagonism of B7-CD28 and B7-CTLA-4 interactions had no effect on costimulation by fibroblasts expressing B7-H2. Further, human B7-H2, which is also known as LICOS, shares 50% of its extracellular domain sequences with murine B7-H2 and can specifically bind *in vitro* to human ICOS. Exposure of mice to endotoxin induces the expression of B7-H2 mRNA in testes, kidney, and peritoneal tissues. Interestingly, *in vitro* differentiated DCs do not express B7-H2, even though they express CD80. As one additional functional test of the importance of B7-H2–ICOS interactions in the growth and differentiation of mature T cells, transgenic mice have been engineered to secrete a soluble form of the B7-H2 molecule into the bloodstream.⁸⁷ These animals develop a lymphoid hyperplasia by 12 weeks of age involving both the T-cell and B-cell compartments. Although it is difficult to ascertain whether this soluble B7-H2 molecule is acting as an agonist or an antagonist for ICOS in this system, the results do support a role for B7-H2–ICOS interactions in the regulation of T-cell homeostasis.

B7-DC/PD-L2 Costimulatory Molecule

PD-L2, also called B7-DC, is another recently identified member of the B7 family. It shares approximately 41% amino acid identity with PD-L1. Human and mouse PD-L2 shares approximately 72% amino acid identity. B7-H1 and PD-L2 are both ligands for PD-1, a member of the CD28/CTLA-4 family. PD-L2, like PD-L1, is widely expressed on normal tissues (e.g., liver, lung, pancreas, heart), but not on resting peripheral blood cells. Expression on

monocytes, macrophages, and DCs is upregulated with IFN- γ stimulation. PD-L2 does not bind to CD28, CTLA-4, or ICOS^{3,73} but has been shown to bind to PD-1 found on activated lymphocytes and has potent costimulatory property for naive T cells.⁷³ B7-DC costimulates T-cell proliferation more efficiently than CD80 and induces a distinct pattern of lymphokine secretion. In particular, B7-DC strongly costimulates IFN- γ but not IL-4 or IL-10 production from naive T cells. These properties of B7-DC may account for some of the unique activity of DCs, such as their ability to initiate potent T-helper-1 (Th1) responses.⁷³ It has been shown that B7-DC fusion protein costimulated higher levels of T-cell proliferation and IFN- γ expression than CD80 costimulation.

B7-H3 Costimulatory Molecule

B7-H3 is the recently identified member of the B7 family of immune costimulatory molecules. It is a type I Ig superfamily transmembrane glycoprotein that shares approximately 24% sequence identity with CD80, 26% with CD86, 28% with B7-H1, 29% with PD-L2, and 29% with B7-H2. Human and mouse B7-H3 shares 88% amino acid identity.³ There are two isoforms of the human B7-H3 protein, which differ in the N-terminal sequence and, most importantly, differ in the number of extracellular Ig domains. It appears that the isoform with four, rather than two, Ig domains is most widely expressed in human tissues and has been designated as B7-H3b. This form is apparently the result of a gene duplication event that occurred in humans. The mouse homologue contains two extracellular Ig domains, as is usually observed in B7 family members.⁸⁸

B7-H3 mRNA is not detectable in peripheral blood mononuclear cells, although it is found in various normal tissues such as the heart, kidney, testis, and colon, and in several tumor cell lines. However, the expression on DCs, monocytes, macrophages, and B and T cells can be induced and upregulated by LPS, cytokines (GM-CSF, IFN- γ) exposure, or a PMA/ionomycin combination.^{12,88} LPS treatment stimulates B7-H3 expression on murine macrophages, DCs, and B cells.⁸⁹ Soluble B7-H3 protein binds a putative counter-receptor on activated T cells that is distinct from CD28, CTLA-4, ICOS, and PD-1. B7-H3 costimulates proliferation of both CD4+ and CD8+ T cells, enhances the induction of cytotoxic T cells, and selectively stimulates IFN- γ production in the presence of TCR signaling.⁹⁰ B7-H3 does not bind CD28, CTLA-4, ICOS, or PD-1. A recombinant B7-H3/Ig fusion protein can bind activated T cells. This suggests that a novel putative counter-receptor is present on activated T cells. This putative receptor has not been identified as of yet. Data obtained with the recombinant fusion protein demonstrates that B7-H3 can mediate T-cell proliferation and IFN- γ production. Collectively, the data suggest that B7-H3 plays a regulatory role after initial T-cell priming.³

Understanding of the function of the B7 family member B7-H3 is at an earlier stage. Dong and colleagues⁷⁸ and Chen and Yi⁹¹ review data that support stimulatory and inhibitory functions for B7-H3, discuss recent work that has identified TLT-2 (triggering receptor expressed on myeloid cells-like transcript 2) as a receptor for B7-H3, and entertain the idea that additional receptor-mediated inhibitory responses for B7-H3 may exist. There is consensus that B7-H4 functions as a coinhibitory molecule, and Dong,⁷⁸ Chen,⁹¹ and Allison⁷⁹ and their colleagues review the inhibitory functions of B7-H4 (B7x/B7S1). Yi and Chen⁹¹ discuss new data that B7-H4 serves as an important negative regulator of innate immunity through inhibition of neutrophil growth. Functional studies of B7S3 and BTNL2 are limited but indicate that both can inhibit T-cell activation. Mutations in the human BTNL2 gene have been identified, which are linked with sarcoidosis, and polymorphisms strongly associated with several human autoimmune diseases,⁹² suggesting that BTNL2 may serve to regulate tolerance or inflammation.

B7-H4 or B7x/B7S1 Costimulatory Molecule

B7-H4 is the newest member of the B7 family and is a negative regulator of T-cell responses. Mouse and human B7-H4 share 87% of amino acid identity, suggesting that there is an important evolutionary conserved function for this molecule. Human and mouse B7-H4 mRNAs are expressed broadly in both lymphoid and nonlymphoid organs. In mice, B7-H4 is expressed constitutively on B220+ B cells and can be induced in peritoneal macrophages and DCs. B7-H4Ig binds to a receptor on activated but not naive T cells. The receptor is distinct from CTLA-4, ICOS, and PD-1, and the receptor for B7-H3. Initially, it was implicated that B7-H4 is a ligand for BTLA, but later it was found that B7-H4Ig binds to wild type but not BTLA^{-/-} cells.¹² Therefore, the receptor for the B7-H4 remains to be identified, and hence, B7-H4 is also an orphan ligand such as B7-H3. B7-H4 has been shown to negatively regulate T-cell activation. Its expression in nonlymphoid tissues suggests that it might mediate tolerance at the tissue level. In a mouse ovarian tumor model, macrophages treated with oligonucleotides to block B7-H4 expression reduced the growth of tumors. This blockade of B7-H4 enhances the activity of tumor-infiltrating T cells, thereby shifting the balance of immunity and immune evasion in favor of tumor destruction.^{93,94} Hence, inhibiting the function of B7-H4 may be an effective strategy to boost antitumor immunity.

Recently, molecular homologues of CD28 and CTLA-4 receptors have been identified. ICOS is a CD28-like costimulatory receptor with a unique B7-like ligand such as B7RP-1, B7-H2, BH-2, and GL-50. PD-1 is an inhibitory receptor, with two B7-like ligands such as B7-H2 and B7DC. Additional members of B7 and CD28 gene families have been proposed. Integration of signals through this family of costimulatory and inhibitory receptors and their ligands is critical for activation of immune responses and tolerance.

CD28 Molecule

CD28 is the most effective and well-characterized T-cell costimulatory receptor discovered to date. The genes encoding CD28 costimulatory receptor are found on chromosome 1 in mice and chromosome 2 in humans. Mouse and human CD28 molecules share approximately 68% amino acid identity. It is a transmembrane cell surface glycoprotein belonging to Ig superfamily. It has a single Ig variable-like extracellular domain in addition to its transmembrane and cytoplasmic domains. CD28 is a 44-kDa, 212-amino acid, disulfide-linked homodimer receptor that is glycosylated at five different sites. Almost all murine T cells express CD28 and bind to the costimulatory ligands CD80 and CD86 to provide a costimulatory signal to T cells.⁵² It is expressed on 95% of the resting CD4+ T cells and 50% of the resting CD8+ T cells in human peripheral blood. After MHC–peptide–TCR complex formation, the ligation of B7 to CD28 provides a critical costimulatory signal to the T cell, without which T cell would become either apoptotic or anergic.³⁹ CD28 ligation can alter the threshold level of TCR ligation required for activation, reduce the time needed to stimulate naive cells, and enhance the magnitude of the T-cell response.⁹⁵ CD28 ligation also induces the antiapoptotic gene BCL-xL; increases cytokine secretion, particularly IL-2; enhances cell adhesion; facilitates reorganization of the T-cell plasma membrane upon binding to an APC; prevents anergy induction; and supports germinal center formation.^{96–98} Generally, CD28 engagement does not have a physiological effect in the absence of TCR signaling. The importance of the role of CD28-mediated costimulation has been demonstrated in a variety of model systems *in vitro* and *in vivo*.^{99–102} Most importantly, in CD28 $-/-$ mice, there was normal development of T cells. These mice showed profound, but selective defects in their immune responses. A putative role for CD28 in the differential regulation of Th1/Th2 CD4+ T-cell subsets was first suggested by studies using clones. Blocking CD28–B7 interactions greatly reduced IL-2 production and proliferation in Th1 cells, whereas Th2 cells remained unaffected.¹⁰³ After ligation, CD28 cell surface expression is downregulated, whereas that of CTLA-4 is upregulated. B7–CD28 interactions are also important in promoting T-cell tolerance because they have a critical role in the homeostasis of CD4+ CD25+ Tregs, which play an essential role in regulating self-tolerance. Hence, CD28 costimulation is necessary for the initiation of most T-cell responses, and blockade of CD28 signaling results in ineffective T-cell activation. This has therapeutic implications in that blockade of CD28 costimulation can be profoundly immunosuppressive, which prevents induction of pathogenic T-cell responses in autoimmune disease models and allows for prolonged acceptance of allograft in models of organ transplantation.

CTLA-4 Molecule

CTLA-4 (CD152) is also a member of the Ig gene superfamily present on T cells. The genes encoding CTLA-4 are closely linked on human

chromosome 2 and mouse chromosome 1. Human and mouse CTLA-4 share approximately 76% amino acid identity.¹⁰⁴ Although CD28 and CTLA-4 share approximately 30% amino acid identity, there has been no evidence that they can exist as a heterodimer. There is only limited conservation between the cytoplasmic domains of the two molecules. Where the CD28 receptor has at least five known N-linked glycosylation sites, CTLA-4 appears to have only one N-linked glycosylation site.^{105,106} CTLA-4 can be found on the surface of both CD4+ and CD8+ T cells within 24 h after activation but has not been detected by monoclonal antibodies (mAbs) in naive T cells.^{107,108} In contrast to the stimulatory effects of CD28 ligation, CTLA-4 acts as an inhibitory receptor that is vital for down-modulation of the immune response. CTLA-4 could inhibit T-cell responses first by competing with CD28 binding to CD28 ligands because CTLA-4 has a markedly higher affinity for shared ligands CD80 and CD86 compared with CD28 (K_d 0.2–0.4 μm versus 4.0 μm) and a 40–100-fold higher avidity.^{109,110} Second, CTLA-4 has a unique expression pattern. Unlike CD28, CTLA-4 is not expressed constitutively on the cell surface of naive T cells.¹¹¹ It is quickly upregulated upon TCR–CD28 engagement, with the biological effects of CTLA-4 cross-linking often being observed before the detection of protein. This appears to be because the majority of CTLA-4 is sequestered inside the cell, it traffics to the site of T cell–APC interaction and is quickly endocytosed into clathrin-coated pits.^{112,113} The distinct affinities of CD28 and CTLA-4 for B7 ligands have led to the proposal of a model in which B7–CD28 and B7–CTLA-4 interactions predominate at distinct stages of an immune response. When B7 ligands are expressed at low levels on resting T cell, B7 molecules may preferentially be engaged by a high-affinity inhibitory receptor CTLA-4. When B7 ligands are upregulated and expressed at higher levels, the predominant interactions could be between CD28 and B7, enhancing T-cell proliferation and differentiation. Once CTLA-4 expression is upregulated following T-cell activation, the inhibitory B7–CTLA-4 interaction could once again predominate, leading to termination of T-cell activation.¹¹⁴

ICOS MOLECULE

ICOS is a costimulatory receptor homologous to CD28 and CTLA-4 costimulatory receptors found on activated, but not resting T cells. Mouse ICOS is a 47–57-kDa, disulfide-linked, N-glycosylated homodimer, whereas human ICOS is a 55–60-kDa, disulfide-linked, glycosylated homodimer found on activated T cells. ICOS gene is closely linked to the genes for CTLA-4 and CD28 on human chromosomes 2q23 and mouse chromosome 1. ICOS shares approximately 30–40% sequence similarity with CD28 and CTLA-4. The ICOS gene maps to the CD28/CTLA-4 locus, suggesting that these arose by gene duplication.^{115,116} ICOS contains several conserved

motifs found in CD28, including the extracellular Ig-V-like domain and the YXXM motif in the cytoplasmic tail.^{97,117} However, ICOS does not have a conserved MYPPPY motif, which is necessary for CD28 and CTLA-4 binding to CD80/CD86 ligands.²⁴ ICOS is not constitutively expressed on naive T cells but is induced on CD4+ and CD8+ T cells following cell activation.^{115–117} Triggering of ICOS significantly costimulates the proliferation of T cells but fails to substitute for CD28 ligation in the induction of IL-2 secretion. ICOS costimulation is equivalent to that mediated by CD28 for the production of the cytokines IL-4, IL-5, IFN- γ , and TNF- α , but shows greater induction of IL-10. ICOS costimulation also enhances CD40 ligand expression that leads to an increased polyclonal Ab secretion in T- and B-cell cocultures.¹¹⁸ This indicates that the ICOS:ICOS-L pathway provides key positive signals that promote T-cell activation, differentiation, and effector responses and T-cell-dependent B-cell responses. Both CD28 and ICOS signaling upregulate Th1 as well as Th2 cytokines;^{117,119} however, ICOS does not upregulate IL-2 production. Thus, ICOS stimulates T-cell effector functions but not T-cell expansion. The role of ICOS in stimulating IL-10 production may contribute to its role in regulating Treg, T-cell tolerance, and autoimmunity.^{120,121} The function of ICOS during *in vivo* immune responses appears to depend on timing/stage of immune response as well as microenvironment because ICOS-L can be expressed on endothelial and epithelial cells as well as on professional APCs. Mutations in the ICOS gene result in impaired B-cell responses in both mice and humans.^{118,122,123} Mice deficient in ICOS are resistant to collagen-induced arthritis,¹²⁴ and antibody blockade of ICOS-L also reduced disease severity in this model. This is consistent with the requirement of ICOS/ICOS-L signaling for productive B-cell responses, as collagen-induced arthritis is dependent on the development of pathogenic autoantibodies to collagen. Similarly, blockade of ICOS-L in NZB/NZW F1 mice, which spontaneously develop a lupus-like autoimmune syndrome characterized by autoantibodies to nuclear antigens, resulted in a delay in development of clinical signs of lupus nephritis. By contrast, ICOS-deficient mice were more susceptible to experimental autoimmune encephalomyelitis (EAE), a T-cell-mediated model of multiple sclerosis, and blockade of ICOS can either exacerbate or ameliorate EAE.¹²⁵ Blockade of ICOS in a diabetes model also augmented disease, and a role for ICOS in maintenance of regulatory T-cell function was proposed as a mechanism for this exacerbation.¹²⁰ These preclinical studies in arthritis and lupus models suggest that blockade of the ICOS pathway may be of greatest benefit in inflammatory or autoimmune diseases with a significant pathogenic B-cell component.

The ICOS/ICOS ligand pathway has critical roles in stimulating effector T-cell responses and T-dependent B-cell responses, and regulating T-cell tolerance.⁷⁸ Although CD28 and ICOS have overlapping functions in early T-cell activation, ICOS has emerged as an important receptor in the immune system to fine-tune T-cell effector functions. In addition, ICOS is important

for generation of chemokine (CXC motif) receptor 5 (CXCR5)+ follicular helper T cells (TFH), a unique T-cell subset that regulates germinal center reactions and humoral immunity. Recent studies in ICOS $-/-$ mice indicate that ICOS can regulate IL-21 production, which in turn regulates the expansion of T-helper type-17 (Th17) cells and TFH. ICOS also plays an important role in controlling IL-10-producing Tregs and peripheral T-cell tolerance. Grimbacher and colleagues¹²⁶ review the discovery of mutations in ICOS in patients with common variable immunodeficiency. Human ICOS deficiency leads to defective IL-10 and IL-17 production, impaired germinal center-affinity maturation, and isotype class switching, which result in profound hypogammaglobulinemia.

The Programmed Death-1

PD-1 is a 50–55-kDa, type I transmembrane receptor related to CD28 and CTLA-4, but it lacks the membrane proximal cysteine that allows these molecules to homodimerize, and hence, PD-1 is monomeric in nature.⁷² PD-1 was identified in a T-cell line undergoing activation-induced cell death.^{83,127} It lacks the MYPPPY motif, a sequence critical for CTLA-4 and CD28 binding to CD80 and CD86. PD-1 is constitutively expressed on a subset of CD4–CD8 thymocytes, immature B cells, and some peripheral T cells, and its expression can be induced on T cells, B cells, monocytes, and myeloid cells following activation.^{128–131} The ligands of PD-1 are the B7 family members PD-L1 (B7-H1) and PD-L2 (B7-DC).^{71,72,132} Resting B cells, monocytes, and DCs do not express either PD-L1 or PD-L2. The overall distribution of PD-L1 and PD-L2 transcript is similar in human and murine tissues, with a high level of expression in placenta, a low expression level in the spleen, lymph nodes, and thymus, and the absence of expression in brain tissue.^{72,132} Expression of PD-L1 and PD-L2 in both lymphoid and nonlymphoid tissues suggests that the PD-L1/PD-L2 pathway may modulate immune responses in the secondary lymphoid organs as well as in peripheral sites. PD-1 coligation with the TCR complex that is induced by PD-1/PD-1 ligand interaction is also shown to inhibit proliferation and cytokine production of both CD4+ and CD8+ T cells. Costimulation with soluble anti-CD28 mAb can overcome PD-1-mediated inhibition by augmenting IL-2 production. The presence of ligands of PD-1, that is, PD-L1 and PD-L2 on parenchymal cells of the heart, lung, and kidney, suggests that the PD-1–PD-L system could provide unique negative signaling to help prevent autoimmune diseases.¹³⁰

PD-1:PD-L1/PD-L2 has critical roles in regulating T-cell activation and tolerance. PD-1 and CTLA-4 have important nonredundant inhibitory functions. CTLA-4 appears to have a more central role in the lymphoid organs, whereas PD-1 has an important role in regulating inflammatory responses in peripheral tissues. PD-1, a receptor for PD-L1 and PD-L2, functions as a negative regulator of immune responses.¹³³ PD-L1 and PD-L2 are expressed in

nonlymphoid cells and tissues in both normal and inflammatory conditions, suggesting that these ligands regulate immune responses in nonimmune peripheral sites. PD-1-deficient mice develop autoimmune disorders, including lupus-like arthritis in the C57BL/6 background and dilated cardiomyopathy in Balb/c mice,¹³⁰ which emphasizes the negative regulatory function of PD-1. Administration of antagonistic anti-PD-1 antibodies in EAE in the NOD model of type I diabetes and in a model of graft versus host disease results in accelerated and more severe disease development.^{77,134} In all cases, worsening of disease was associated with increased IFN- γ production, suggesting that the PD-1/PD-L pathway is a negative regulator of this cytokine *in vivo*. PD-1 blockade in tumor models enhances effector T-cell function and impairs metastasis in B16 melanoma and CT26 colon cancer models.¹³⁵ Thus, development of agonistic and antagonistic anti-PD-1 antibodies is an approach for the treatment of autoimmune disorders and cancer, respectively. Although antagonistic antibodies have been identified for PD-1, agonistic antibodies for coinhibitory receptors such as PD-1 and CTLA-4 have yet to be reported.

CD40 Costimulatory Molecule

CD40 is an approximately 50-kDa, type I transmembrane glycoprotein belonging to the TNFR superfamily. It is expressed on a variety of cells, including B cells, follicular DCs, activated monocytes, macrophages, endothelial cells, vascular smooth muscle cells, and several tumor cell lines.^{136–139} Recent studies have indicated that CD40–CD154 interaction can upregulate costimulatory molecules, activate APCs, and influence T-cell priming and T-cell-mediated effector functions. This interaction can activate macrophages, natural killer (NK) cells, and endothelial cells, as well as participate in the pathogenic processing of chronic inflammatory diseases, such as autoimmune diabetes, graft rejection, atherosclerosis, and cancer.^{140,141} During a normal T-cell response, CD40 on DC is engaged by CD40L, which is transiently expressed on activated CD4+ T cells. CD40 molecules are therefore cross-linked, which induces DC maturation and enables them to efficiently present Ag to T cells.¹⁴² CD40L–CD40 interaction is of great significance in induction of immune responses because mice and humans that lack CD40 or CD40L (CD154) genes have reduced Ab production and Ig class switching and are unable to mount effective responses against infectious agents. However, novel and recent data provide ample evidence that the functions of the CD40 molecule may extend well beyond their role in humoral immune response. CD40 signaling in macrophages induces the activity against an intracellular pathogen independently of IFN- γ and reactive nitrogen intermediates. In BCG-infected DCs, it was shown that CD40 stimulation not only promotes their ability to secrete IL-12 but increases the release of other inflammatory mediators such as IL-1 α , IL-1 β , and IL-6.¹⁴³ This shows that CD40 stimulation may

potentiate the development of inflammatory responses, which play a critical role in antimycobacterial immunity.

CD40L/CD154 Costimulatory Molecule

CD40 ligand (CD40L), also known as CD154, TNFSF5, TRAP, or gp39, is a 260-amino acid, type II transmembrane glycoprotein belonging to the TNF superfamily. Murine CD40L consists of a 22-amino acid cytoplasmic domain, a 24-amino acid transmembrane domain, and 214-amino acid extracellular domains bearing a single glycosylation site. CD40L is expressed predominantly on activated CD4 + T lymphocytes and some other types of cells such as NK cells, mast cells, basophils, and eosinophils.^{138,144,145} Murine CD40L shares 78% amino acid sequence identity with human CD40L. Both forms of CD40L, that is, membrane bound and soluble, induce similar effects on B cells. Although all monomeric, dimeric, and trimeric forms of soluble CD40L can bind to CD40, the soluble trimeric form of CD40L has the most potent biological activity through oligomerization of cell surface CD40, a common feature of TNFR family members. Expression of CD154 was detectable 4 h after activation, peaked between 6 and 8 h, and returned to near resting levels between 24 and 48 h. The CD154–CD40 interaction regulates the activation of APCs by upregulating the expression of other costimulatory molecules and by inducing the production of cytokines. Recent studies have suggested that CD40–CD154 interactions regulate the production of IL-12 by DCs and macrophages. IL-12 is required for the development of Th1 type of response. Further, the role of anti-CD154 Ab in the prevention of Th1-cell-mediated autoimmune diseases is also reported.¹⁴⁶ CD40L mediates a range of activities on B cells, including induction of activation-associated surface antigen, entry into cell cycle, isotype switching, Ig secretion, and memory generation. CD40–CD40L interaction also plays important roles in monocyte activation and DC maturation.

PATHWAYS IN THE TNFR/TNF FAMILY

TNFR family members can recruit TNFR-associated factor (TRAF) adapter proteins and activate the nuclear factor κ -B signaling pathway, making them fundamentally distinct from costimulators such as CD28 or ICOS. CD40 and its ligand, CD154, were the first costimulatory molecules to be identified as members of the TNFR/TNF superfamily and are crucial for the functions of B cells and DCs. CD40 signals into DCs and B cells as well as other cell types, including tumor cells. Noelle and colleagues¹⁴⁷ discuss downstream signaling pathways initiated by CD40 through TRAF proteins and the essential functions of CD4–CD40L interactions in controlling humoral, cellular, and tumor immunity.⁷⁹

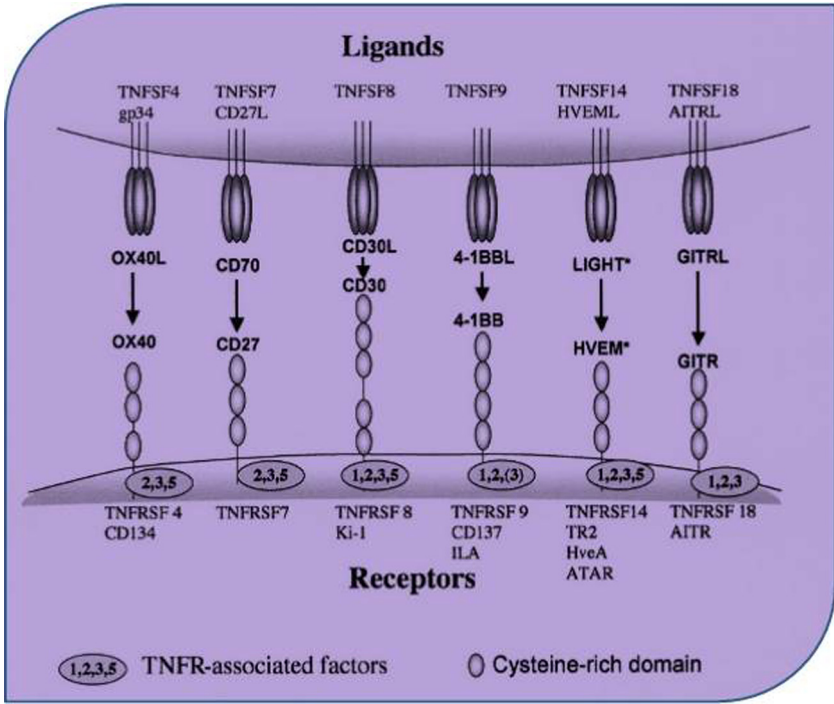


FIGURE 1.6 Summary of TNFR/TNF family ligands and their receptors. The names of receptors and ligands are indicated, as well as a brief summary of predominant expression patterns for each. The TNFR family costimulatory members such as OX40, CD27, CD30, 4-1BB, HVEM, and GITR are type I transmembrane proteins characterized by cysteine-rich motifs in their extracellular domains. The ligands for the costimulatory TNFR family members contain TRAF2-binding motifs consisting of the major conserved motif (P/S/A/T)x (Q/E)E or the minor motif PxQxxD. TRAF trimers are recruited to TNFRs upon receptor ligation. Remember TRAF3 association is observed with human but not mouse 4-1BB.

Studies of the CD27/CD70, CD30/CD30L, OX40/OX40L, 4-1BB/4-1BBL, glucocorticoid-induced TNF receptor (GITR)/GITR ligand, and HVEM/lymphotoxin-like have revealed that these molecules exhibit inducible expression, compete with herpes simplex virus glycoprotein D for HVEM, which is a receptor expressed by T lymphocytes (LIGHT) pathways, and indicate that these TNFR/TNF family members can provide important costimulatory signals. With the exception of CD27, which is constitutively expressed on naive T cells, the other TNFRs are expressed only upon T-cell activation. In addition, memory T cells and Tregs constitutively express certain family members. All the members of TNFR/TNF costimulatory molecules are depicted in Figures 1.2 and 1.6. These expression patterns have suggested that the TNFR/TNF family may be important in controlling effector and memory responses.

Recent work indicates that OX40/OX40L and 4-1BB/4-1BBL interactions have key roles in regulating the balance between effector and Treg responses. As reviewed by Croft et al.,¹⁴⁸ OX40 promotes effector T-cell expansion and survival, as well as the generation, reactivation, and maintenance of memory T cells. OX40 also blocks natural Treg activity and antagonizes the generation of inducible Tregs. Thus, OX40 can promote effector and memory T-cell responses directly by stimulating these cells and indirectly by inhibiting Tregs. OX40 has become an attractive therapeutic target; OX40 blockade can attenuate autoimmunity and inflammation, whereas stimulating OX40 can enhance antitumor immunity. As reviewed by Watts and colleagues,¹⁴⁹ the major role of 4-1BB is for survival of activated and memory T cells, with preferential effects on CD8⁺ T cells.

However, when other costimulatory signals are limiting, 4-1BB signals can cooperate with TCR-induced signals to enhance proliferation and development of effector function, perhaps by allowing T cells to survive through rounds of division. In addition, 4-1BB can exert immunosuppressive effects in some settings. Agonistic anti-4-1BB mAbs can not only expand CD8⁺ T cells but also inhibit humoral immunity and ameliorate autoimmunity. 4-1BB is expressed on Tregs, and ligation of 4-1BB on Tregs can enhance their expansion and possibly their suppressive function. 4-1BB also can induce IFN- γ and CD8⁺ T-cell-dependent suppression. Further work is needed to understand how 4-1BB controls the balance between effector and regulatory cell function. This is a clinically relevant issue, as agonistic anti-4-1BB mAbs are in clinical trials.

The CD27/CD70 pathway is also important for generation of effector and memory T-cell responses and promotes proliferation and survival of activated T cells.¹⁵⁰ The potency of costimulation by CD27/CD70 interactions is illustrated by the development of immunopathology in transgenic mice that constitutively express CD70 or CD27. These findings give impetus to studies examining the impact of CD70/CD27 interactions during chronic antigen stimulation, such as chronic virus infection and autoimmunity. HVEM sits at a pivot point, connecting the TNFR and Ig superfamilies with a complicated molecular cross talk among its receptors: LIGHT, LTA, CD160, and BTLA. LIGHT can influence T-cell activation both directly and indirectly by engaging receptors expressed on T cells and other cell types. The binding of LIGHT or LTA to HVEM delivers a costimulatory signal,¹⁵¹ whereas the binding of BTLA or CD160 to HVEM delivers a coinhibitory signal.¹⁵² Thus, HVEM is a bidirectional switch regulating T-cell activation in a costimulatory or coinhibitory fashion depending upon which ligand is engaged. LIGHT or LTA1b2 binding to lymphotoxin-bR, which is expressed by DCs and stromal cells in lymph nodes and other tissues, induces chemokine production by stromal cells resulting in the development of lymphoid structures and DC expansion. Thus, LIGHT has the capacity to serve as both a costimulatory ligand for T cells and a stromal activator to induce chemokine

production and lymphoid recruitment. For these reasons, it has become an attractive target for immunotherapy, and methods for intratumoral delivery of LIGHT are being studied. Freeman and Cai¹⁵² discuss the inhibitory consequences of HVEM interactions with CD160 and BTLA, and provide insight into how engagement of HVEM can lead to stimulatory or inhibitory outcomes. HVEM has distinct domains for binding its stimulatory ligand LIGHT and its coinhibitory ligands CD160 and BTLA. Therapies targeting the cysteine-rich domain-1 of HVEM to block BTLA and CD160 binding are being developed to enhance immune responses and vaccination.

4-1BB/4-1BBL

4-1BB (CD137) is another costimulatory receptor expressed on activated T cells (CD4+, CD8+, and NKT), activated NK cells, DCs, eosinophils, and mast cells. 4-1BB is a member of the TNFR superfamily that is inducibly expressed on T cells following stimulation through the TCR complex.^{153–155} The ligand of 4-1BB (4-1BBL or CD137L), a member of the TNF family, is expressed on activated macrophages, DCs, and B cells.^{156,157} When coupled to a strong signal through the TCR, engagement of 4-1BB can induce IL-2 production independently of CD28 ligation.¹⁵⁸ Early studies also demonstrated that ligation of 4-1BB by either cell surface 4-1BBL or specific antibodies provides a costimulatory signal particularly to CD8+ T cells, enhancing proliferation, cytokine production, and particularly survival.^{159–161} More recent data point to the role of 4-1BB engagement in augmenting rather than initiating T-cell response and in sustaining effector functions.¹⁶² *In vivo*, the first use of anti-4-1BB antibodies as an antitumor therapy was reported by Melero et al.,¹⁶³ who demonstrated the therapeutic potential of this strategy to promote rejection of P815 mastocytoma and Ag104A sarcoma tumors.¹⁶³ Further studies using either 4-1BB agonistic antibodies or gene-modified tumor cells expressing 4-1BBL confirmed their efficacy to expand tumor-reactive T cells, suppress tumor growth, and in some cases induce regression of preestablished tumor in different tumor models.^{164–166} CD8+ T cells constitute the main effector subset involved in tumor rejection following treatment by 4-1BB mAb because CD4+ T cells, NK, and NKT cells were not required for the antitumor effects.¹⁶⁷ Interestingly, several reports in rodent models have shown synergistic effects of agonistic anti-4-1BB mAb in combination with anti-TNF-related apoptosis-inducing ligand and CD40 mAbs,¹⁶⁸ with intratumoral introduction of the IL-12 gene,¹⁶⁹ or with chemotherapy or radiotherapy.^{170,171} These observations provide a guide for developing combination studies in the clinic. An unusual feature of anti-4-1BB mAb therapy is the almost paradoxical observation that, although improved immunity is seen in tumor settings, diminished pathology has been observed in autoimmune models. Fu and colleagues¹⁷² showed that same agonist antibody to 4-1BB used to promote antitumor immunity resulted in ameliorating both the incidence and

the severity of EAE. Many other studies have confirmed the beneficial effects of anti-4-1BB mAb in various autoimmune disease models, including rheumatoid arthritis, EAE, and systemic lupus erythematosus.¹⁷³ In a transplantation model, anti-4-1BB also was shown to inhibit rejection of intestinal allografts in mice.¹⁷⁴ Although a complete understanding of these differential effects is lacking, the promotion of Tregs development and activity is believed to play a role.¹⁷⁵ Additional potential explanations for immune suppression in some settings include the apparent ability to delete CD4+ T cells, retard B-cell function, and upregulate indoleamine-2,3-dioxygenase and IFN- γ .^{166,173} Together these results suggest that the possibility of both immune-potentiating and immune-suppressive effects of 4-1BB engagement should be taken into consideration and monitored in clinical/translational studies in patients. A human antihuman 4-1BB mAb (BMS-663513) is currently under investigation by Bristol-Myers Squibb in a number of phase I and II trials in patients with metastatic or locally advanced solid tumors.^{165,176}

OX40/OX40L

OX40, also known as CD134, is another costimulatory molecule of the TNFR superfamily that was identified in 1987¹⁷⁷ and that is expressed mainly on activated CD4+ but also CD8+ T cells. Its ligand OX40L is mainly expressed on professional APCs such as DCs, B cells, and macrophages, particularly after stimulation with TLR ligands or CD40L.^{178–180} OX40L also has been observed on T cells and endothelial cells.^{181,182} OX40 functions as a late costimulatory receptor and artificial engagement by OX40L stimulates proliferation, cytokine secretion, and survival of T cells, in part by increasing the expression of antiapoptotic molecules of the Bcl-2 family.¹⁸³ Several systems have shown that OX40 expression is not dependent on CD28 but that CD28 can augment the level of OX40 expression,^{183,184} suggesting that the two molecules cooperate in a sequential manner. Therefore, a model was proposed whereby OX40 signals act in a temporal manner after CD28 signals, supporting continued survival and proliferation of effector.¹⁸⁵ Interestingly, recent data have also shown that OX40 ligation may suppress the function and generation of CD4+ Foxp3+ Tregs.^{186,187} *In vivo*, OX40L transduction of tumor cells and engagement of OX40 by an agonist Ab or an OX40-Ig chimeric protein has been shown to increase tumor immunity against different murine tumor models, including B16 melanoma and hepatic colon metastases.¹⁶⁴ Interestingly, it was shown that agonistic anti-OX40 mAb given in combination with agonistic anti-4-1BB and intratumoral injection of an adenovirus expressing IL-12 showed a synergistic effect to induce long-term survival of mice bearing large chemically induced colon carcinoma.¹⁸⁸

The first phase I clinical trial using a murine antihuman OX40 mAb has been initiated in patients with advanced cancer. This trial is essentially a proof of concept, as neutralizing human antimouse antibodies are expected

to be limiting, and a humanized or fully human antibody should be pursued. A humanized OX40 agonist was recently developed that could be tested in future clinical studies.¹⁸⁹

HVEM/LIGHT

HVEM is another member of the TNFR superfamily but in contrast to others is expressed by both resting and activated T cells. The principal ligand for HVEM is the TNF superfamily member 14 (LIGHT/TNFSF14), and ligation by LIGHT acts as a positive costimulatory signal. LIGHT expression is induced on activated T cells themselves, but also by NK cells, monocytes, and DCs.¹⁶⁴ In addition to its binding to HVEM, LIGHT also can bind the LT- β receptor, which can be expressed by stromal cell populations in lymph nodes and other tissues, including the tumor microenvironment. Evidence supporting the costimulatory effect of LIGHT came from experiments, showing that blocking of LIGHT inhibits early T-cell proliferation and cytokine secretion in an allogeneic mixed lymphocyte reaction.^{190–192} However, *in vivo* experiments suggest that a major impact of LIGHT is through induction of chemokine production by stromal cells that results in recruitment of T cells and DCs, leading to the generation of lymph node-like structures. Transgenic mice expressing LIGHT driven by the CD2 promoter show an increased number of activated T cells, higher proportions of effector and memory T cells, and signs of autoimmunity.^{193,194} Transgenic mice expressing LIGHT driven by the rat insulin promoter develop autoimmune diabetes mellitus, with pancreatic islets showing a lymph node-like morphology.¹⁹⁵ LIGHT knockout mice also have been generated. These mice show defective expansion of superantigen-reactive CD8+ T cells and defective CTL generation after peptide priming *in vivo*,^{196,197} which supports a positive immunoregulatory role.

The attractive properties of LIGHT to serve as both a costimulatory ligand for T cells and a stromal cell activator to induce chemokine production and lymphoid recruitment made it attractive to consider for intratumoral application. Introduction of a LIGHT cDNA in P815 tumor cells induced regression of established tumors with the maximal therapeutic effect requiring both CD4+ and CD8+ T cells. The antitumor effect of LIGHT was shown to be CD28 independent in this model.¹⁹⁸ Introduction of LIGHT in Ag104 sarcoma cells expressing the alloantigen Ld has been shown to promote direct antigen presentation, *in vivo* chemokine production, T-cell recruitment, and tumor rejection.¹⁹⁹ LIGHT also appears to promote recruitment of NK cells into tumor sites, which may participate in a bridge to adaptive immunity.²⁰⁰ There is growing interest to bring therapies involving LIGHT forward into the clinic. As the maximal impact is expected to be at the level of altering the tumor microenvironment, strategies for intratumoral targeting would be desirable. Along these lines, an adenoviral vector encoding LIGHT injected intratumorally was shown to promote T-cell recruitment,

control of the injected tumor, and elimination of noninjected metastases in mice.²⁰¹ Another potential strategy for clinical translation would be *in vivo* electroporation directly into tumor metastases, as has recently been reported using an IL-12 cDNA in patients with melanoma.²⁰²

ICOS, GITR, and CD27/CD70

Other receptors that have been described as having positive costimulatory functions and could be attractive to consider for augmenting antitumor immunity include ICOS, a member of the B7 family, and GITR, and CD27, members of the TNFR superfamily. ICOS and GITR are mainly expressed by activated CD4+ and CD8+ T cells but constitutive expression by CD25+ CD4+ Foxp3+ Tregs has also been demonstrated.²⁰³ CD27 is expressed on naive CD4+ and CD8+ T cells, and also by NK cells. The ligands for these receptors, GITR ligand (GITRL), CD27 ligand (CD70), and ICOS ligand (B7h or LICOS), are predominantly expressed on populations of APCs (B cells, macrophages, and DCs).^{164,185} Interestingly, recent studies have confirmed the critical role for CD70 expression on activated DCs to promote optimal T-cell activation *in vivo*. Transgenic expression of CD70 on DCs enabled T-cell priming *in vivo* in the absence of adjuvant,²⁰⁴ which argues that it is perhaps one of the most critical of the costimulatory ligands for adaptive immunity. Animal models targeting these additional costimulatory pathways in the tumor context are under evaluation but earlier in their development. B7h expression on tumor cells was demonstrated to promote regression in the Sa1/N fibrosarcoma and J558 plasmocytoma models via a CD8-dependent mechanism.^{205,206} Like for B7-1 and B7-2, this strategy was ineffective when weakly immunogenic tumors were investigated.²⁰⁷ Other studies have shown that CD27/CD70 can enhance tumor rejection, with CD70 transfection enhancing both NK-dependent and T-cell-dependent mechanisms of tumor elimination.^{208–210} The administration of an anti-GITR agonistic antibody (clone DTA-1) also has been shown to induce the rejection of several murine syngeneic tumors through an IFN- γ -dependent mechanism and without obvious autoimmune manifestations.^{211,212} Anti-GITR mAb promoted the activation of effector CD4+ and CD8+ T cells, and altered the intratumoral ratio of Tregs/T effectors. More recently, observations made by Zhou et al.²¹³ have suggested that the antitumor effects of anti-GITR antibody were mainly driven by its positive costimulatory signal in effector CD4+ T cells rather than by its possible effect to abrogate the suppressive functions of Tregs.

SUMMARY

The modern concept of costimulation has shed new light on the mechanisms involved in the regulation of T-cell activation and differentiation. The identification of several new costimulatory molecules and their receptors supports the

idea of fine-tuning T-cell functions via a multitude of simultaneously or consecutively expressed T-cell molecules. The joint action of various costimulatory receptors on T cells adds not only to the complexity of this mechanism but also to the availability of ligands on different cell types and in different organs. The identification of the orphan ligands B7 homologues 3 and 4 shows that there may be more costimulatory molecules to be identified. [Table 1.3](#) categorizes the currently known costimulatory molecules that were introduced in this chapter according to functional aspects. With a growing insight into costimulatory mechanisms, a vast array of potential targets for the modulation and redirection of T-cell responses becomes available. Accordingly, many experimental studies using animal models have explored the efficacy of targeting costimulatory molecules for the inhibition, or even prevention of, the development of the different diseases.

Among these, allergen-induced sensitization and airway disease are of the most interest, because they are clearly mediated by misled T-cell responses against common environmental antigens. Most of the aforementioned molecules have been studied in murine models of allergic airway inflammation. Chapter 2 will give, therefore, an overview on these studies in murine models as well as discuss preliminary data in the human system. Finally, Chapter 3 will discuss the utilization and feasibility of costimulatory molecules for novel treatment strategies of allergic airway disease.

In sum, our understanding of costimulation has evolved significantly from the two-signal model for T-cell activation. There are both stimulatory and inhibitory second signals, and the inhibitory signals play important roles in the induction and maintenance of T-cell tolerance. In addition, we now realize that costimulatory pathways not only regulate the initial activation of naive T cells but also control effector, memory, and Tregs. These pathways may control tolerance by preventing effector T-cell responses and by promoting/maintaining Tregs. Costimulatory and coinhibitory molecules can be expressed on nonhematopoietic cells as well as professional APCs, and expression on parenchymal cells may regulate T-cell responses in specific microenvironments. Although costimulatory molecules are generally viewed through their effects on T cells, many receptor ligand interactions may lead to bidirectional signaling with important consequences for the non-T-cell partner. Thus, molecular definition of T-cell costimulatory pathways is providing insights into the exquisite regulation of T-cell activation and tolerance. Cosignaling pathways are major regulators of the critical balance between tolerance and immune responses to pathogens that are effective with an acceptable level of tissue damage. Therefore, understanding the relationships between these costimulatory ligands and their receptors in different immune effector pathways will surely help us in improving immunomodulatory therapeutics, development of improved vaccines, and avoidance of unintended tissue injury.

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Chapter 2

Concept of Reverse Costimulation and Its Role in Diseases

INTRODUCTION

The development of immune response, especially in case of naive T cells, needs at least two distinct signals for full activation to proliferate and differentiate. The first signal is provided by the specific antigen [major histocompatibility complex (MHC)–peptide complex] recognition by the T-cell receptor (TCR) and the second signal is provided by the costimulatory molecules. This requirement for the second signal explains why adaptive immune responses are stimulated by microbes but not by most self-antigens, which are not normally recognized by the innate immune system and therefore do not elicit adaptive immune responses. The second signal, which is not delivered via TCR and is not antigen specific, has been termed as costimulatory signal because, although essential, it does not by itself induce any response in T cells. However, when a T cell has its receptor ligated and receives a costimulatory signal, the T cell will proliferate and differentiate into an effector cell. Moreover, T cells that bind antigen but do not receive a costimulatory signal are thought to die or to become anergic, a state in which the cell cannot be activated even if it receives both of the signals required to activate a T cell.^{1–4} Thus, an encounter with antigen can lead to two quite distinct outcomes: proliferation and differentiation into effector cells, or inactivation or death, which outcome occurs is determined by the appropriate delivery of costimulatory signals.

Because the original proposal of a two-signal model (Figure 2.1) for lymphocyte activation, it has become firmly established that costimulatory signals mediated primarily via the conventional CD28-B7 and CD154/CD40 pathways, are necessary for full T-cell activation and play a crucial role in mediating allograft rejection, immune tolerance, infections, and so on. In several transplantation

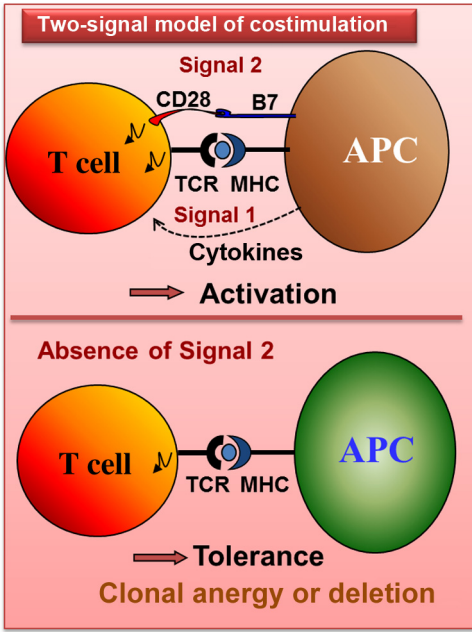


FIGURE 2.1 Two-signal model of T-cell activation. Two signals are needed for optimal activation of T cells. Signal 1 which is specific is delivered through TCR on T cells and Signal 2 which is antigen independent and nonspecific is delivered through costimulatory molecules present on APCs. Signal 2 is required for sustained T-cell proliferation, effector/memory cell generation, and prevention of anergy or apoptosis. If costimulatory signal is absent, the cell will undergo clonal anergy (unresponsiveness) or deletion. Expression of costimulatory molecules on professional APCs ensures that only pathogen-specific T cells are activated and minimize the chances of acquired immune responses being mounted against self.

models, CD28-B7 or CD154-CD40 blockade has been shown to prevent acute rejection, induce long-term allograft survival, and prevent the development and progression of chronic rejection. With the continuous discovery of new costimulatory molecules on antigen-presenting cells (APCs) and their receptors on T cells, the story is very much an evolving one. However, to date, CD80 and CD86 remain the best-defined costimulatory molecules on APCs, which provide the second signals for activation of naive T cells. The question remains: Can CD80 and CD86 deliver bidirectional costimulation?

CONCEPT OF BIDIRECTIONAL COSTIMULATION

The best-defined costimulatory molecules to date are two structurally related proteins, CD80 (B7-1) and CD86 (B7-2). These two B7 ligands, CD80 and CD86, can augment immune responses by binding to CD28 and downregulate responses by binding to CTLA-4. It has been suggested that CD86 participates in initiating immune response, whereas CD80 may be more important in sustaining or regulating immune responses (Table 2.1). The upregulation of CTLA-4 on activated T cells parallels the kinetics of CD80 expression. This property, together with the higher avidities of CD80 ligands for CTLA-4 than those of CD28, has raised the possibility that CD80/CTLA-4 interactions predominate late to terminate immune responses.⁵⁻⁷ Although both of these B7 ligands play a major role in providing costimulation to T cells by binding to CD28,

TABLE 2.1 B7 Family of Costimulatory Molecules, Their Expression Patterns, and Functions

Ligand	Expression	Receptor	Function
B71 [CD80]	DCs, activated macrophages, and B cells	CD28/ CTLA-4	Activation of naive T cells, T-cell proliferation, IL-2 production, T-cell survival/inhibition of T-cell responses
B72 [CD86]	DCs, activated macrophages, and B cells	CD28/ CTLA-4	Activation of naive T cells, T-cell proliferation, IL-2 production, T-cell survival/inhibition of T-cell responses
B7-H1 [PD-L1]	DCs, monocytes, macrophages, B cells, activated	?	Effector T-cell functions, T-cell proliferation, cytokine [IL-10, IFN- γ] production, apoptosis of CTLs, DCs, and monocytes, inhibition of T-cell responses
	T cells, endothelial cells, and tumor cells		
B7-DC [PD-L2]	DCs, monocytes, macrophages, B cells, activated	PD-1	Effector T-cell functions, T-cell proliferation, cytokine [IL-10, IFN- γ] production, apoptosis of CTLs, DCs, and monocytes, inhibition of T-cell responses, regulation of DC biology
	T cells, endothelial cells, and tumor cells		
B7-H2 [B7-H2, B7RP-1, GL50, LICOS]	DCs, monocytes, Langerhan's cells, B cells, fibroblasts, endothelial cells, etc.	ICOS	Effector T-cell function, T-cell proliferation, Ig isotype class switching, cytokine [IFN- γ , IL-10, IL-4] production
B7-H3	Induced on DCs and monocytes	? [orphan ligand]	T-cell proliferation, IFN- γ production, enhancement of CTL activity
B7-H4	Induced on T cells, B-cells, DCs, and macrophages	BTLA [?]	Inhibition of T-cell proliferation, cytokine production, cell cycle arrest

Note: DC: dendritic cells; IFN- γ : interferon- γ ; PD-1: programmed death-1; IL: interleukin; CTL: cytotoxic T lymphocyte.

which leads to their proliferation, cytokine production, and development of effector functions. The binding to CTLA-4 provides the inhibitory signals required for downregulation of the immune response.⁸ The ability of APCs to deliver the costimulatory signal to T cells by B7 molecules is very well

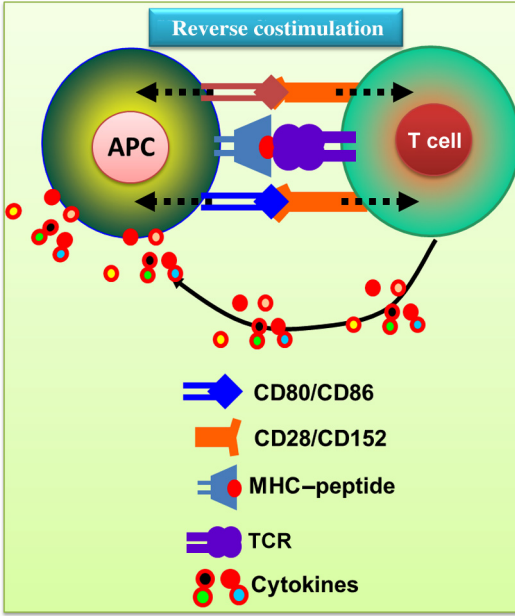


FIGURE 2.2 CD80/CD86 deliver bidirectional signaling for the activation of T cell and APC. Cognate association between T cell and APC leads to interaction between MHC and peptide with TCR and CD80/86 with CD28/CD152. This steers toward the activation and differentiation of T cell. Activated T cell secretes cytokines that ultimately help in the activation and differentiation of APC. The arrows shown in the diagram depict that there occurs as well an alternative pathway of bidirectional signaling through costimulatory molecules CD80/CD86 and CD28/CD152 that activate both T cell and APC.

established.^{9–11} However, despite the fact that both CD80 and CD86 play a major role in providing costimulation to T cells, these molecules can also serve as counter-receptors that transduce signals to APCs upon engagement with CD28/CD152 (Figure 2.2). CD80 and CD86 could also serve as counter-receptors that transduce distinct signal to the APCs upon engagement by CD28 or CTLA-4. The intracellular domains of CD80 and CD86 are quite distinct and could mediate differential signal transduction. Such signaling could influence the ability of APCs to function as effector cells.^{12–14} However, whether the engagement of CD80 and CD86 molecules with CD28 and CD152 affects the function of the APC has been poorly documented until recently.^{15,16} There was indirect evidence suggesting that CD28/CD152/B7 signaling pathways may affect B-cell responses and immunoglobulin (Ig) synthesis.^{14–16} Studies done with CD80^{-/-} and CD86^{-/-} mice do indicate the role of CD80 and CD86 in delivering bidirectional costimulation. Mice lacking CD80 and CD86 were found to be profoundly deficient in their ability to generate *in vivo* germinal center formation, Ig class switching, memory formation, and affinity maturation through somatic hypermutation.^{15–18} Thus, a role of CD80 and CD86 in the activation of B cells is plausible and needs to be systematically addressed.

However, for the first time Suvas et al.¹⁹ from our lab gave a concept of bidirectional costimulation and demonstrated that costimulation through CD80 specifically inhibits the proliferation and IgG secretion by lipopolysaccharide (LPS)-stimulated B cells and B-cell lymphoma by upregulating the expression

of proapoptotic molecules caspase-3, caspase-8, Fas, FasL, Bak, and Bax, and downregulating the levels of antiapoptotic molecule Bcl-x(L).¹⁹ In contrast, costimulation through CD86 augmented the level of antiapoptotic molecules Bcl-w and Bcl-x(L), and decreased the levels of caspase-8. Thus, the costimulatory signals not only influence the activation of T cells, but can also affect the activity of B cells by reverse costimulation. CD80 has been shown to be involved in bidirectional costimulation. Signaling through anti-CD80 Ab in LPS-activated B cells declined the proliferation and production of IgG1 and IgG2a antibodies (Abs). Signaling through CD80 molecule retarded the growth of B cells and mainly augmented the levels of proapoptotic molecules, that is, caspase-3, caspase-8, Fas, FasL, Bak, and Bax, and downregulated the expression of Bcl-x(L) and Bfl-1, which suggests that CD80 signaling induces apoptosis by upregulating proapoptotic molecules and therefore restricting their proliferation.¹⁹ These findings clearly establish that CD80 signaling not only costimulates T cells on ligation with CD28/CD152 but can also influence B cells through bidirectional costimulation. Signaling through CD86 molecule has been shown to play an important role in the bidirectional costimulation of B cells. Cross-linking of CD86 on LPS-activated B cells not only enhanced the proliferation and production of IgG1 and IgG2a Abs but also increased the expression of antiapoptotic molecules Bcl-w and Bcl-x(L) and downregulated the expression of proapoptotic molecules such as caspase-8, thereby promoting their survival by enhancing the expression of antiapoptotic proteins.¹⁹

After the report published by Suvas et al.,¹⁹ many groups have endorsed the concept of bidirectional costimulation. Schmidt et al.²⁰ has shown that targeting death receptor-6 on B cells can enhance B-cell expansion, survival, and humoral responses. Consistent with this, increased nuclear levels and activity of nuclear factor kappa-B (NF- κ B) transcription factor, c-Rel, and elevated Bcl-x(L) expression were observed.²¹ Nguyen et al. have shown that cross-linking B7-DC with the monoclonal antibody (mAb) directly potentiates dendritic cell (DC) function by enhancing antigen uptake and DC presentation of MHC-peptide complexes, promoting DC survival, and increasing secretion of interleukin (IL)-12p70.^{17,20} Radhakrishnan et al.²¹ have demonstrated that B7-DC cross-linking restores antigen uptake and augments APC function by matured DCs. Lumsden et al. have shown that CD80/86 and CD40 are required on B cells for T-dependent Ab responses.¹⁷ Mukherjee et al. have shown that CD80 and CD86 expression on mouse CD4 + T lymphocytes enhance their cell cycle progression and survival.²² Grohmann et al. have shown that upon recognition of CTLA-4 on T cells by B7 expressed on APCs, both molecules could become activated, leading to changes in the functional state not only of T cell but also of the APCs.²³ NF- κ B, a family of transcription factors, consists of five major subunits, namely, NF- κ B (P105/P50), NF- κ B2 (P100/P52), Rel-A (P65), Rel-B, and C-Rel, that form various homo- and heterodimeric complexes.^{24,25} CD86 cross-linking on IL-4 and CD40L-activated B cells increased the rate and

level of IgG1 transcription, nuclear localization of NF- κ B p50 subunit, and phosphorylation of Rel-A (p65) and I κ B α . It also increased octamer-binding transcription factor 2 (oct-2) expression and binding to the 3'Ig H enhancer. These results were not observed in CD86 $-/-$ B cells.²⁵ All these results clearly establish that CD86 does not only costimulate T cells but can also deliver reverse signals into B cells and modulate their activity. NF- κ B remains sequestered in the cytoplasm of the resting cell in a complex with I κ B proteins. Upon activation of the classical NF- κ B pathway, I κ B proteins are phosphorylated, polyubiquitylated, and degraded within the proteasome, which causes the release of NF- κ B dimers that translocate to the nucleus and regulate gene activity. Recent reports indicate that CD86 stimulation on a B cell activates the classical NF- κ B pathway, which leads to a protein kinase C-3 (PKC-3)-independent phosphorylation and degradation of I κ B α and subsequent nuclear localization of P50/P65. CD86 is also reported to increase the phosphorylation of P65 in a PKC-dependent manner.²⁶ It has been reported that cross-linking CD40 on B cells rapidly activates NF- κ B.²⁷

REVERSE COSTIMULATION OF APCs

Many studies since then have shown the significance of CD80 and CD86 in influencing the activity of B cells, stem cells, DCs, and macrophages. Recently, many reports have appeared in literature highlighting the role of bidirectional signaling (reverse costimulation) through different costimulatory molecules into APCs and in particular DCs. Reverse signaling through CD80, CD86, B7-DC, CD40L, and OX40 has been shown to regulate the immune functions of DCs. It has been shown that cross-linking of B7-DC, a member of B7 family on DC, modulates its biology by increasing survival, antigen presentation, IL-12p70 production, and migration to draining lymph nodes.^{19–21,28} This indicates that B7-DC can deliver directly to DCs. As we know, DCs has two states: the immature DCs that perform sentinel functions, sampling for antigen, and danger signals, and the mature DCs that perform enhanced APC functions but are no longer capable of acquiring the antigen. However, recently a new DC activation phenotype has been achieved when mature DCs were cross-linked with B7-DC. This B7-DC cross-linking restored the antigen-acquiring function of immature DCs and augmented APC function of matured DCs.²² During a normal T-cell response, CD40 on DC is engaged by CD40L, which is transiently expressed on activated CD4 $+$ Th cells. CD40 molecules are thereby cross-linked, which induce DC maturation and enable them to efficiently present antigen to T cells. Agonistic ligation of anti-CD40 Ab effectively helps the Th cells in DC maturation. Hence, the ligation of CD40 induces maturation of DCs and could be a useful target for vaccines.²⁹ Ligation of CD40 on DCs promotes upregulation of the costimulatory molecules CD80 and CD86,³⁰ IL-12 secretion, and release of chemokines such as IL-8, macrophage inflammatory protein (MIP)-1 α , and MIP-1 β .^{31,32} DCs also

up-regulate OX40L, another member of the tumor necrosis factor (TNF) super family in response to CD40 ligation. OX40L binding to OX40 costimulatory molecule enhances production of cytokines such as TNF- α , IL-12, IL-1, and IL-6 by DCs, and also induces the expression of costimulatory molecules.^{32,33}

Very recently, it has been shown that CD28 induces immunostimulatory signals in DC via CD80 and CD86. Logue and Sha have indicated that CD28-B7 bidirectional signaling is a two-way street to activation of DC.³⁴ Bidirectional signaling along the B7-CTLA-4 coreceptor pathway enables reciprocal stimulation of T cells and DCs. Binding of CD28 to CD80/CD86 ligands leads to enhanced production of IL-6 by DCs.²³ However, binding of CD152 to CD80/CD86 augments the secretion of interferon- γ (IFN- γ), which in turn upregulates the expression of the enzyme indolamine 2,3-dioxygenase (IDO) in DCs, resulting in tryptophan catabolism and suppression of T-cell proliferation.²³ Further, ligation of CD86 leads to antiapoptotic signals.³⁵ It has been shown that CD86 is recruited to lipid rafts upon T-cell–DC interaction. In this bidirectional interaction between CD28 on a T cell and CD86 on the DC, naive CD4 + T cells receive their costimulatory signal and the DCs appear to respond by reorganizing their CD86 to lipid rafts.^{36,37} Although the functional significance of CD86 response remains to be determined, the fact that the ligation and recruitment of CD86 to lipid rafts are required for the appearance of phosphorylated serines in rafts supports the idea that this is a mutually beneficial interaction that modulates not only T cell but also DC signaling. These findings on DCs also further confirm the interaction between CD80/CD86 and CD28/CD152 that delivers a bidirectional costimulation, which activates both T cells and DCs.

ROLE OF CD80 AND CD86 IN THE COSTIMULATION OF B CELLS

Although many costimulatory molecules, such as CD80, CD86, CD40, heat-stable antigen, intercellular adhesion molecule, vascular cell adhesion molecule, B7-DC/programmed death ligand-2 (PD-L2), B7-H1/programmed death ligand-1 (PD-L1), are known to be expressed on the surface of APCs. However, the best-defined costimulatory molecules are two structurally related proteins known as CD80 (B7-1) and CD86 (B7-2). Both CD80 and CD86 have their ligands, CD28 and CTLA-4, expressed on T cells (**Table 2.2**). It has been established recently that CD80 and CD86 play a role in the activation of APCs.¹⁹ Stimulation via CD86 in B cells can modulate their proliferation, IgG secretion and expression of proapoptotic and antiapoptotic molecules, nuclear localization of NF- κ B (p50) subunit, and phosphorylation of ν -rel reticuloendotheliosis viral oncogene homolog A (Rel-A) (p65) and I κ B α ; and increase oct-2 expression.^{18,25–27} We have investigated the impact of CD80 and CD86 on B cells.¹⁹ Signaling in LPS-stimulated B cells was delivered through CD80 and CD86 molecules by their respective Abs. Exciting features observed during the study were that cross-linking of CD86 enhanced the proliferation and

TABLE 2.2 Characteristics of CD80 and CD86 Costimulatory Molecules

Characteristic	CD80 (B7-1)	CD86 (B7-2)
Expression	B cell, macrophage, DC	B cell, macrophage, DC
Kinetics of expression	Induced slowly and expressed for a longer duration (4–5 days)	Expression is modulated rapidly but stays for short duration (48 h)
Ligand	CD28, CD152	CD28, CD152
CD152 association	Faster association and slower dissociation rates	Faster association and dissociation rates
CD152 binding affinity	high	low
Sequence Identity	25%	25%
Structure	Dimer	Monomer
Location	Transmembrane	Transmembrane
Antibody blockade	Enhances immune response	Attenuates immune response

production of IgG1 and IgG2a Abs. In contrast, anti-CD80 Ab could reduce the proliferation and production of IgG1 and IgG2a Abs. Importantly, anti-CD80 Ab could also retard the growth of B cells and upregulate the expression of proapoptotic molecules (Figure 2.3). The involvement of Fas and FasL expression in B cells in inducing apoptosis was demonstrated by flow cytometry. In addition, the association of CD80 and CD86 molecules in the regulation of the activation of pro- and antiapoptotic molecules [Fas, FasL, FADD, FAP, FAF, TRAIL (Fas2L), TNFR (p55), TRADD, RIP, Bcl-w, Bfl-1, Bcl-x(L), Bak, Bax, Bcl-2, Bad, caspase-3, caspase-8] was examined by ribonuclease protection assay in B cells.²⁶ We observed that signaling through CD80 molecule mainly augmented the levels of proapoptotic molecules, that is, caspase-3, caspase-8, Fas, FasL, Bak, and Bax, and downregulated the expression of Bcl-x(L) and Bfl-1 (Figures 2.4). This suggests that CD80 signaling induces apoptosis via mechanisms involving proapoptotic molecules rendering the cells more vulnerable to apoptosis and therefore restricting their proliferation. In contrast, signaling through CD86 increased the expression of antiapoptotic molecules Bcl-w and Bcl-x(L) and downregulated the expression of caspase-8 (Figures. 2.4). Thus, there may be a possibility that ligation of CD86 on B cells may promote their survival by increasing the expression of antiapoptotic proteins. It is worth mentioning here that we demonstrated the role of CD80 and CD86 not only in

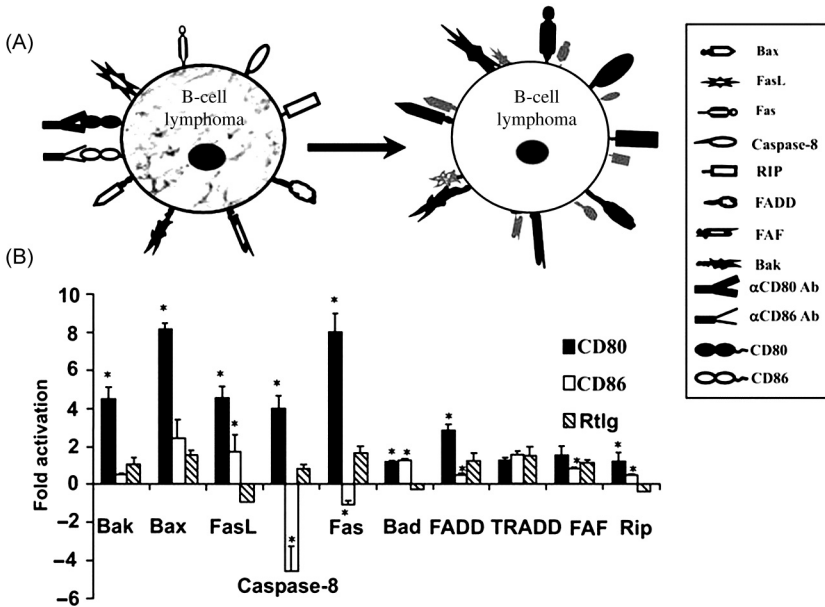


FIGURE 2.3 Modulation of the expression of proapoptotic molecules by CD80 and CD86 signaling. (A) The normal expression of proapoptotic molecules (open symbols) on B-cell lymphoma. Signaling through CD80 significantly upregulates Fas, FasL, caspase-8, Bak, Bax, and FADD (solid symbols) and triggering through CD86 downregulates caspase-8 (gray color symbols). (B) The WEHI-279 cells were cultured with anti-CD80 Ab, anti-CD86 Ab, and Rtlg, and cross-linked with anti-Rtlg Ab. Total RNA was extracted from the cells by the guanidinium isothiocyanate-phenol-chloroform method using Trizol reagent. The expression was determined using the RiboQuant multiprobe RNase protection assay system. The figure depicts the summary of results of Suvas et al.¹⁹ The resulting resolved bands were imaged using a phosphor imager. The normalized quantity for each band was obtained by dividing with L32 housekeeping gene control. The old change was calculated by dividing the value of normalized quantity of the experimental samples with that of cells cultured with medium alone and is expressed as mean ± SD obtained from two experiments. “***” indicates that $P < 0.001$ employing Student’s *t* test.

case of B cells isolated from mouse splenocytes but also for B-cell lymphomas.¹⁹ Because B-cell lymphomas (WEHI-279 and A20) are 100% B cells, this finding therefore rules out any involvement of cytokines secreted by contaminating T cells in activating B cells. After our report,¹⁹ many reports in literature started accumulating which demonstrate the role of CD80 and CD86 in bidirectional signaling of APCs.^{25–28,34,38–44} CD86 cross-linking studies on CD40 ligand and IL-4-activated B cells have shown that there is increase in the rate and level of IgG1 transcription, nuclear localization of NF-κB p50 subunit, and phosphorylation of Rel-A (p65) and IκBα. It also increased oct-2 expression and binding to the 3'Ig H enhancer. These effects do not occur in CD86^{-/-} B cells.^{25,26,41,42} It has also been demonstrated that increased expression of Bcl-x(L)

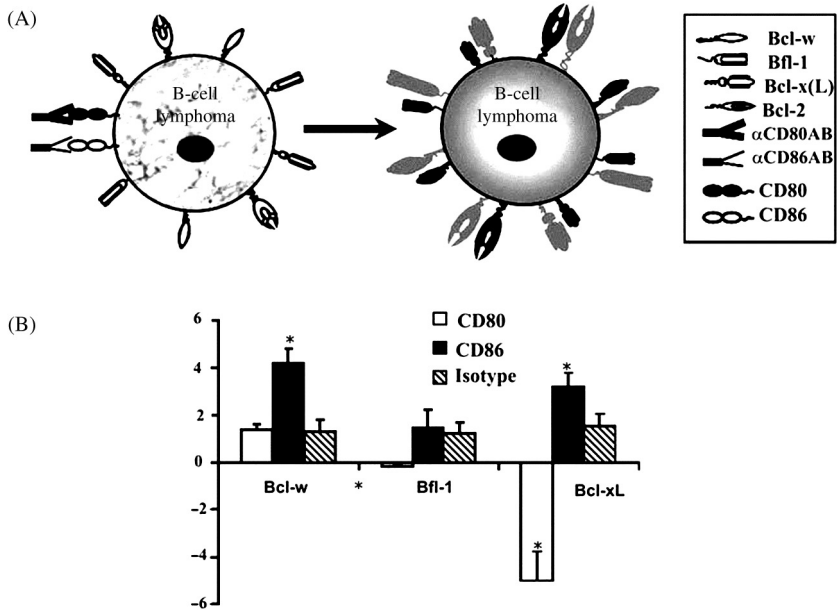


FIGURE 2.4 Modulation of the expression of antiapoptotic molecules by CD80 and CD86 signaling. (A) The normal expression of antiapoptotic molecules (open symbols) on B-cell lymphoma. Signaling through CD80 significantly downregulates Bcl-x(L) and Bfl-1 (solid symbols) and triggering through CD86 upregulates Bcl-w and Bcl-x(L) (gray color symbols). (B) The WEHI-279 cells were cultured for 72 h with anti-CD80 Ab, anti-CD86 Ab, and Rtlg, and cross-linked with anti-Rtlg Ab. Total RNA was extracted from the cells by the guanidinium isothiocyanate-phenol-chloroform method using Trizol reagent. The expression was determined using the RiboQuant multiprobe RNase protection assay system. The figure depicts the summary of results of Suvas et al.¹⁹ The resulting resolved bands were imaged using a phosphor imager. The normalized quantity for each band was obtained by dividing with L32 housekeeping gene control. The -fold change was calculated by dividing the value of normalized quantity of the experimental samples with that of cells cultured with medium alone and are expressed as mean ± SD obtained from two experiments. “*” indicates that $P < 0.001$ employing Student’s *t* test.

but not Bcl-2 can prevent the apoptosis in B-cell lymphoma. Moreover, signaling through CD40 upregulated Bcl-x(L) and Bfl-1, and protected B-cell lymphoma from apoptosis.^{45,46} Thus, this finding accurately establishes that CD80 and CD86 not only costimulate T cells on ligation with CD28/CD152 but can also influence B cells through bidirectional costimulation, consequently not only modulating the activity of T cells but also of B cells.

ROLE OF BIDIRECTIONAL COSTIMULATION IN THE ACTIVATION OF DCS

Many studies have now established the role of reverse costimulation through CD80/CD86 and B7-DC costimulatory molecules in the modulation of

activity of DCs. It has been shown that CD86 is recruited to lipid rafts upon T-cell–DC interaction. In this bidirectional interaction between CD28 on a T cell and CD86 on the DC, naive CD4+ T cells receive their costimulatory signal and the DCs appear to respond by reorganizing their CD86 to lipid rafts.^{36,37} Although the functional significance of CD86 response remains to be determined, the fact that the ligation and recruitment of CD86 to lipid rafts are required for the appearance of phosphorylated serines in rafts supports the idea that this is a mutually beneficial interaction that modulates not only T cell but also DC signaling. Cross-linking of B7-DC, a member of B7 family on DCs, modulates the DC biology by increasing their survival, antigen presentation, IL-12p70 production, and migration to lymph nodes.^{28,43,44} These findings on DCs further confirm the interaction between CD80/CD86 and CD28/CD152 that delivers a bidirectional costimulation, which activates both T cells and APCs. Logue and Sha have indicated that CD28-B7 bidirectional signaling is a two-way street to activation of DC. Engagement of B7 molecules on DCs by the CTLA-4–immunoglobulin (CTLA-4–Ig) fusion protein or by CTLA-4 expressed on regulatory T cells induces expression of the T-cell–tolerogenic enzyme IDO in DCs, which catabolizes tryptophan and leads to inhibition of T-cell function.^{23,47} In an unexpected twist to those studies, in this issue of *Nature Immunology*, Orabona et al.³⁸ have revisited those experiments using CD28-Ig to engage B7 molecules on DCs and report notably different effector outcomes for treated DCs. CD28-Ig treatment, in contrast to CTLA-4–Ig treatment, leads to production of IL-6 from DCs that squelches the previously described IFN- γ -driven immunosuppressive pathway of tryptophan catabolism in DCs. Engagement of B7 molecules on DCs by CD28-Ig thus results in activated DCs, whereas engagement of B7 molecules on DCs by CTLA-4–Ig results in tolerogenic DCs. Induction of tolerogenic IDO in DCs by CTLA-4–Ig treatment involves multiple steps. Engagement of B7 molecules by CTLA-4–Ig leads to the activation of several intracellular signaling pathways in DCs: the transcription factor NF- κ B, the kinase p38 mitogen-activated protein kinase (MAPK), and the transcriptional activator STAT1. These activated DCs secrete IFN- γ , which, in an autocrine or paracrine way, promotes the subsequent expression of IDO by DCs.²³ IFN- γ does not seem to be absolutely required for IDO expression, as IFN- γ receptor-deficient DCs can still upregulate IDO in response to CTLA-4–Ig treatment. IDO expression by DCs promotes tolerance by catabolizing tryptophan, leading to a local decrease in tryptophan and an increase in catabolic by-products that inhibit T-cell proliferation and may also induce apoptotic death. In contrast to results with CTLA-4–Ig treatment, Orabona et al.³⁸ report that DCs treated with CD28-Ig secrete IL-6, in addition to IFN- γ , and do not undergo upregulation of IDO expression (Figure 2.5). This failure to induce the IDO immunosuppressive pathway by CD28-Ig treatment does not seem to be due to inadequate engagement of B7 molecules on DCs, but instead depends

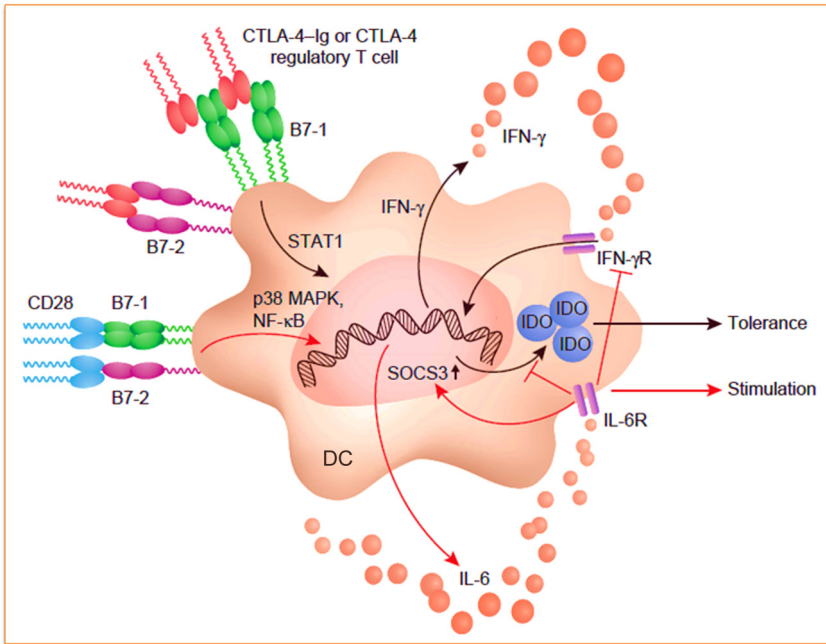


FIGURE 2.5 Distinct effector responses in DCs treated with CTLA-4-Ig and CD28-Ig. CTLA-4-Ig engagement of B7 molecules on DCs leads to activation of STAT1, p38 MAPK, and NF- κ B, which stimulates the DC to produce IFN- γ . In an autocrine or paracrine way, IFN- γ promotes upregulation of IDO, an enzyme that catabolizes tryptophan and leads to potent inhibition of T-cell responses. CD28-Ig engagement of B7 molecules on DCs, however, results in concomitant secretion of both IL-6 and IFN- γ . IL-6, in an autocrine or paracrine way, inhibits the IFN- γ -driven upregulation of IDO through downregulation of the IFN- γ receptor (IFN- γ R) and upregulation of suppressor of cytokine signaling-3 (SOCS-3). IL-6 effects are thus dominant in CD28-Ig-treated DCs and lead to DCs capable of enhanced cell-mediated immunity. IL-6R, IL-6 receptor.

critically on IL-6 secretion. Blockade of IL-6 signaling restores IDO expression in CD28-Ig-treated DCs, indicating that IL-6 effects are dominant over IFN- γ effects in CD28-Ig-treated DCs. The basis for the unique induction of IL-6 secretion in DCs treated with CD28-Ig but not those treated with CTLA-4-Ig is unclear, as both fusion proteins activate MAPK and NF- κ B pathways, albeit with different kinetics. The three different *in vivo* models used by Orabona et al.³⁸ demonstrate the potential for CD28-Ig to be used as an adjuvant. This adjuvant activity is clearly demonstrated, as CD28-Ig conditioning of DCs results in enhanced reactivity in a skin test assay using poorly immunogenic peptides and also allows a normally nonprotective DCs pulsed with hyphae to become protective in an infectious model using *Candida albicans*. This adjuvant effect of CD28-Ig is further demonstrated in a tumor model in which CD28-Ig injection allows for the eradication of a poorly immunogenic tumor through the enhancement of the antitumor

cytotoxic T lymphocyte (CTL) response and reduction of regulatory T-cell-suppressive activity. As with the *in vitro* studies of CD28-Ig, the *in vivo* adjuvant activity of CD28-Ig in all three models seems to depend critically on IL-6 signaling. These data suggest that CD28-Ig treatment could be a valuable tool to enhance immune responses during vaccination or to induce antitumor immune responses. In addition to the potential clinical implications for using CD28-Ig therapeutically, the findings by Orabona et al.³⁸ raise several intriguing questions. Does the adjuvant effect of CD28 as a fusion protein also extend to bidirectional activation of DCs by T cells expressing endogenous CD28? The original observation of CTLA-4-Ig inhibition of DCs has since been extended to endogenous regulation in studies, demonstrating that regulatory T cells, which have high constitutive CTLA-4 expression, can activate IDO expression on DCs through a CTLA-4-dependent mechanism.⁴⁷ Orabona et al.³⁸ found that IL-6 was also secreted when CD28 expressed on the surface of Jurkat cells was used to stimulate DCs, suggesting that the stimulatory effects of CD28 could also be relevant to natural cellular interactions between T cells and DCs. Although it is clear that engagement of B7 molecules leads to alterations in DC function, the most proximal signaling events elicited by CD28-Ig and CTLA-4-Ig treatment remain murky. In contrast to CD28 and CTLA-4, whose function as receptors is better understood, the cytoplasmic tails of B7 molecules have so far yielded very few insights into potential biochemical signaling mechanisms at the level of modification or association with other signaling molecules. The need to address this issue is emphasized by the demonstration in the article of Orabona et al.³⁸ that engagement of B7 molecules on DCs by CD28 and CTLA-4 fusion proteins can lead to notable functional differences in DCs.

Studies have begun to demonstrate previously unrecognized structural and biophysical differences between CD28 and CTLA-4, as well as between B7-1 and B7-2, that may underlie functional differences mediated by these costimulatory molecules.⁴⁸ In addition to the approximately 20-fold higher affinity of CTLA-4 versus CD28 for B7 molecules, it is now recognized that although both CD28 and CTLA-4 are structurally dimers, dimeric CD28 can bind only one B7 molecule, whereas CTLA-4 can bind two B7 molecules.⁴⁹ It is unclear whether these differences in valency and affinity of B7 binding underlie the functional differences observed by Orabona et al.³⁸ between CTLA-4-Ig and CD28-Ig treatment. B7-1 and B7-2 can also exert an unexpected degree of control over localization of CD28 and CTLA-4 during immunological synapse formation. Because B7-1 is a dimer, whereas B7-2 is a monomer, CTLA-4 will preferentially form extensive arrays with B7-1 but not with B7-2.^{50–52} These affinity differences between CD28 and CTLA-4 binding to B7-1 and B7-2 allow selective recruitment of CTLA-4 to the immunological synapse to be mediated principally by B7-1 and selective recruitment of CD28 to the immunological synapse to be mediated by B7-2.⁵³ In sum, the cell-extrinsic adjuvant effect of CD28-Ig on DCs

reported by Orabona et al.³⁸ provides an unexpected symmetry to the cell-intrinsic stimulatory effect of CD28 on T cells and indicates that CD28 and CTLA-4 as ligands of B7 molecules can signal DCs to initiate distinct effector responses. The potential for costimulatory molecules to exert both cell-intrinsic and cell-extrinsic regulation of effector responses is likely to also be relevant to newer members of the CD28–B7 family costimulatory molecules.⁵⁴ Cross-linking of the B7 family member PD-L2 on DCs, for example, has also been shown to activate the immune function of DCs.²⁸ Furthermore, the broader expression of more recently discovered CD28–B7 costimulatory molecules, such as the inhibitory PD-1 receptor expressed on activated T and B cells, suggests the potential for previously unknown immune regulation through bidirectional signaling. Further, it has been shown that CD28 induces immunostimulatory signals in DC via CD80 and CD86. Bidirectional signaling along the B7–CTLA-4 coreceptor pathway enables reciprocal stimulation of T cells and DCs. Binding of CD28 to CD80/CD86 ligands lead to enhanced production of IL-6 by DCs.^{38,55} However, binding of CD152 to CD80/CD86 augments the secretion of IFN- γ , which in turn up-regulates the expression of the enzyme IDO in DCs, resulting in tryptophan catabolism and suppression of T-cell proliferation.^{23,56,57}

ROLE OF REVERSE COSTIMULATION ON MACROPHAGES

We have investigated the impact of costimulation on macrophages. In this study, we have attempted to analyze the distinct role of CD80, CD86, and CD40 costimulatory molecules in the activation of macrophages and their influence on the survival of intracellular pathogens. The following five major findings have emerged from this study: (1) Costimulation through CD80, CD86, and CD40 augmented the release of nitric oxide (NO), IL-1, IL-6, and TNF- α ; (2) CD80, CD86, and CD40 worked synergistically with IFN- γ in the production of NO, IL-1, IL-6, TNF- α , and IL-12; (3) CD80 and CD86 modulated the expression of CD14, CD206, CD80, CD86, MHC-I, and MHC-II molecules; (4) triggering of CD86 and CD40 but not CD80 increased the uptake of soluble and particulate antigens; and (5) engagement of mainly CD80 reduced the survival of intracellular pathogens.⁵⁸ The enhanced secretion of NO by macrophages on costimulation by CD86 molecule will help in the eradication of intracellular infections. Augmented release of NO by macrophages is known to be beneficial in case of many intracellular infections. These results further substantiate our earlier findings¹⁹ and are now endorsed by other groups as well; the finding that signals delivered through CD80 and CD86 can affect the function of APCs. Because B7-1/B7-2 activates the induction of TNF- α as well as reactive oxygen species and bactericidal activity in macrophages, anti-B7-1/anti-B7-2 mAb treatment can be used to tailor immune responses to induce cytotoxicity against intracellular pathogens that reside inside the macrophages.⁵⁸ This approach

may be limited not only to *Mycobacterium tuberculosis* but also to other infections, including AIDS, leishmania, and malaria. This novel strategy can also be effectively exploited to develop immunotherapy using humanized Abs against B7-1 and B7-2 or CD28 fusinogenic proteins for the treatment of intracellular pathogens because this approach is based on modulating the immune system of the host rather than targeting the pathogen; hence, it diminishes the chances of emergence of drug-resistant strains of bacteria. Our results have important implications for the efforts to establish a vaccine against intracellular pathogens, such as *M. tuberculosis* and *Salmonella typhimurium*, because they suggest that vectors that induce/upregulate the expression of costimulatory ligands on APCs such as macrophages (which are also the host for *M. tuberculosis*) will help to generate a protective IFN- γ -dependent immune response, which is desired in tuberculosis.

In another study, the functional significance of costimulatory interactions between decidual macrophages (DMs) and T cells with a particular focus on B7-H1:PD-1 signaling was investigated. It was observed that human DMs suppress IFN- γ production by T cells through costimulatory B7-H1:PD-1 signaling in early pregnancy.⁵⁹ On analysis, the expression profile of B7 ligands on human DMs revealed that B7-H1 was present on DMs isolated from early but not term pregnancies. B7-H1 was not expressed on the peripheral monocytes (PMs) of pregnant women. In response to IFN- γ , B7-H1 expression was induced on PMs and was enhanced on DMs, suggesting that this cytokine might be a key factor in the control of B7-H1 expression in the decidua. The majority of decidual T cells were noted to exhibit robust expression of PD-1, whereas the expression was limited to a small subpopulation of circulating T cells. Functional assays demonstrated that DMs are able to suppress T-cell IFN- γ production via B7-H1–PD-1 interactions. This suppressive property was not observed for PMs, which lack B7-H1. B7-H1 on DMs may function as a key regulator of local IFN- γ production and therefore contribute to the development of appropriate maternal immune responses to the fetus in early pregnancy.⁵⁹

In another study, it was observed that there was early induction of suppressor of cytokine signaling-1 (SOCS-1), and the downregulation of toll-like receptors (TLRs) 7 and 9 induces tolerance in costimulated macrophages.⁶⁰ In this study, mouse macrophages were simultaneously stimulated with the TLR7 agonist, gardiquimod, and the TLR9 agonist, CpG ODN 1826, to examine the mechanism and effector functions of macrophage tolerance. Compared with individual stimulation, the costimulation of both TLRs reduced the secretion of TNF- α and IL-6 through the delayed activation of the NF- κ B pathway; notably, IL-10 remained unchanged in costimulated macrophages. This tolerance reflected the early induction of SOCS-1, according to the detection of elevated TNF- α secretion and restored NF- κ B signaling in response to the small interference RNA-mediated abrogation of SOCS-1 signaling. In addition, the restimulation of each TLRs using the same ligand significantly reduced the expression of both TLRs in

endosomes. These findings revealed that the costimulation of TLR7 and TLR9 induced macrophage tolerance via SOCS-1, and the restimulation of each receptor or both TLR7 and TLR9 downregulated TLR expression through a negative feedback mechanism that protects the host from excessive inflammatory responses. Moreover, the insufficient and impaired immune response in chronic viral infection might also reflect the repeated and simultaneous stimulation of those endosomal TLRs.⁶⁰

APPLICATION OF REVERSE COSTIMULATION IN DISEASES

Many reports on *in vivo* studies reveal the distinct role of CD80 and CD86 in diseases. We have for the first time reported that signals delivered via CD80 into B-cell lymphomas restrict their growth by modulating the expression of antiapoptotic molecules.¹⁹ Thus, providing a novel insight into the mechanism where triggering through CD80 could control the growth of lymphomas. These observations constitute a rationale for development of a therapy using anti-CD80 Ab for cancer cells that express CD80 molecule. Interestingly, after our report, clinical studies evaluated anti-CD80 Ab as a targeted therapy for lymphomas with promising results.^{61,62} Reverse costimulation has been used in CD80-positive hematologic malignancies [e.g., diffuse large B-cell non-Hodgkin's lymphoma (NHL), chronic lymphocytic leukemia, Hodgkin's disease, and multiple myeloma] and has showed promising results.^{61,62} CD80 and CD86 have been shown to have differential effects on the development of spontaneous autoimmune diabetes in nonobese diabetic mice. Ab neutralization of CD80 has been shown to exacerbate diabetes, whereas neutralization of CD86 was shown to abrogate the development of diabetes.⁶³ Differential effects of signaling by CD80 and CD86 have also been demonstrated in other autoimmune models, including lupus nephritis in the Murphy Roths large (MRL)-lpr/lpr mice and experimental autoimmune encephalomyelitis (EAE). The animals lacking CD86 showed diminished renal Ig deposition and attenuated pathology, but CD80-deficient animals developed more severe nephritis.^{64,65} Recently, a novel regulatory role for CD80-mediated intracellular signals in CD4⁺ T cells has been shown to have important implications in the relapse of EAE.²⁶ It has been established that *in vivo* blockade of CD86 costimulation could suppress maternal immune attack to the fetus.⁶⁶ Further, cross-linking of CD80 on the surface of neural stem cells *in vitro* enhances apoptosis.⁴⁰ Besides this, CD80 and CD86 have been implicated to have opposing roles in regulation of xenotransplantation rejection, where CD80 drives cell-mediated rejection (CMR) and attenuates acute vascular rejection (AVR), whereas CD86 drives AVR. Remarkably, indefinite xenograft survival can be achieved by suppressing AVR with CD86 neutralization in combination of cyclosporine A therapy, which inhibits CMR.⁶⁷ An important question arises in the studies using Abs,

whether the influence of anti-CD80 and CD86 Abs arises due to obstruction of the interaction of T cells with APCs or due to signaling delivered by Abs. It is evident from the recently published work that triggering through CD80 can *vis-à-vis* deliver signals as well.^{16,19,61,62}

Based on the pleiotropic activities of the CD28/CD152–CD80/CD86 pathway mentioned above, it can thus be exploited for the potential clinical usefulness in immune intervention in diseases such as autoimmunity, transplant rejection, and allergy, and in the elimination of tumors that evade immune surveillance.^{67–72} Hence, the interest lies in understanding the therapeutic manipulation of these costimulatory molecules. These molecules may provide a means to either promote immune responses against cancer or reduce graft rejection and autoimmunity. CD80/CD86 signaling/blockade by CTLA-4–Ig has been studied in clinical trials as treatment for severe psoriasis vulgaris, rheumatoid arthritis, multiple sclerosis, lupus nephritis, and renal transplantation.^{71,72} Importantly, the experimental data showing distinct functions of CD80 and CD86 in regulating the immune response in various disease models underscore the need to design tailor-made therapeutic strategies for humans for preventing transplant rejection and treatment of autoimmune diseases. Some of the studies done recently have established the therapeutic potential of Abs against CD80 and CD86.^{19,61,62,72} However, it is imperative to evaluate their clinical significance in other diseases as well. In the case of autoimmune diseases and allergy, it will be worth studying the role of signaling through CD80/CD86 in promoting clonal deletion of highly pathogenic auto-reactive B and T cells (Figure 2.6).

REVERSE COSTIMULATION THROUGH CD80 (B7-1)

CD80 is a costimulatory molecule known for its role in T-cell activation and also in regulating the activity of normal and malignant B cells.¹⁹ Surface CD80 is expressed transiently on activated B cells, macrophages, and DCs. Surprisingly, CD80 is downregulated on most of the cancer cells, and the loss of CD80 alone is sufficient to allow them to escape the attack of the immune system and to impart anergy and apoptosis in tumor-infiltrating T cells.⁷³ In the absence of costimulation, recognition of antigens by T cells may not cause any response, even if tumor cells express MHC molecules and tumor-specific antigens (Figure 2.7). Hence, most human malignancies that lack CD80 expression have been suggested to evade immune surveillance and therefore contribute for failure of immune recognition.⁷⁴ However, follicular lymphomas express CD80 and therefore could potentially be targeted by CD80 immunotherapies. Lack of either CD80 or MHC-I must be sufficient to allow tumor cells to evade immune response, but it has been shown in some tumors that high expression of MHC-I and absence of costimulatory molecules render them resistant to lysis by CTLs (Figure 2.7A). It has been shown that high expression of MHC-I but lack of CD80 does not

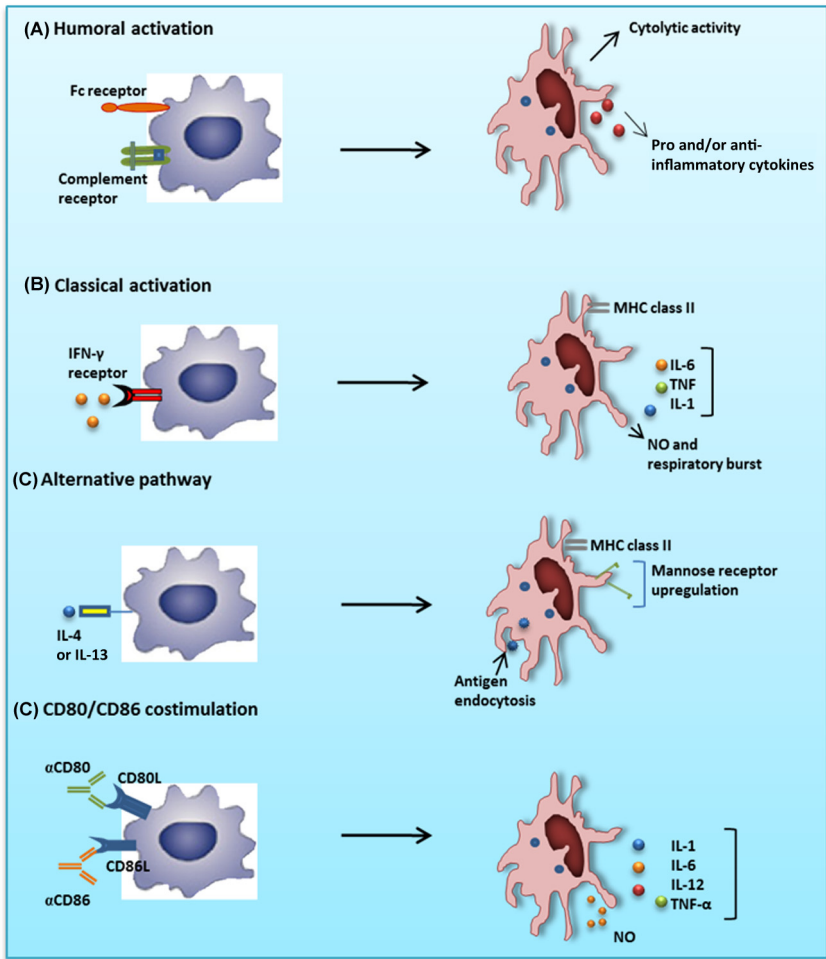


FIGURE 2.6 Reverse costimulation activation pathways of macrophages. Macrophages are activated by pathways of innate and acquired immunity. Innate activation of macrophages occurs by microbial stimuli that are recognized TLRs, CD14 (LPS receptor), and a range of nonopsonic receptors. (A) Humoral activation is mediated by some Fc and complement receptors. (B) Classical activation is mediated by the priming stimulus of IFN- γ , followed by a microbial trigger. (C) Alternate activation is mediated by IL-4 and IL-13 acting through a common receptor chain IL-4R α . (D) Costimulatory activation of macrophages occurs by delivery of signals through CD80 and CD86 molecules. This whole figure depicts that in addition to well-known innate, humoral, classical, and alternate activation, macrophages can also be activated by delivery of signals through CD80 and CD86 molecules.

allow cytotoxicity of target cells by CTLs *ex vivo*. In contrast, CD80-transfected tumor cells become susceptible to lysis by CTLs *ex vivo*. Although priming of antitumor CTLs can also take place in absence of CD80, the effector T cells are not produced without the expression of CD80 on tumor cells.⁷⁵

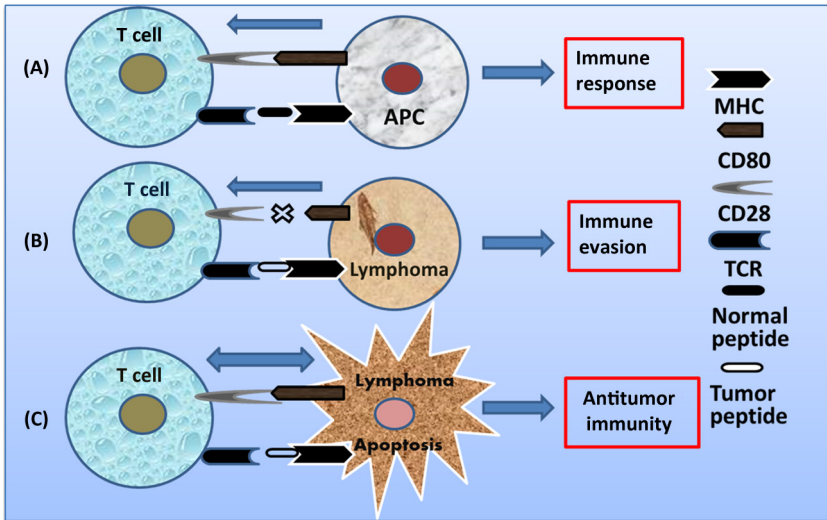


FIGURE 2.7 Reverse costimulation through CD80 in modulating the activity of tumor cells. When a normal cell presents MHC–peptide molecule complex to T cells, the delivery of CD80 costimulatory signal generates an effective immune response (A). Efficient presentation of peptide in association with MHC but in absence of costimulation induces immunological ignorance in T cells and immune evasion in cancer cells (B). Presentation of tumor-peptide–MHC complex and expression of CD80 costimulatory molecule on tumor cells leads to its apoptosis and elicitation of antitumor immunity by the activation of T cells and reverse costimulation (\leftrightarrow) through the CD80 molecule (C).

Interestingly, CD80 has also been known to enhance the memory response in CTLs.⁷⁶ This suggests that CD80 expression on tumor is important for antitumor CTL effector function. Expression of CD80 on tumor cells confers direct presentation capacity to prime naive CTLs *in vivo*.⁷⁷ Transfection of tumor cells with CD80 will help in generating long-lasting immunity. Therefore, this approach may lead to the development of a successful vaccine against cancer.

CD80 expression is needed for the sustained predominance of CD8+ T cells within a tumor (Figure 2.7C). Transfection of CD80 cDNA into human erythroleukemia DR+ cells induces the allogeneic response of purified T cells from both the cord blood and the peripheral blood of adult donors, demonstrating that CD80 expression could lead to accessory cell-independent activation of naive T cells. Furthermore, in the P815 tumor system, CD86 was substituted for CD80; interestingly, no specific CTL activity was observed. Even though in the tumors that express CD86 there is a need for CD80 expression by host APCs for efficient eradication.⁷⁸ Thus, it is apparent that transfection of CD80 in tumor cell serves as a cofactor for IL-2 production and in prevention of anergy by TCR ligation.⁷⁹ This suggests that

expression of CD80 on cancer cells contributes toward the activation of T cells responsible for imparting anticancer immunity. The expression of CD80 on tumor cells also enhances natural killer (NK) cell recognition and lysis of tumors, which plays an important role in tumor immunity.⁸⁰ CD80 costimulation can direct the CD8 T cells to produce IL-2, proliferate, and acquire cytolytic activity.⁸¹ This helper-independent generation of CTLs may have practical application in the development of tumor-specific immunotherapy.⁸² Therefore, it is clear that downregulation of CD80 costimulatory molecule by various cancer cells acts as an effective immune evasion mechanism (Figure 2.7B). This has been reported in multiple myelomas, carcinomas, leukemias, and transplantable malignancies. Some of the colon carcinomas such as MC38 and melanomas such as B16 also lack the expression of CD80. This is evident from the fact that silencing of CD80 expression results in tumorigenicity and transfection of CD80 leads to decreased tumorigenicity.⁸³ CD80 expression triggers *in vitro* NK cell-mediated cytotoxicity. The CD80 gene product functions as a triggering signal for NK cell-mediated cytotoxicity and the strength of this response is such that it overrides the inhibitory signals mediated by MHC-I molecules.^{4,84} The efficient control of solid allogeneic tumors by NK cells depends on the codelivery of both CD80 and MHC-I on the tumor cells. The codelivery is required for optimal expansion and effector function of NK cells in response to both melanoma and plasmocytoma that express allogeneic MHC-I. The two signals required for T-cell function can also regulate NK cell immunity and reveal an important similarity between the innate NK cell response and the adaptive T-cell response. Furthermore, it has been reported that NK cells and CD8 + T cells can eliminate CD80-expressing tumors that are resistant to IL-12 gene therapy.^{85,86} Hence, expression of CD80 on tumor cells may also have an important bearing on NK cell effector function.

REVERSE COSTIMULATION THROUGH CTLA-4

Recent evidences indicate that engagement of CTLA-4 onto B7 molecules expressed on APCs influence immune responses by delivering suppressive signals into DCs.²³ It was shown that both a soluble form and a membrane-anchored form of CTLA-4 on regulatory T cells can activate the immunosuppressive pathway of tryptophan catabolism in DCs. Engagement of CD80/CD86 on murine DCs by CTLA-4-Ig activates STAT1 and downregulates the SOCS-3 pathway and ultimately stimulates DCs to produce IFN- γ , which acts in an autocrine or paracrine fashion to induce IDO, an enzyme that degrades tryptophan to by-products and inhibit T-cell proliferation.²³ Further, IDO production by this way has been shown to play an important role in maintaining maternal–fetal tolerance and suppressing T-cell responses to MHC-mismatched allograft, tumors and self-antigens.⁸⁷ The ligation of CTLA-4 on mouse CD4 + CD25 + regulatory T cells can

mediate the same IDO-inducing effect *in vitro*.⁴⁷ CTLA-4-Ig, a fusion protein containing the extracellular domain of CTLA-4 linked to an IgG1 Fc region, can block the interaction of CD28 with B7-1 and B7-2 and suppress immune responses in multiple preclinical models of autoimmune and inflammatory disease.⁸⁸ However, exacerbations of disease are also seen in these preclinical models after treatment with CTLA-4-Ig,^{88,89} complicating predictions for therapeutic use. Exacerbations might result from the blockade of B7-1 and B7-2 interactions with the inhibitory CTLA-4 receptor or from the loss of regulatory T-cell function, which is dependent on CD28 signaling.^{90,91} The use of CTLA-4-Ig was first reported in patients with psoriasis vulgaris; some efficacy was observed, although statistical limitations of the trial design precluded identification of a robust therapeutic dose.⁹² More recently, CTLA-4-Ig has been tested in combination with methotrexate (MTX) in clinical trials of rheumatoid arthritis.⁹³ In these studies, 2 or 10 mg/kg doses of CTLA-4-Ig were administered to patients on days 1, 15, and 30, and monthly thereafter for a total of 6 months; patients also received approximately 15 mg MTX per week. Evaluation at 6 months showed that a higher percentage of patients receiving 10 mg/kg CTLA-4-Ig/MTX had improved clinical scores compared with those on MTX alone ($P < 0.001$). Further studies will be needed to assess long-term efficacy and safety for CTLA-4-Ig, and to elucidate its mode of action in humans. However, these promising results suggest that manipulation of costimulatory signals can ameliorate inflammatory disease. CTLA-4 is a negative regulator of T-cell activation, and Abs that block CTLA-4 can enhance T-cell responses.⁴ In a recent clinical trial, stage IV melanoma patients were treated with antihuman CTLA-4-Ab in combination with a gp100 melanoma-associated antigen peptide vaccine.⁹⁴ Some tumor regression was observed, although patient numbers were too small to draw conclusions on efficacy. Several patients developed manifestations of autoimmune responses in their skin, liver, or intestines, which appeared to resolve after discontinuation of Ab therapy. The autoimmune manifestations limited patient accrual but suggested that the anti-CTLA-4 Ab-enhanced immune responses. Similar autoimmune adverse events were seen in a second small clinical trial.⁹⁵ Alternative clinical approaches have explored the use of vaccination strategies with tumor cells expressing high levels of B7-1 or B7-2 to enhance tumor immunity, although robust clinical protocols have yet to emerge.^{96–98}

B7-DC Reverse Costimulation

B7-DC does not only costimulate T-cell proliferation and IFN- γ production by T cells, but its cross-linking on DC also modulates DC biology. B7-DC cross-linking on DCs enhanced antigen presentation and IL-12p70 production *in vitro*. Furthermore, anti-B7-DC treatment increased the survival of

DCs *in vitro* and the migration of adoptively transferred DCs reaching draining lymph nodes *in vivo*.²¹ B7-DC cross-linking has also been shown to restore the antigen-acquiring function of immature DCs and augments APC function of matured DCs.²⁸ This shows that B7-DC can function as a means for the communication of signals directly to DCs from ligands in the environment. One of the challenges in targeting the CD28/CTLA-4/B7 pathway has been the shared receptor–ligand interactions. Although CTLA-4–Ig can block stimulatory CD28 signals, it might also interfere with inhibitory CTLA-4 signals. Recent data also indicate that CTLA-4–Ig delivers “reverse signals” mediated by B7 ligands directly into the APC.^{23,99} Alternative strategies to block B7 ligands using anti-B7 Abs have been promising in preclinical studies, including primate studies in transplantation¹⁰⁰ and many rodent autoimmune and inflammatory disease models.⁹⁰ Recent data indicate that B7-1 and B7-2 can signal into DCs, generating tolerogenic or immunogenic signals depending upon whether the B7s are engaged by CTLA-4 or CD28,³⁸ respectively, further suggesting that blockade using Abs to B7-1 and B7-2 may differ from using CTLA-4–Ig. Strategies to identify blocking Abs to CD28 for clinical use must consider potential immune activation mediated by unintentional cross-linking of CD28; this would also apply to use of blocking Abs to inducible costimulator (ICOS).³⁸ Although small molecules have been reported that block the interaction of B7-1 with CD28,^{101,102} these are of limited potency, and there are no reports of small molecules that block the interactions of both B7-1 and B7-2 with CD28. Thus, new approaches that specifically target CD28 are needed. Additionally, approaches that spare CD28-dependent regulatory T cells would allow preservation of these important cells in inflammatory and autoimmune diseases. These results suggest that bidirectional B7–CTLA-4 interactions may participate in down-regulation of T-cell responses and induction of T-cell tolerance. Therefore, CTLA-4 not only regulates TCR and CD28 signals in T cells but also delivers signals through CD80/CD86 into DCs to induce IDO and the subsequent biological effects. Binding of CD28 to B7 ligands on DCs leads to up-regulation of IL-6 production by DCs and subsequent immunostimulatory effects on T-cell activation.³⁸ However, binding of CTLA-4 to B7 ligands upregulates IFN- γ , which then upregulates the expression of the enzyme, IDO, resulting in tryptophan catabolism and suppression of T-cell proliferation.^{23,87}

Reverse Costimulation Through CD40L/CD40

The CD40–CD40L receptor–ligand interaction plays a central role in regulating the immune responses by activating and modulating DCs, macrophages, and T cells. CD40–CD40L interaction has been shown to induce CD80 and CD86 expression on other APCs, thereby promoting their costimulatory capacity. CD40 ligation with CD40L on DCs promotes the

upregulation of the costimulatory molecules such as CD80 and CD86,^{103,104} IL-12 secretion and release of chemokines such as IL-8, MIP-1 α , and MIP-1 β .^{105–107} CD40 ligation has also been shown to upregulate the expression of OX40L, another member of the TNF super family on DCs. CD40L–CD40 interactions mediate B-cell proliferation without further costimulus. Central functions of the B cells, such as proliferation, up-regulation of membrane activation markers, isotype switching (in conjunction with different cytokines), regulation of apoptosis, and memory B-cell formation, are all dependent on signals provided through the CD40–CD40L interactions.^{108,109} In order for the B cells to enter the cell cycle, produce Ig, and undergo Ig class switching or somatic hypermutation, it is necessary for them to receive the appropriate T-cell help through CD40–CD40L interaction.¹¹⁰ It has been reported that CD28 ligation stabilizes CD40L mRNA, allowing for more rapid translation and transport of CD40L onto the T-cell surface, and therefore results in increased B-cell responses.¹¹¹

The interrelationship between the CD28/B7 and CD40L/CD40 pathways plays an important role in the ability of B cells to present antigen. Resting B cells express little if any CD80 or CD86, but CD40 ligation induces their expression. B cells activated with anti-CD40 Ab stimulated an allogeneic MLR more efficiently, and this stimulation was blocked by human cytotoxic T-lymphocyte antigen-4–Ig. Further investigations have revealed that CD40 cross-linking by Ab or the interaction of CD40 with its ligand CD40L upregulated CD80 and CD86.¹¹² Incubation of resting B cells with activated T cells is another potent means of inducing CD80 and CD86. In this system, CD80 upregulation was inhibited by either anti-CD40 or CD40L Abs, consistent with the notion that CD40L/CD40 regulates CD80 expression. However, the CD40L antagonist did not completely inhibit the ability of activated T cells to upregulate CD86. Therefore, other signals mediated by MHC-II engagement or cytokines are also involved in CD86 regulation by activated T cells.

CD40 is an important regulator of macrophage function as it stimulates the production of TNF- α and other factors.¹¹³ In the presence of IFN- γ , CD40 signaling enhances NO production and anti-*Leishmania* and anti-*Toxoplasma gondii* activity of macrophages.^{114,115} However, it is not known whether CD40 can replace IFN- γ as the priming signal for induction of macrophage antimicrobial activity. However, it has been recently shown that CD40 signaling induces the antimicrobial activity of macrophages against intracellular pathogen *T. gondii* despite the lack of two central features of classically activated macrophages, that is, priming with IFN- γ and production of reactive nitrogen intermediates. This CD40 signaling primes macrophages in response to TNF- α to acquire antimicrobial activity.

Interaction between the CD40 receptor on APC and its ligand (CD40L) on activated T cells plays a critical role in immunity to intracellular pathogens by upregulating the production of IL-12.¹¹⁶ CD40- or CD40L-deficient

mice have an increased susceptibility to *Leishmania* infection and show an impaired priming of Th1-type cells, correlating with a lack of activation of the macrophage effector functions required for parasite clearance.¹¹⁴ CD40 stimulation of bacillus calmette guerin-infected DCs not only promotes their ability to secrete IL-12 but increases the release of other inflammatory mediators such as IL-1 α , IL-1 β , and IL-6.¹¹⁷ These may potentiate the development of inflammatory responses, which play a critical role in anti-mycobacterial immunity.

REVERSE SIGNALING THROUGH CD137L/CD137

CD137L is expressed in APCs and other myeloid cells (B cells, macrophages, DCs, mast cells, and eosinophils) and nonhematopoietic cells (endothelial cells, fibroblasts, and epithelial cells).¹¹⁸ Evidence supporting that CD137L signals play an *in vivo* physiological role in inflammation is just being emerged, even though accumulating evidence has demonstrated the existence of CD137L signals at molecular and cellular levels. For example, CD137L signaling mediates cellular functions ranging from cell differentiation, proliferation, and survival to the production of inflammatory mediators in a variety of cells.¹¹⁹

It is now becoming clear that CD137L signaling is critical in multiple phases of inflammation. Inflamed vessels express CD137 and CD137L, and CD137L signaling in endothelial cells leads to the production of proinflammatory cytokines and chemokines.^{120,121} Further, CD137L signaling may facilitate transendothelial migration of leukocytes through upregulation of cell adhesion molecules on endothelial cells.^{120,121} However, CD137L signaling increases the expression of cell adhesion molecules on monocytes and promotes their extravasation. Because endothelial cells express both CD137 and CD137L, CD137–CD137L interactions between endothelial cells and leukocytes may amplify inflammation in such a way that endothelial cells induce sustained production of inflammatory mediators and prime leukocytes before they arrive at inflamed tissue territories. In the tissues, it seems that CD137L signaling in recruited leukocytes, residential cells, and parenchymal cells is also critical in the amplification of inflammation. Macrophages express CD137L on exposure to an inflammatory environment and produce high levels of proinflammatory cytokines and chemokines in response to CD137L signals.^{122–124} In collaboration with other inflammatory inducers, CD137L signaling results in the production of inflammatory mediators in macrophages in a synergistic manner,¹²⁵ an indication that CD137L signaling is an amplifier of inflammation. It is noteworthy that CD137L can sustain TLR signaling by binding to TLRs without engagement of CD137.¹²⁴ Recently, we have identified a novel inflammatory pathway involving CD137L signaling in epithelial cells.¹²⁶ In kidney ischemia–reperfusion injury, CD137 expressed on infiltrated NK cells stimulates CD137L in

tubular epithelial cells to produce CXCR1 and CXCR2 that are required for recruitment of neutrophils in the kidney. Because the kidney ischemia–reperfusion injury does not occur without NK cells or neutrophils, it is thought that the axis of NK cell tubular epithelial cell neutrophils is the major pathogenic pathway for kidney ischemia–reperfusion injury.¹²⁷

There are few reports on the roles of CD137L signaling in disease context. As mentioned above, CD137L signaling is indispensable for kidney ischemia–reperfusion injury.¹²⁶ Considering that CD137L signaling is critical in inflammatory responses, it is predicted that milder inflammatory diseases will occur in the absence of CD137L signals. Indeed, we have demonstrated that blocking of CD137L can inhibit inflammatory responses and prevent mortality in *C. albicans*-induced sepsis (unpublished data). In this model, CD137 signaling enhances the phagocytic activity of neutrophils, whereas CD137L signaling induces massive cytokine production by macrophages following *C. albicans* infection. In this model, agonistic anti-CD137 mAb has dual beneficial effects on *C. albicans*-induced sepsis by promoting fungal clearance by neutrophils and downregulating cytokine production by macrophages.

Emerging evidence suggests that both CD137 and CD137L signals play an important role in the evolution of inflammation. CD137 signaling in lymphoid cells such as NK cells, NK T cells, and T cells promotes inflammation by producing IFN- γ and TNF- α . However, the outcomes of CD137 signaling in regulatory T cells and neutrophils are connected to the suppression of inflammation. CD137 signaling in regulatory T cells and NK cells induces production of immunosuppressive cytokines that directly decrease inflammation (unpublished result). CD137 signaling in neutrophils can facilitate pathogen clearance and thus contributes to rapid resolution of inflammation (unpublished results, Mir and Agrewala). CD137L signaling seems to always have proinflammatory actions.

Role of Reverse Costimulation in Cancer

Clinical studies have demonstrated that administration of anti-CD80 Ab can decrease tumor burden in patients suffering from lymphomas.⁴³ CD80 is expressed transiently on the surface of activated B cells and other APCs, including DCs, but is constitutively expressed on a variety of NHLs, including follicular lymphomas. Thus, CD80 is an attractive target for lymphoma therapy. Targeting CD80 on NHLs with anti-CD80 Ab can arrest their growth and therefore can serve as a therapeutic intervention for treating cancer. Engaging costimulatory pathways, as well as blocking coinhibitory pathways, has therapeutic potential for treatment of cancer. However, for tumor immunotherapy, there are multiple barriers to the antitumor response, and coinhibitory pathways participate in shielding tumors from immune eradication. The differential expression of coinhibitory molecules (such as B7-H4) on certain cancers provides an opportunity for selective immunotherapeutic

intervention. Interference with multiple coinhibitory pathways may be needed for optimal therapeutic benefit. Despite the expression of antigens by tumor cells, spontaneous immune-mediated rejection of cancer seems to be a rare event. TCR engagement by peptide–MHC constitutes the main signal for the activation of naive T cells but is not sufficient to initiate a productive generation and maintenance of effector cells. Full activation of T cells requires additional signals driven by costimulatory molecules present on activated APCs but rarely on tumors. Following the discovery of B7-1 (CD80), several other costimulatory molecules have been shown to contribute to T-cell activation and have relevance for improving antitumor immunity. Moreover, increasing the understanding of coinhibitory receptors has highlighted key additional pathways that can dominantly inhibit antitumor T-cell function. Improving positive costimulation, and interfering with negative regulation, continues to represent an attractive immunotherapeutic approach for the treatment of cancer. The pathways with the highest potential for clinical application and translation to the clinic summarizing clinical trials aimed at boosting positive costimulation and antitumor immunity with agonistic CD40, 4-1BB, or OX40 Abs, and clinical trials that block the coinhibitory receptors CTLA-4 and PD-1. However, challenges are posed by the development of inflammation or autoimmunity following immune intervention.

A significant number of patients relapse or do not respond to rituximab due to intrinsic or acquired resistance. Hence, mAbs targeting other cell surface antigens on B-cell lymphomas are being studied. CD80 serves as an attractive target in the continued development of mAbs against lymphoma. Preclinical studies with galiximab, an anti-CD80 primatized mAb, have been encouraging and have demonstrated antitumor activity against various B-cell lymphoma models, both as a single agent and in combination with rituximab. Data were reviewed from a PubMed literature search from 1975 to 2009 and also included a review of abstracts from published proceedings of annual meetings from the American Society of Hematology and International Conference of Malignant Lymphoma, Lugano.

Galiximab (Biogen Idec, Inc., San Diego, California) is a chimeric, primatized IgG1 mAb consisting of human constant and primate (cynomolgus macaque) variable regions targeting CD80. It binds specifically to CD80, which is transiently expressed on the surface of activated B cells and APCs, including DCs, but is constitutively expressed on a variety of NHLs, including FL. Thus, CD80 serves as an attractive target for the development of antilymphoma therapy. Galiximab is emerging as an interesting and effective therapy for the treatment of patients with NHL, especially in combination with rituximab. Its favorable toxicity profile makes it an attractive alternative to existing agents. This “primatized” Ab is structurally indistinguishable from human Abs and, therefore, is not significantly immunogenic in humans. Although a number of *in vivo* as well as *in vitro* studies point toward possible “unique” mechanisms of action of galiximab, more studies are needed to

fully understand this agent's antitumor activity, as well as to establish strategies by which to optimize its efficacy. However, to further assess the therapeutic role of galiximab in the treatment of patients with follicular and other lymphoma subtypes, as monotherapy or in combination with rituximab or systemic chemotherapy, additional studies are under active consideration.

As established by clinical and experimental results, it is quite evident that anti-CD80 Ab treatment is effective against relapsed and refractory follicular lymphoma. Future studies in other CD80-positive hematological malignancies (e.g., diffuse large B-cell NHL, chronic lymphocytic leukemia, Hodgkin's lymphoma, and multiple myeloma) should also be taken under consideration. Thus, anti-CD80 Ab immunotherapy may have a potent role in the treatment of CD80-bearing cancer cells because their binding can modulate the key molecules in the signaling pathway and enhance antitumor response. Considering the importance of CD80 signaling in the regulation of immune responses against cancer, the manipulation of this signaling pathway to increase immunity against cancer represents a potential therapeutic approach. Understanding the mechanisms by which tumor cells escape immune surveillance will help us to establish new and effective approaches to vaccination and immunotherapy. One of the most important immune evasion mechanism employed by tumor cells is the downregulation of the CD80 costimulatory molecule. The success of novel cancer therapies depends on the identification of functional targets that play an essential role in tumor growth and metastasis, survival, and evasion from immunosurveillance. Anti-CD80 therapy can be used to target tumors expressing CD80. The clinical success of anti-CD80 Ab for the treatment of refractory follicular lymphoma has stimulated great interest in the promise of Ab therapeutics for cancer. The qualities of galiximab, such as long half-life and high specificity and safety compared with other cancer therapeutics together with its ability to bind to and modulate key players in signaling pathways that drive malignant transformation and enhance antitumor immune functions, makes it a highly desirable therapeutic agent.

Most tumors do not regress but continue to grow in spite of the presence of spontaneous or antigen-induced immune responses, due to downregulation of costimulatory molecules such as CD80. The existence of systemic immune responses may not by itself be sufficient to deal with the complex nature of tumor–host interactions because factors such as insufficient costimulation to induce T-cell response may further contribute to the lack of effective immunity. It is now well established that T cells are rendered anergic due to the lack of costimulatory molecule(s) expression by tumor cells. Tumor cells are able to induce antigen-specific tolerance or anergy on the basis of MHC-I-restricted antigen presentation without the expression of costimulatory molecules. This unresponsiveness, however, can be reversed when tumor cells are genetically modified to express costimulatory molecules. A plethora of studies suggest that the insertion of genes encoding CD80 into tumors generally

increases their immunogenicity and can be used as vaccine. Recently, fusogene vectors have been developed to encode multiple gene products such as CD80 with cytokines or MHC molecules as fusion proteins from a single cistron to increase the immunogenicity of target tumor cells. The vectors generated could be used in immunotherapy for the treatment of multiple myeloma, leukemias, and other cancers, as they have been shown to stimulate allogeneic mixed lymphocyte proliferation and augment increases in CTL and NK cell responses and IFN- γ release. Considering the importance of costimulation in the regulation of immune responses against relapsed cancer, the manipulation of this pathway to increase immunity represents a potential therapeutic approach. Furthermore, signaling either by anti-CD80 Abs or through CD28-bearing T cells regresses the growth, augments the expression of proapoptotic molecules, and induces apoptosis of CD80 + lymphomas. Therefore, immunotherapy utilizing anti-CD80 Abs is a promising future treatment, especially in the case of relapse and refractory lymphomas.

Therapeutic modalities targeting the B7 and CD28 families of ligands and receptors are showing promise in the clinic. The encouraging results in targeting the B7/CD28/CTLA-4 pathway in psoriasis and rheumatoid arthritis underscore the importance of this costimulatory pathway and validate the pathogenic contributions of T cells in the etiology of these diseases. The costimulatory ICOS pathway might also provide therapeutic opportunities, particularly in diseases in which pathogenic Abs contribute to disease progression. Although the negative regulators CTLA-4 and PD-1 are critical for attenuating normal immune responses, agonistic compounds targeting these proteins to “tune down” inflammatory responses have not yet been identified. This is probably a result of the requirement for coengagement of PD-1 or CTLA-4 with the antigen receptor for delivery of inhibitory signals. However, antagonists of these inhibitory receptors might have utility in enhancing immune responses for cancer, infectious diseases, or vaccine development. The CD28 receptor/B7 ligand families are not yet complete, as receptors for the new B7-H3 and B7-H4 ligands have not been identified, and some data support additional receptors for PD-L1 and PD-L2. Many receptors and their ligands belonging to B7 and TNF/TNFR families of molecules through whom reverse costimulation can be delivered for the activation of APCs and the cells expressing them (Figure 2.8). The role of CD28 in the development of regulatory T cells is further explored, and it has been shown that CD28 is not only required for IL-2 production but also for differentiation of regulatory T cells, independently of IL-2 production. Further support for the costimulatory role of B7-H3 has been demonstrated in allograft rejection models. Treatment of B7-H3-deficient mice with cyclosporin A, rapamycin, or anti-CD154 Ab reduced the incidence of rejection of cardiac or allografts compared with treated wild-type mice. It is postulated that B7-H3 contributes to allograft rejection by induction of IFN- γ or by chemokines induced by IFN- γ . Despite the complex roles and interactions within

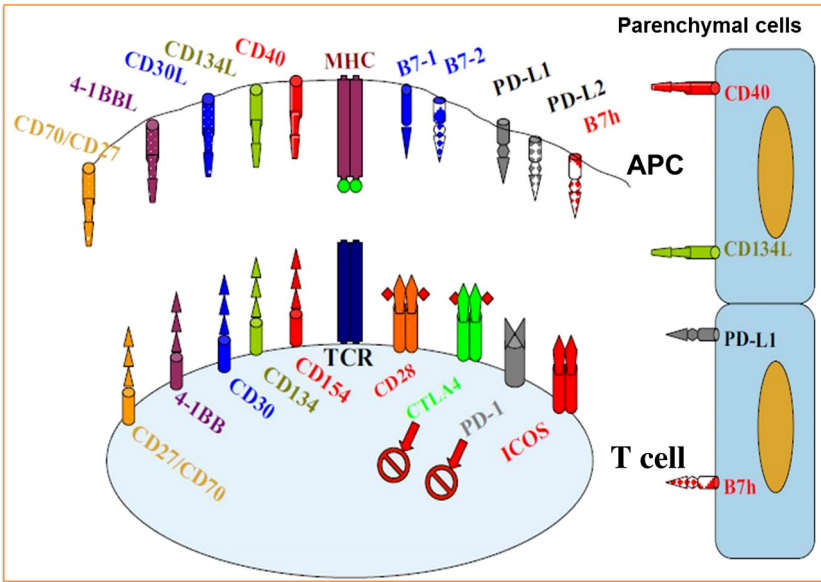


FIGURE 2.8 Schematic representation of costimulatory molecules through which reverse costimulation can be delivered. Signal 1 is represented by the TCR interacting with the antigen in the MHC groove. Signals from the CTLA-4 and PD-1 are inhibitory to the T cell. 1BB, CDw137; 1BBL, CDw137 ligand; CTLA-4, cytotoxic T-lymphocyte associated gene-4; ICOS, inducible costimulator; PD, programmed death; TCR, T-cell receptor.

the CD28 and B7 families, we predict that novel approaches targeting these families will yield new therapies for the treatment of inflammation, autoimmunity, transplantation, cancer, and infectious diseases.

Many studies published recently have now established that the CD80 and CD86 costimulatory molecules expressed on the surface of the APCs do not merely have a unidirectional function of stimulating T cells on engagement with CD28 and CD152, but this interaction delivers bidirectional signals affecting the activity of APCs as well. Moreover, based on the kinetics of differential expression, affinity, association and dissociation rates, and structure, indicates that CD80 and CD86 can differentially regulate the cells of immune system. In particular, CD80 delivers similar signals as delivered by CD152 for inhibiting the action of T cells. In contrast, CD86 costimulate APCs in an analogous manner as done by CD28 in activating T cells. Interestingly, the inhibitory role of CD80 on B cells, DCs, neural stem cells, and lymphomas has shown to induce apoptosis by upregulating the expression of proapoptotic molecules and downregulating the expression of antiapoptotic molecules. This inhibitory role of CD80 has been efficiently exploited to treat patients suffering from relapsed and refractory lymphomas. In addition, signaling through CD80 can be explored for

inducing tolerance/apoptosis in autoimmune diseases, hypersensitivity reactions, allergies, and transplantation.

Considering the importance of costimulation in the regulation of immune responses against relapsed cancer, the manipulation of this pathway to increase immunity, regresses the growth, augments the expression of proapoptotic molecules, and induces apoptosis of lymphomas represents a potential therapeutic approach. Therapeutic modalities targeting the B7 and CD28 costimulatory families of ligands and receptors are showing promise in the clinic. The encouraging results in targeting the B7/CD28/CTLA-4 pathway in the autoimmune diseases such as psoriasis, multiple sclerosis, and rheumatoid arthritis underscore the importance of this costimulatory pathway and validate the pathogenic contributions of T cells in the etiology of these diseases. This novel strategy of costimulation activation/inhibition can be effectively exploited to develop immunotherapy using humanized Abs against CD80, CD86, and CD40 or CD28 fusogenic proteins for the treatment of intracellular pathogens such as *M. tuberculosis*, HIV, *Leishmania donovani*, and *Trypanosoma cruzi*. This strategy can also be used as an alternative strategy or in combination with the drugs. This approach is based on modulating the immune system of the hosts rather than targeting the pathogen; hence, it significantly diminishes the chance of emergence of drug-resistant strains of pathogens and if applied properly may overcome the rising menace of infectious diseases. To develop alternative or adjunct (with drugs) therapies using costimulatory molecules, an intensive effort has been undertaken in the last decade to understand how intracellular pathogens exploit costimulatory molecules, which are the *tour de force* of the immune system. The potent role of costimulatory molecules is aptly established in the optimum activation of T cells and APCs, the cells that play a cardinal role in curbing the infections. Hence, immunotherapy involving costimulatory molecules can be a breakthrough strategy to treat various diseases, to minimize side effects inflicted by drug therapies, and to restrict the emergence of drug resistance.

To translate this field into the clinic, it is urgent to develop novel methods to target currently appreciated costimulatory pathway and deeply understand pathophysiology of these diseases involving costimulatory molecules. Despite the complex roles and interactions within the CD28 and B7 costimulatory families, we predict that novel approaches targeting these families will yield new therapies for the treatment of inflammation, autoimmunity, transplantation, cancer, and infectious diseases.

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Chapter 3

Costimulation Immunotherapy in Infectious Diseases

INTRODUCTION

The immune system is a sophisticated and complex weapon that has evolved to destroy invading pathogens. Multicellular organisms possess very sophisticated defense mechanisms that are designed to effectively counter the continual microbial insult of the environment within the vertebrate host. However, successful microbial pathogens have in turn evolved complex and efficient methods to overcome innate and adaptive immune mechanisms, which can result in disease or chronic infections. Although the various virulence strategies used by viral and bacterial pathogens are numerous, there are several general mechanisms that are used to subvert and exploit immune systems that are shared between these diverse microbial pathogens. The success of each pathogen is directly dependent on its ability to mount an effective anti-immune response within the infected host, which can ultimately result in acute disease, chronic infection, or pathogen clearance.¹ The three biggest global infectious disease threats to humans are human immunodeficiency virus (HIV), tuberculosis, and malaria, each killing one to two million people worldwide each year. Each of these three causative agents (which represent a virus, a bacterium, and a parasite) have developed highly effective mechanisms to subvert the human immune system, which explains why developing vaccines and controlling these pathogens have been so difficult. Successful pathogens have evolved a range of anti-immune strategies to overcome both innate and acquired immunity (Table 3.1), which play critical roles in their abilities to cause disease. Although at first glance the immunomodulatory mechanisms used by viruses and bacteria might appear quite different, there are a surprising number of similarities and shared mechanistic concepts. Both types of pathogens have to overcome the same host immune mechanisms, and it is illustrative to see how they have developed parallel strategies to neutralize host immunity. Moreover, viral and bacterial diseases are often linked, exploiting weaknesses in host defenses that are caused by another pathogen.^{2–5}

TABLE 3.1 Various Strategies Used by Pathogens (Bacteria and Viruses) to Evade Effective Immune Responses of the Host

Strategy Used by Pathogens	Examples from Bacteria	Examples from Viruses
Subvert or kill immune cells/phagocytes	<ul style="list-style-type: none">– Superantigens– Avoid phagolysosomal fusion– Block inflammatory pathways by injecting effectors– Replicate within and overrun immune cells	<ul style="list-style-type: none">– Infect and kill immune cells (DCs, APCs, lymphocytes, macrophage, etc.)– Inhibit CTL/NK cell killing pathways– Alter immune cell signaling, effector functions, or differentiation– Express superantigens
Block acquired immunity	<ul style="list-style-type: none">– IgA proteases– Block antigen presentation	<ul style="list-style-type: none">– Downregulate MHC-I or -II– Block antigen presentation/proteasome– Prevent induction of immune response genes
Inhibit complement	<ul style="list-style-type: none">– Proteases to degrade complement– Produce capsules and long chain LPS to avoid complement deposition and MAC attack	<ul style="list-style-type: none">– Soluble inhibitors of complement cascade– Viral Fc receptors
Inhibit cytokines/interferon/chemokines	<ul style="list-style-type: none">– Block inflammatory pathways– Activate alternate pathways– Secrete proteases to degrade	<ul style="list-style-type: none">– Inhibit ligand gene expression– Ligand/receptor signaling inhibitors– Block secondary antiviral gene induction– Interfere with effector proteins
Modulate apoptosis/autophagy	<ul style="list-style-type: none">– Inhibit apoptosis– Activate death signaling pathways– Alter apoptotic signaling pathways	<ul style="list-style-type: none">– Inhibit or accelerate cell death– Block death signaling pathways– Scavenge free radicals– Downregulate death receptors or ligands– Inactivate death sensor pathways

Interfere with TLRs	<ul style="list-style-type: none"> – Alter TLR ligands to decrease recognition – Bind to TLR to dampen inflammation – Inject effectors to inhibit downstream inflammation signaling 	<ul style="list-style-type: none"> – Block or hijack TLR signaling – Prevent TLR recognition
Block antimicrobial small molecules	<ul style="list-style-type: none"> – Secrete proteases to degrade – Alter cell surface to avoid peptide insertion – Use pumps to transport peptide – Directly sense small molecules to trigger defense mechanism 	<ul style="list-style-type: none"> – Prevent inducible nitric oxide synthase (iNOS) induction – Inhibit antiviral RNA silencing
Block intrinsic cellular pathways	<ul style="list-style-type: none"> – Alter ubiquitin pathway – Alter transcriptional programs 	<ul style="list-style-type: none"> – Inhibit RNA editing – Regulate ubiquitin/ISGylation pathways
Secreted modulators or toxins	<ul style="list-style-type: none"> – Many toxins – Proteases 	<ul style="list-style-type: none"> – Ligand mimics (virokines) – Receptor mimics (viroceptors)
Modulators on the pathogen surface	<ul style="list-style-type: none"> – Lipid A of LPS – Carbohydrates such as capsules – Outer membrane proteins – Adhesins and invasins 	<ul style="list-style-type: none"> – Complement inhibitors – Coagulation regulators – Immune receptors – Adhesion molecules
Hide from immune surveillance	<ul style="list-style-type: none"> – Avoid phagolysosomal fusion – Inhibit phagocytosis 	<ul style="list-style-type: none"> – Latency – Infect immunoprivileged tissues
Antigenic hypervariability	<ul style="list-style-type: none"> – Vary many surface structures – Pili, outer membrane proteins, LPS – Strain to strain variation 	<ul style="list-style-type: none"> – Express error-prone replicase – Escape from antibody recognition – “Outrun” T-cell recognition

For intracellular bacteria, entry into host cells represents the central requirement for survival in, as well as elimination by, the host. Host-cell-directed uptake, called “phagocytosis,” is a feature of the so-called professional phagocytes that comprise polymorphonuclear granulocytes and mononuclear phagocytes. Entry induced by the pathogen and termed invasion allows entry into nonphagocytic cells (nonprofessional phagocytes). Contact between host cells and pathogens proceeds either directly via receptor–ligand interactions or indirectly via deposition on the surface of the pathogen of host molecules for which physiologic receptors exist on the target cell. Depending on the cellular target, the final outcome of host cell entry varies markedly. Nonprofessional phagocytes are nonphagocytic, and hence entry depends on expression of surface receptors that can be misused for invasion. Benefits of having a well-performing immune system are clearly shown by the pathologies associated with congenital and/or acquired immunodeficiency. The protective function of the immune system resides in the capacity of immune cells to discriminate between self- and nonself-antigens. Major histocompatibility complex (MHC) molecules expressed on the surface of all nucleated cells (class I), and “professional” antigen-presenting cells (APCs) (class II) are an essential tool for the recognition of nonself-antigens.^{6,7} The vertebrate immune system is highly evolved to combat and eliminate pathogens (Figure 3.1), however, some pathogens can successfully subvert the host immune system to establish their intracellular survival via strategies like immunosuppression, molecular mimicry, disguise or sequestration of antigens, circumvention of complements and cytokines cascade, blockade of antigen presentation, escape from apoptosis and autophagy, and modulation of costimulatory signals.¹ It is well known that development of immune response, especially in case of naive T cells needs at least two distinct signals for full activation to proliferate and differentiate.⁸ The first signal is provided by the specific antigen (MHC–peptide complex) recognition by the T-cell receptor (TCR) and the second signal is provided by the costimulatory molecules.⁹ This requirement for second signal explains why adaptive immune responses are stimulated by microbes but not by most self-antigens, which are not normally recognized by the innate immune system and therefore do not elicit adaptive immune responses. The second signal, which is not delivered via the TCR and is not antigen specific, has been termed as costimulatory signal because, while essential, it does not by itself induce any response in T cells. However, when a T cell has its receptor ligated and receives a costimulatory signal, the T cell will proliferate and differentiate into an effector cell. Moreover, T cells that bind antigen but do not receive a costimulatory signal are thought to die or to become anergic,¹⁰ a state in which the cell cannot be activated even if it receives both of the signals required to activate a T cell. Thus, an encounter with antigen can lead to two quite distinct outcomes: proliferation and differentiation into effector cells, or inactivation or death, which outcome occurs is determined by the appropriate delivery of costimulatory signals (Figure 3.2). With the continuous discovery of new

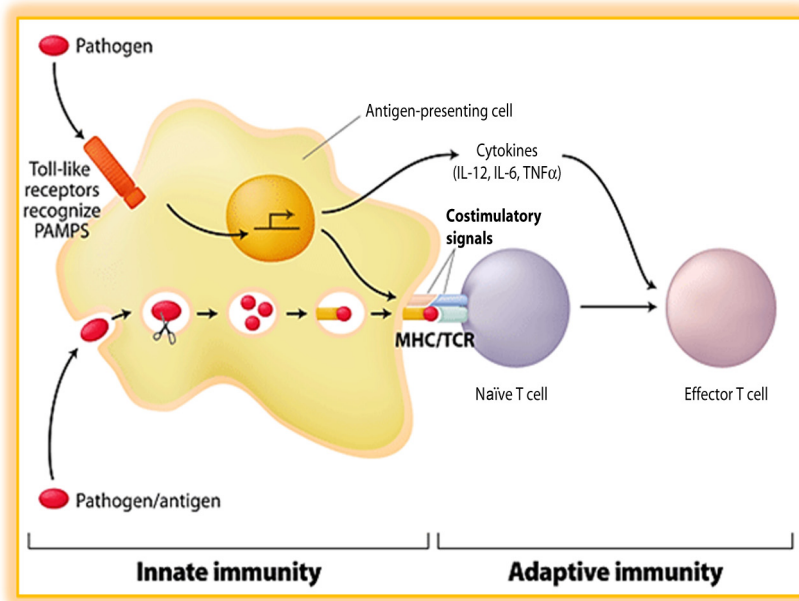


FIGURE 3.1 Processing of pathogens (antigens) by vertebrate immune system to generate immune response against pathogens. Pathogens have PAMPs on their surface that are recognized by PRRs on host cells which phagocytose the pathogen. After chopping down the pathogen in the phagosome–lysosome system, they present the antigens to T cells in context with their own MHC molecules to be recognized by naive T cells specific for that antigen so the T cell can mount an effective immune response.

costimulatory molecules on APCs and their receptors on T cells, the story is very much an evolving one. But to date, B7-1 (CD80) and B7-2 (CD86) remain the best-defined costimulatory molecules on APCs¹¹ which provide the second signals for activation of naive T cells.^{12,13} For the first time Suvas et al. from our lab gave a concept of bidirectional costimulation and demonstrated that costimulation through CD80 and CD86 not only influence the activation of T cells but by bidirectional costimulation they can also affect the activity of B cells.¹⁴ Since then many studies have shown the significance of CD80 and CD86 in influencing the activity of B cells, stem cells, and dendritic cells (DCs) but nothing is known about the role of these molecules in the case of macrophages. For instance, it is not known how these molecules regulate the activation of macrophages, which are the important cells of the immune system and the host for a plethora of intracellular pathogens,^{6,15} especially the causative agent of tuberculosis. So what is needed is an integrated picture of how signals delivered through major costimulatory molecules CD80 and CD86 expressed on the surface of macrophages regulate the type of immune response generated during infection. Hence the main aim of the present study is to

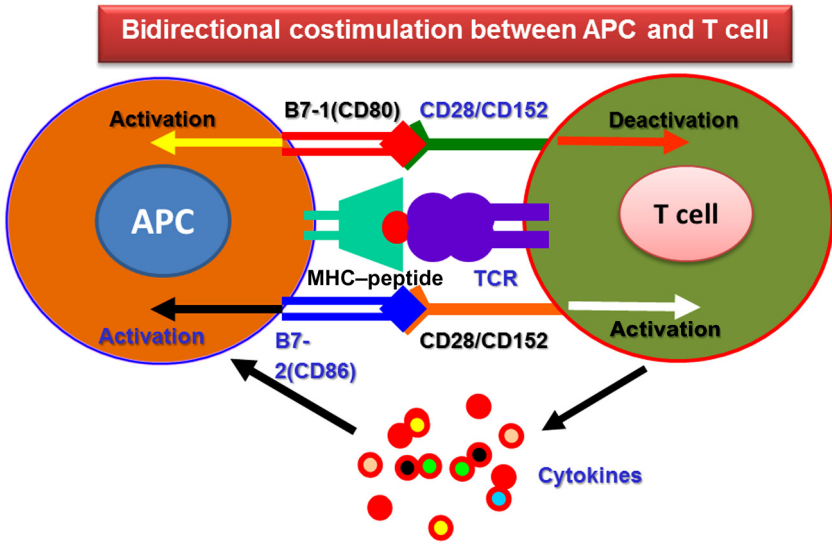


FIGURE 3.2 Innate and adaptive arms of immune responses against pathogens. Pathogens have PAMPs on their surface which are recognized by PRRs on professional APCs like DCs which then move to lymph nodes where they interact with the naive or memory T cell generating the humoral or cell-mediated immune response against the pathogen. Without the involvement of costimulatory signals the naive T cells will not be activated as the costimulatory signals are important for their activation.

monitor whether bidirectional costimulation can affect the performance of macrophages. Further, we will investigate whether costimulatory signals can be selectively delivered to activate and inhibit the function of macrophages. Further, we will investigate whether the delivery of costimulatory signals can curb the growth of intracellular pathogens. The use of agents that block or activate costimulatory pathways will undoubtedly require a better understanding of how the diverse biologic activities of these pathways are orchestrated. Also we will seek to find the better biomarkers to predict outcomes of the manipulation of T-cell and APC costimulation in humans and to find preclinical studies with established predictive value. Signaling through CD80 and CD86 can be explored for inducing tolerance/apoptosis in autoimmune diseases, hypersensitivity reactions, allergies, transplantation, and in killing the intracellular pathogens.

Some of the most dreadful pathogens of humans such as *Mycobacterium tuberculosis* (*Mtb*), HIV, and *Leishmania donovani* cause chronic infections and remain a serious global threat to humanity¹⁶ and a challenge to scientific community to find a solution for their control. These three organisms are representatives of successful pathogens from the class of bacteria, virus, and parasites. Innovative strategies have been developed by these pathogens to evade immune responses such as subversion of phagocytosis, antigenic shift

and drift, induction of immune regulatory pathways, interference with antigen processing/presentation, and more often the manipulation of the costimulatory molecules. Exploitation of the behavior of costimulation pathways provides evolutionary incentive to the pathogens and thereby abates the functioning of immune system. Impairment by pathogens in the signaling events delivered by costimulatory molecules may be responsible for defective T-cell responses; consequently organisms grow unhindered in the host cells. These molecules expressed on the surface of various cells play a decisive role in the initiation and sustenance of immunity. These pathogens have devised various convergent devices that pathogens employ to tune and tame the immune system using costimulatory molecules. Studying host–pathogen interaction in context with costimulatory signals may unveil the molecular mechanism that will help in understanding the survival/death of the pathogens. Hence, the very same pathways can potentially be exploited to develop immunotherapeutic strategies to eliminate intracellular pathogens.

The organisms mentioned above are representatives of successful pathogens from the class of bacteria, virus, and parasites. It appears that these pathogens adopt a common evolutionarily convergent mechanism to evade host immune reaction. In this chapter we will discuss the mechanisms by which pathogens suppress host immunity by modulating the expression of costimulatory molecules. [Table 3.2](#) shows the modulation of costimulatory molecules by intracellular pathogens. We therefore, in this chapter, suggest avenues of therapeutic intervention by exploiting costimulatory pathways for treating infections caused by pathogens.

Costimulation Biology Modulation by Pathogens

Costimulatory molecules of CD80/CD28, tumor necrosis factor (TNF)/TNFR, and T-cell immunoglobulin mucin (TIM) superfamilies have revealed a plethora of the possible ligand–receptor interactions that has elaborated the understanding of regulatory mechanisms of the immune responses mediated by APCs and T cells. For example, a positive regulator like CD40L (on T cells) when associated with CD40 (on APCs), not only activates T cells but also results in the activation of DCs; a process that is popularly called “T-cell licensing.”^{17,18} Similarly, ligation of CD28 with CD80 and CD86 is known to induce the secretion of interleukin-6 (IL-6) and interferon- γ (IFN- γ) by DCs and activation, proliferation, and differentiation of B cells.^{19,20} It is reported that 4-1BBL expressed on DCs, binds to 4-1BB on T cells, to bolster DCs help to T cells.²¹ Many reports have highlighted the inhibitory roles of cytotoxic T lymphocyte-associated antigen (CTLA-4; CD152) and programmed death (PD)-1 (expressed on T cells) with ligands CD80/CD86 and PDL-1/PDL-2 (on APCs), respectively.^{22,23} It clearly suggests that costimulation not only amplifies the magnitude of the activation of T cells and APCs, but fine tunes the immune response as well, thereby controlling the hyper activation.

TABLE 3.2 Modulation of Costimulatory Molecules by Various Intracellular Pathogens such as Protozoa, Bacteria, and Viruses

Intracellular Pathogen Type	Name of the Intracellular Pathogen	Costimulatory Molecule Upregulated/Downregulated	Loss of Function
Viruses	HIV	PD-1 ↑, PDL-1 ↑, CTLA-4 ↑, CD80 ↓, CD86 ↓, CD33 ↓, CD40 ↓, 4-1BB ↓, OX40 ↓	Blocks IL-2 but augments IL-10 secretion, induces exhaustion of T cells, defective CTLs, respectively, and hampers antigen uptake ability of APCs
	HBV	PDL-1 ↑, PD-1 ↑	Induces IL-10 secretion, enhances apoptosis and anergy in T cells
	HCV	CD83 ↓, CD86 ↓	Reduces stimulatory capacity of DCs
	Measles	CD40 ↓, CD80 ↓, CD86 ↓, CD25 ↓, CD83 ↓, CD69 ↓	Abnormal DCs differentiation improper CD8 T-cell proliferation
	Herpes simplex virus	ICAM-1 ↓	Blocks APCs T-cell communication
Protozoa	<i>L. donovani</i>	CD80 ↓	Inefficient T-cell response
	<i>T. gondii</i>	CD80 ↓	Inhibits T-cell stimulatory activity
Bacteria	<i>Mtb</i>	CD80 ↓, CD86 ↓, CD40 ↓, PDL-1/PDL-2 ↑, PD-1 ↑	Hampers effective T-cell activation; induces anergy or apoptosis in T cells, paralyzes IL-2 and chemokines secretion, and inhibits NK cell function
	<i>M. leprae</i>	CD80 ↓, CD28 ↓	Blockade of IL-12 secretion, defective T-cell response
	<i>S. typhimurium</i>	ICAM-1 ↓	Impedes antigen uptake ability of APCs
	<i>B. anthracis</i>	CD40 ↓, CD80 ↓, CD86 ↓	Impairment of antigen-specific B- and T-cell immunity, suppresses the function of DCs
	<i>H. pylori</i>	PDL-1 ↑, CTLA-4 ↑	Exhaustion of DCs, obstructs cytokines secretion, induces anergy in T cells
	<i>B. bronchiseptica</i>	CD40 ↓	Hampers maturation of DCs
	<i>B. pertussis</i>	CD40 ↓, ICAM ↓	Promotes differentiation of Tregs

Note: Symbol (↑) means upregulation or enhancement in the expression while symbol (↓) means downregulation or suppression of expression.

T-cell interaction with APC involving TCR and costimulatory molecules activates a plethora of downstream signaling molecules, leading to the induction of the expression of CD40L, PD-1, and CD28.²⁴

Intracellular pathogens utilize an array of mechanisms to manipulate costimulatory molecules. Upon infection, pathogens trigger IL-10 secretion by inhibiting p38 mitogen-activated protein kinases (MAPKs) and promoting ERK phosphorylation.²⁵ IL-10 blocks the degradation of I κ B- α which inhibits the NF- κ B activation eventually inhibiting the expression of costimulatory molecules.²⁶ Interference in Toll-Like Receptor (TLRs) signaling by the intracellular pathogens is considered to be a foremost event in the suppression of costimulatory molecules. Mannosylated lipoarabinomannan (ManLAM) of *Mtb* binds to DC-SIGN and compromises the lipopolysaccharide (LPS)-induced activation of DCs by interfering in the TLRs' signaling.²⁷ Binding of *Mtb* early secreted antigenic target protein 6 (ESAT-6) to TLR-2 activates AKT and prevents interaction between the MyD88 and IRAK4, consequently abrogating NF- κ B activation that suppresses CD80 expression.²⁸ Similarly, HIV upregulates PDL-1 on DCs and monocytes by attenuating the signaling of TLR-7 and TLR-8.²⁹ In addition, HIV-mediated PI3K activation upregulates PDL-1 on APCs and thus suppresses the activation of HIV-specific CD8+ T cells.³⁰ Rapid endocytosis of costimulatory molecules upon infection is suggested as one of the mechanisms to interrupt the function of APCs. For instance, Nef protein of HIV modulates the actin-dependent trafficking mechanism to remove CD80/CD86 from the monocytes surface, making them inefficient to activate T cells.³¹

TCR signaling is central for the optimum expression of costimulatory molecules and effector function of T cells. Therefore, it may serve as an important target for pathogens to paralyze T-cell activity.³² *Mtb* glycolipids interfere with TCR signaling and block the activation of CD4+ T cells.³² HIV gp120 interacts with CD4 (a coreceptor for TCR) that inhibits intracellular signal transduction through TCR, leading to decreased hydrolysis of polyphosphoinositide (PI), Ca²⁺ influx, activation of protein kinase C (PKC), and eventually failure of nuclear factor of activated T cells (NFAT) translocation. These events culminate in the inhibition of CD40L expression. Decline of CD40L on T cells abrogates the bidirectional signaling and reduces the exhibition of CD80 on APCs.³³ In contrast, Nef activates NFAT and stimulates IL-2 release to overcome the exogenous requirement of IL-2 to promote T-cell proliferation, consequently disseminating the infection.³⁴ However, Nef retards CD28 and CD4 expression, thereby making T cells incapable of promoting cell-mediated immunity (CMI).³⁵ Corroborative with these observations, Nef-mutated HIV is unable to cause persistent infection. Similarly, core protein of HCV inhibits TCR signaling and upregulates PD-1.³⁶ The above-mentioned mechanisms imply that pathogens can effectively utilize various costimulatory pathways to subvert immune response to persist in the host.

MODULATION OF COSTIMULATORY MOLECULES BY VIRUSES

The vertebrate immune system has evolved complex antiviral innate and adaptive immune mechanisms like production of IFNs which limit the viral replication by the host cells. Other wonderful examples involve the natural killer (NK) cell-mediated lysis and apoptosis of the infected cells^{37,38} to counter viral infestation. Adaptive antiviral immunity relies greatly on the lysis of infected cells by cytotoxic CD8+ T cells and neutralizing antibodies secreted by B cells. HIV is a retrovirus that is responsible for 1.9 million deaths annually. It predominantly infects CD4+ T helper (Th) cells. Further, it invades DCs, monocytes, and macrophages expressing CD4 and one of its coreceptors CCR5 or CXCR4. HIV can also bind to DC-SIGN and mannose receptors.^{39,40} However, the virus preferentially replicates in the activated HIV-specific CD4+ T cells. Indeed, excessive loss of CD4+ T cells is the hallmark of HIV infection.

Like many other intracellular pathogens, HIV efficiently exploits the costimulatory molecules to override the immune responses. Its infection is associated with decreased expression of CD40L on CD4+ T cells.⁴¹ Upon activation, CD4+ T cells from individuals with progressive disease show very little upregulation of CD40L, which corroborates with their inability to help APCs and failure to induce IL-12 in DCs.⁴² Further, CD40–CD40L interaction has been demonstrated to be important in engendering a robust HIV-specific CD8+ T-cell response. Furthermore, HIV interferes in the CD40 signaling pathway in B cells and hinders T cell help, thereby impairing the secretion of immunoglobulin G (IgG) and IgA antibodies.⁴³ AIDS patients suffer from a defective humoral immunity (HI), which may be due to loss of T-cell function or B-cell intrinsic defects.⁴⁴ HIV upregulates Fas and FasL (members of TNF superfamily) on CD8+ T cells and APCs, respectively, which leads to the apoptosis of the interacting CD8+ T cells.^{45,46} Thus, in HIV infections, the hunter becomes the hunted! During viral infections, continual expression of CD80/CD86 on DCs is decisive to maintain the effector function of CD8+ T cells.⁴⁷ Intriguingly, HIV downregulates the expression of CD80/CD86 and their ligand CD28 on infected APCs and T cells, respectively. The expression of costimulatory molecules such as 4-1BBL, CD70, OX40, and OX40L is affected during HIV-1 infection.⁴⁸ Measles, herpes, and hepatitis C viruses (HCVs) retard the expression of CD80, CD86, CD25, CD83, and CD40 that leads to poor CD8+ T-cell priming.^{49,50} In addition, herpes virus suppresses intracellular cell adhesion molecule (ICAM)-1 on APCs, thereby obstructing immunological synapse with T cells.⁵¹

Chronic viral infections are associated with loss of function in T cells, a phenomenon popularly known as T-cell exhaustion. Exhausted T cells highly express PD-1 and have poor effector function.⁵² PD-1 expression on HIV-specific T cells is associated with T-cell exhaustion and disease progression.⁵³ PD-1 is known to make CD8+ T cells more susceptible to Fas-mediated

lysis.⁵⁴ Signaling through PD-1 can suppress IL-2 secretion by CD8+ T cells. IL-2 is known to rescue T cells from anergy and boost the memory response.⁵⁵ In addition, upregulation of PDL-1 on APCs during HIV or hepatitis B virus (HBV) infection supports the survival of pathogens⁵⁶ (Table 3.2). In essence, viruses can exploit costimulatory molecules in restraining the function of both T cells and APCs.

MODULATION OF COSTIMULATORY MOLECULES BY BACTERIA

The bacterial intracellular pathogens like *Salmonella typhi* (*S. typhi*), *Helicobacter pylori* (*H. pylori*), *Mtb*, *Mycobacteria avium*, *M. leprae*, etc., infect both DCs and macrophages. These APCs have pathogen recognition receptors (PRRs) on their surface which recognize bacterial components known as pathogen-associated molecular patterns (PAMPs) thus triggering innate immunity that initiates antimicrobial defense mechanisms involving autophagy, apoptosis, release of antimicrobial compounds like IFN- γ and TNF- α , etc.⁵⁷ It is followed by activation of adaptive immunity which is highly specific and has immunological memory. The adaptive immune response involves the clonal selection of B cells and T cells that confer HI and CMI to pathogens, respectively (Figure 3.3). Pathogen-specific Th1 cells release cytokines, especially IFN- γ and TNF- α , which plays an imperative role in activating infected macrophages and restraining the microbial growth.⁵⁸ Moreover the interaction of costimulatory molecule ligand receptor between CD80/CD86 with CD28 is essential for preventing apoptosis and anergy in T cells. Further, interaction of CD40/CD40L costimulatory receptor ligand results in efficient IL-12 production together with an upregulation of costimulatory molecules, in addition to enhanced antigen presentation by APCs thereby leading to an enhanced T-cell response.^{18,59} Hence, interference in the downmodulation of any of these pathways by the pathogens would be detrimental to the host. *Mtb* is one of the most successful pathogens in the history of mankind.

There are numerous reports indicating the role of mycobacteria in downregulating the expression of CD80, CD86, and CD40 on APCs.⁶⁰ An elegant study demonstrated that the expression of costimulatory molecules and MHC are downregulated in macrophages infected with fluorescent reporter bacteria.⁶¹ A recent study showed, albeit for BCG, that MHC-II, CD80, CD86, and CD40 are downregulated during chronic phase of infection. In such studies, it is important to delineate the expression of these molecules on infected versus noninfected cells, as by-stander inflammation could interfere in the interpretation of results. Defect caused by bacterial infections in APCs in the CD80/CD86 (downregulation) and CTLA-4 (upregulation) signaling pathway in APC and T cells is known to induce anergy/apoptosis of interacting T cells, and to paralyze the release of IL-2, which may compromise the generation of T-cell memory.^{62,63} Impediment in CD28 signaling interferes in IFN- γ production

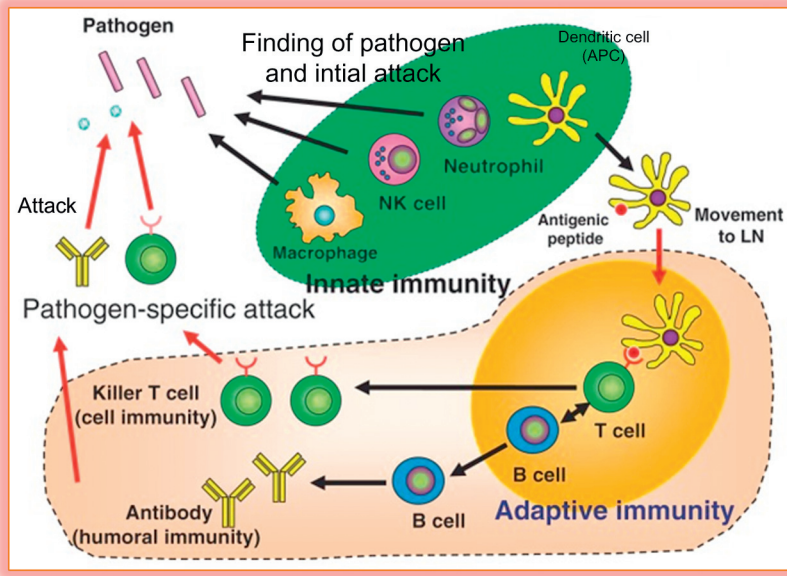


FIGURE 3.3 CD80 and CD86 deliver bidirectional signaling for the activation of T cell and APC. Cognate association between T cell and APC lead to interaction between MHC–peptide with TCR, and CD80/86 with CD28/CD152. This steers toward the activation and differentiation of T cell. Activated T cell secretes cytokines that ultimately help in the activation and differentiation of APC. The arrows shown in the diagram depict that there occur as well an alternative pathway of bidirectional signaling through costimulatory molecules CD80/CD86 and CD28/CD152 that activates both T cell and APC.

and hence promotes the survival of pathogens. *M. leprae* has been shown to obstruct the CD28/B7 signaling pathway for rendering antigen-specific T cell unresponsive in lepromatous leprosy patients. Protective CMI is always associated with the release of chemokines and migration of immune cells to the site of infection. *Mtb* impairs the secretion of chemokines by interfering with the CD28-CD80/CD86 signaling pathway thereby obstructing the surveillance of immune cells and enhancing the propagation of the bacterium.^{64,65} Recently, the importance of CD80/CD86 in controlling mycobacterial infection has been demonstrated in CD80/CD86 double knockout mice.⁶⁶ The down-modulation of CD80/CD86 in chronic phase of the infection suggests that mycobacteria may actively exploit this pathway to anergize the T cells. It has been shown that the most abundant cell wall lipid trehalose 6,6'-dimycolate of *Mtb* and MTSA-10 inhibit the expression of costimulatory molecules on the surface of the macrophages.^{67,68} Similarly, CD40–CD40L interaction is very important in mediating efficient protection against mycobacteria.⁶⁹ Indeed, it has been shown *in vitro* and in lepromatous patients that CD40 is downregulated by

M. leprae.⁷⁰ There is indirect evidence indicating the involvement of *Mtb* manipulating CD40/CD40L expression. In humans, CD40L expression on Th1 cells of tuberculosis patients has been correlated with the intensity of IFN- γ secretion. However, CD40 but not CD40L knockout mice are susceptible to *Mtb* infection. It is reported that the heat shock protein 70 (HSP70) of *Mtb* acts as an alternate ligand for CD40. Corroboratively, overexpression of HSP70 interferes with long-time persistence of *Mtb* and allows its clearance.⁷¹ Interestingly, in chronic mycobacterial infections, CD40 is suppressed on infected cells.^{61,72} Therefore, it is intriguing to speculate that in the chronic phase of infection, mycobacteria may hamper CD40 expression or manipulate CD40L signaling through binding with HSP70 instead of CD40L in order to evade host defense mechanisms.

The expression of PD-1 and its ligands PDL-1/PDL-2 is upregulated by the mycobacteria.⁷³ Blockade of PD-1 interaction with PDL-1/PDL-2 enhances immune response. Several studies have shown that T cells and NK cells from tuberculosis patients have increased PD-1 expression.^{74,75} These observations suggest that mycobacteria may exploit PD-1 and PDL-1/PDL-2 pathways to dampen the host immune responses. In contrast, some studies have suggested that these pathways may be involved in controlling exuberant T-cell responses in tuberculosis and hence may be beneficial for both the host and the pathogen.^{76,77} *Mtb* retards the development and maturation of monocyte-derived DCs to limit the immune response.⁷⁸ These immature DCs have a tolerogenic effect *in vivo* that can result in the generation of regulatory T cells (Tregs). Tregs can restrain the proliferation of naive and memory T cells, thereby suppressing preexisting T-cell immunity.^{79,80}

Not only mycobacterial species but many other bacterial pathogens can exploit CD80/CD86-CD28/CTLA-4 pathways for their persistence (modulation of costimulatory pathways; [Table 3.2](#)). *S. typhi* is known to suppress ICAM-1 and as a consequence reduces the antigen uptake by APCs and inadequate T-cell response.^{81,82} *Yersinia pseudotuberculosis* decreases CD86 expression on B cells and impedes the function of both B and T cells.⁸³ *H. pylori* causes chronic infection in the gut resulting in peptic ulcers. Further, it is known to induce the expression of CTLA-4, resulting in the anergy of T cells and poor clearance of the bacteria.⁸⁴ *H. pylori* diminishes the expression of CD40L on T cells and therefore employs CD40/CD40L pathway for its survival. Furthermore, it upregulates PDL-1 expression on gastric epithelial cells and inhibits the activation of T cells recruited to gastric mucosa.⁸⁵ In addition, it has been reported that PDL-1 upregulation not only blocks T-cell proliferation and IL-2 secretion, but also promotes the development of Tregs.⁸⁶ *Bordetella pertussis* and *B. bronchiseptica* decrease the manifestation of CD40 and ICAM-1 on DCs, and subsequently, render DCs tolerogenic and promote chiefly Tregs but not Th1 cells.⁸⁷ Hence, costimulatory molecules could serve as attractive targets for bacteria to modulate in order to prolong their own survival.

MODULATION OF COSTIMULATORY MOLECULES BY PROTOZOAN PARASITES

T cells play a significant role in controlling the protozoa-inflicted diseases like malaria, visceral leishmaniasis, and trypanosomiasis. Humoral responses play a less important role. However, complement-mediated killing or opsonization is responsible for control of pathogen during the intermittent extracellular phase of its replication cycle within the vertebrate host.⁸⁸ Parasites like *Leishmania donovani*, *L. chagasi*, *Toxoplasma gondii* (*T. gondii*), *T. cruzi*, and *Plasmodium falciparum* can manipulate the costimulatory molecules for evading immune system. *L. donovani* and *L. chagasi* infect macrophages and are reported to downregulate both CD80 and ICAM-1⁸⁹ Such APCs fail to optimally activate T cells. *T. gondii* and *T. cruzi*, selectively dampen the exhibition of CD80 and CD28, respectively to impair the function of T cells.⁸⁹ CD40 deficient mice succumb to plasmodium infection. *Plasmodium* interferes in the signaling mechanism of CD40 in DCs.⁹⁰ This signifies that this pathway is detrimental in providing sterilizing immunity to parasites, as it activates APCs and protective Th1 responses. CD40-CD40L signaling is crucial for Th1 immunity against *L. major*, because CD40L/CD40 knockout mice sparsely secrete IL-12, favoring Th2-biased response. Interestingly, *L. major* can differentially modulate the expression of CD40 and thus anergizes T cells and promotes Tregs population.⁹¹ In nutshell, parasites can efficiently enable the immune system by dampening the expression of costimulatory molecules.

COSTIMULATORY MOLECULES IN THERAPEUTIC VACCINATION

In order to develop successful vaccines against chronic viral diseases it is widely thought that a cell-mediated immune response is required to eliminate or control intracellular viruses. For a protective immune response it is important to induce long-term immunological memory. The initial activation of T cells requires both an antigen-specific signal, involving the recognition of a peptide–MHC complex by the TCR as well as additional costimulatory signals. Only when the T cell recognizes both the antigen and a costimulatory signal on the DCs is an immune response initiated. The idea of a therapeutic vaccine is that the natural immune response, although present, may be suboptimal. By providing antigens and costimulatory molecules together, one may be able to provide a strong enough response to eliminate, or at least better contain, the virus.

Coinhibitory pathways also have become therapeutic targets for combating chronic viral infections. Blockade of the PD-1/PD-L pathway identified the key role of inhibitory receptors in mediating T-cell exhaustion. Exhausted CD8+ T cells express multiple inhibitory receptors, including PD-1, lymphocyte-activation gene 3 (Lag-3), CD160, 2B4 (CD244), and CTLA-4. The coexpression of multiple inhibitory receptors has been associated with

greater T-cell exhaustion and more severe infection. Co-blockade of PD-1 and LAG-3 synergistically improved T-cell responses and reduced viral load, demonstrating that distinct inhibitory pathways can uniquely regulate T-cell exhaustion. Further studies are needed to define the unique and overlapping functions of coinhibitory pathways in exhausted T cells and to test the therapeutic benefits of targeting multiple coinhibitory pathways during chronic viral infection. Recent technological advances in structural biology provide new opportunities for understanding the structural and chemical features of costimulatory receptors and ligands, and their mechanisms of costimulation. The crystal structures of costimulatory receptors and ligands in the Ig, TNF, and TIM families, illustrate how structural understanding can provide important mechanistic insights through defining chemical and physical features that underlie receptor/ligand specificity, affinity, and oligomeric state. These approaches also may guide the development of improved therapeutic agents and reveal new therapeutic strategies for targeting costimulatory pathways.⁹²

We have recently shown that signals delivered through costimulatory molecules:

- i) not only are responsible for optimum activation of T cells but through bidirectional signaling can also influence macrophages;
- ii) can modulate the secretion of proinflammatory molecules;
- iii) can regress the growth of intracellular pathogens like *M. tuberculosis*, *M. microti*, etc.;
- iv) this novel strategy can be effectively exploited to develop immunotherapy either using humanized antibodies against CD80, CD86 and CD40 or CD28 fusogenic proteins for the treatment of intracellular pathogens like *M. tuberculosis*, *HIV*, *L. donovani*, *T. cruzi*, etc.;
- v) since this approach is based on modulating the immune system of the hosts rather than targeting the pathogen; hence it significantly diminishes chance of emergence of drug resistant strains of bacteria. It may be concluded from the results that macrophages are not only activated and kills intracellular pathogens by cytokines secreted by T cells but also through bidirectional costimulation through signals delivered by costimulatory molecules.^{92a,92b,92c}

Latest reports suggest that 4-1BBL can provide a costimulatory signal to the human CD28-T cells, leading to cytokine production, cell survival and the upregulation of molecules associated with cellular cytotoxicity.⁹³ Memory CD8+ T cells specific for chronic viral pathogens (HCV, HIV, cytomegalovirus, Epstein–Barr virus) are found among the CD28-T cell population in human blood.^{94,95} The proportion of T cells lacking CD28 is increased in people with chronic viral infections or other persistent conditions such as multiple myeloma. The observation that 4-1BBL can stimulate CD28-T cells makes it an attractive candidate for boosting immunity in chronic viral infection, where the lack of CD28 on the memory cells will

make this population of cells insensitive to B7 stimulation. As will be discussed below, stimulatory anti-4-1BB antibodies have been tested *in vivo* in mouse models. Provision of anti-4-1BB can systemically increase both anticancer and antiviral immunity.⁹⁶

Further work has shown that the systemic administration of stimulatory anti-4-1BB antibodies can correct the defect in CD28^{-/-} mice when a single dose is provided during the primary influenza infection. This results in full restoration of the memory T-cell response, suggesting that for CD8⁺ T cells, CD28 is only required for initial T-cell activation and not for recall responses. Conversely, a single dose of anti-4-1BB antibody corrects the defect in the CD8⁺ T cell recall response in 4-1BBL^{-/-} mice only when added at the time of viral challenge, arguing that the 4-1BB costimulatory signal is more important during recall responses (unpublished data from Mir and Agrewala et al.) Why the immune system switches from CD28 to 4-1BB as a costimulatory molecule during primary versus secondary responses is not clear. Interestingly, administration of anti-4-1BB (100 µg) during viral challenge results in a two fold increase in the recall response to influenza virus, even in wild-type mice. The physiological levels of 4-1BBL in mice appear to be very low, as the ligands are difficult to detect except after extensive stimulation of the cells *in vitro*.⁹⁷ This is consistent with the finding that the provision of extra 4-1BBL or stimulatory anti-4-1BB antibodies is immune stimulatory even in immunocompetent mice.⁹⁸ These findings suggest that 4-1BBL, rather than B7, may be the better choice of immune stimulatory agents to use in an immunotherapeutic regimen for antiviral immunity.

THERAPEUTIC VACCINATION USING COSTIMULATORY MOLECULES IN HIV

With chronic viral diseases such as HIV there may well have been a vigorous initial immune response to the pathogen, but the combination of viral escape variants and viral interference with the immune system results in the establishment of a chronic infection. One model for applying a therapeutic vaccine to the problem of HIV is to first use antiviral therapy to reduce viral load as much as possible. This would be followed by the interruption of antiviral therapy and provision of HIV antigens together with costimulatory molecules, in the hope that the enhanced immune response could eradicate the residual virus and prevent further erosion of the immune system. A key issue becomes how to deliver the antigen and costimulatory molecules. As has been demonstrated in mouse models of cancer or acute viral infection, systemic administration of stimulatory antibodies against receptors involved in immune triggering is one possible approach. However, the need to produce large amounts of protein product for therapy could be prohibitively expensive. On the other hand, the recent success of a recombinant protein

therapy for arthritis (soluble TNFR) suggests that such approaches might be feasible.⁹⁹ An alternative approach that may be cheaper to administer would be to use recombinant replication-defective viruses containing both viral epitopes and the genes encoding costimulatory molecules as a therapeutic vaccine. Replication-defective adenoviruses, modified canary pox, and other viral vectors are being developed for immunotherapy.^{100,101} Recombinant replication-defective adenoviruses are particularly attractive as there are two regions of their genome that can readily accommodate foreign DNA so that one could independently incorporate both antigens and costimulatory molecules. Furthermore, multiple serotypes of a virus should allow for more than one immunization.

Another approach being considered for HIV therapy is known as adoptive immunotherapy. In this approach, patient lymphocytes are obtained by leukaphoresis and stimulated *in vitro* before reinfusion into the same patient. In a recent example, Levine and colleagues used anti-CD3 (a component of the TCR) and anti-CD28 coated beads to stimulate HIV patient T cells in a nonspecific way. The T cells were then infused back into the patients and their CD4+ T cell counts were found to improve, at least temporarily.¹⁰² The advantage of this approach is that the stimulatory agents are not delivered systemically, which reduces toxicity concerns. This approach, using antigen-specific activation together with costimulatory molecules, such as 4-1BBL, could be used to generate useful T cells for reinfusion into patients. This approach could also use a replication-defective viral delivery vector containing HIV epitopes and costimulatory molecules delivered to the patient's APCs *ex vivo*. These would be used to activate the patients' T cells, and the activated T cells are subsequently reinfused, again avoiding systemic delivery of the virus. Although great strides have been made over the last few years in our understanding of T-cell activation and responses during infection, many challenges remain in the development of therapeutic vaccines for chronic viral infections. These advances suggest that we will see a number of new therapeutic vaccine approaches developed over the next few years.

B7 COSTIMULATORY MOLECULES IN INFECTIOUS DISEASES

Because microbial products can affect the expression of costimulatory molecules, these molecules participate in the pathogenesis of infectious diseases. They also participate in the pathogenesis of autoimmune diseases, the rejection of grafted organs, and graft-versus-host disease, and antagonists of these molecules can prevent or ameliorate these conditions. Although B7 molecules are important in the pathogenesis of graft rejection, graft-versus-host disease, infections, and autoimmune diseases; in addition, they participate in immunity against tumors,¹⁰³ but here we will focus on the role of B7 molecules in infectious diseases. The B7:CD28 costimulatory pathway participates in the pathogenesis of several bacterial, parasitic, and viral diseases.

Certain bacteria or their products can increase the expression of B7. One of these bacterial products, LPS, increases the expression of B7-1 and B7-2 by APCs.^{103,104} The incubation of macrophages with heat-killed *Listeria monocytogenes* also causes increased production of B7.¹⁰⁵ A series of proteins isolated from pathogenic strains of neisseria, *Neisseria meningitidis* and *N. gonorrhoeae*, upregulate the expression of B7-2, which may explain the adjuvant effect of these neisserial proteins in antibody production.¹⁰⁶ In contrast, *Mtb* deficient expression of costimulatory molecules and the intracellular protozoan *L. donovani* inhibit the expression of B7-1 molecules after they infect macrophages *in vitro*.¹⁰⁷ The toxic shock syndrome and food poisoning are serious side effects of the immune system's defense against extracellular bacteria. These complications are caused by a class of microbial products, called superantigens, that can activate all T cells that express a particular family of T-cell antigen-receptor genes. The immune response to certain bacterial superantigens, such as the staphylococcal enterotoxins and the related toxic shock syndrome toxin, also depends on costimulation by B7-1 and B7-2.¹⁰⁸ Only a few studies have addressed the role of the B7:CD28 pathway in viral infections. After infection with the HIV, the expression of CD28 is reduced in both CD4 and CD8 T cells.^{109,110} It is not known whether stimulation of the remaining CD28 molecules promotes or inhibits¹¹¹ viral infection. In mice carrying a targeted mutation in the CD28 gene, the response of helper T cells to infection with vesicular-stomatitis virus is diminished, but cytotoxic T-cell responses to infection with lymphocytic-choriomeningitis virus are normal.¹¹²

These findings suggest that the B7:CD28 pathway plays a more important part in viral infections in which viral clearance depends on antibody responses. Infection with the HBV may be one of these, but this issue has not yet been examined. Infection with *Leishmania major* perhaps best exemplifies the consequences of the dominance of a particular subgroup of helper T cells on the immune response. Strains of mice that use type-1 helper T cells to respond to *L. major* produce large amounts of IFN- γ , and thus resist the parasite. By contrast, strains of mice that use type-2 helper T cells, and consequently produce IL-4 and IL-10, succumb to the organism. CTLA-4-Ig, if administered early after infection, reduces the production of IL-4 and halts the progression of disease in susceptible BALB/c mice but has no effect in the genetically resistant C57BL/6 strain of mice. These different effects in susceptible and resistant mice indicate that the priming of type-2 helper T cells is more dependent on the B7:CD28 pathway than is the priming of type-1 cells,¹¹³ and provides evidence for the interpretation of analogous results in mice deficient in CD28. Infection with the trematode helminth *Schistosoma mansoni* may be an instance in which the levels of B7 expression on APCs correlate with, if they do not dictate, the severity of disease. *S. mansoni* antigens sensitize type-1 and type-2 helper T cells to form large granulomas in the liver around eggs of the parasite. However, the response of the type-1 helper T cells and the intensity

of granulomatous inflammation spontaneously subside after the acute stage of the infection. There is substantial expression of B7-2 in cells of granulomas during the acute schistosomal disease; it rapidly diminishes thereafter.¹¹⁴ It is likely that the downregulation of B7 expression and of the granulomatous reaction are precipitated by IL-10 given that macrophages from egg granulomas secrete IL-10 and induce unresponsiveness in type-1 helper T cells.¹¹⁵ Neutralization of IL-10 by specific antibodies increases the expression of B7-2 by these macrophages and thus restores costimulatory activity.¹¹⁶ A similar process has been observed in humans infected with *S. haematobium*. Taken together, these findings suggest that the inhibition of B7 expression may permit therapeutic control of schistosomal egg granulomas. In fact, injection of IL-10 at a time when its endogenous expression is low or absent results in decreased formation of granulomas and decreased cytokine production by type-1 helper T cells.¹¹⁷ The observation that, as compared with type-2 effector CD4 T cells, type-1 effectors depend more on costimulation and are more susceptible to anergy has implications for chronic infections in which immune defenses depend on type-1 helper T cells. Very recently we have shown that macrophages can be activated not only through classical and alternate pathways but also by costimulation. Upon costimulation macrophages produce nitric oxide and proinflammatory cytokines and also upregulate the cell surface molecules thereby showing a state of activation. These costimulated macrophages effectively inhibit the growth of *Mtb* see [Figure 3.4](#). In tuberculosis, leprosy, schistosomiasis, filariasis, and leishmaniasis, the waning of the B7:CD28 costimulatory pathway may change the clinical form of the disease by depressing CMI and inducing T-cell anergy. To what extent this downregulation of T-cell immunity is caused by the organisms themselves or mediated by cytokines is not known.

COSTIMULATORY MOLECULES IN ALLERGIES AND ASTHMA

The prevalence of allergy is increasing worldwide, especially in Western countries.^{118,119} This trend may be due to increased hygiene which prevents early exposure to microorganisms in infancy and increases the risk of developing an allergic disease in the future.¹²⁰ Air pollution may also favor the development of allergies.¹²¹ Among allergic diseases, asthma is of particular concern as it can evolve into a life-threatening chronic pulmonary disease. It is now recognized that inhalation of aeroallergens by sensitized patients initiates a massive Th2 cell response resulting in the production of IL-4, IL-5, and IL-13 by CD4+ T cells.¹²² This triggers the development of typical signs of bronchial asthma including airway hyperresponsiveness, bronchoconstriction, mucus secretion, and airway remodeling. The recruitment, differentiation, and activation of Th2 cells play a cardinal role in the development of allergic disease. This Th2 polarization is mainly controlled by APCs that provide signals capable of supporting the differentiation of naive T cells into Th2 cells. Initially,

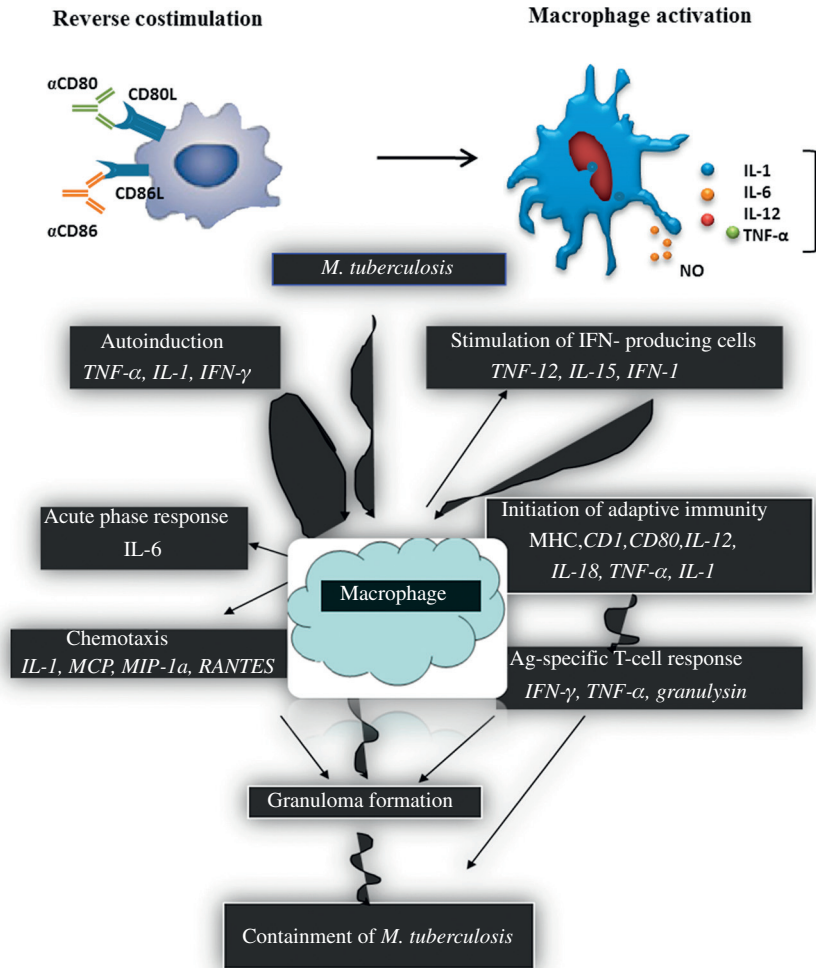


FIGURE 3.4 Activation of macrophages through costimulation to enhance the production of cytokines and nitric oxide. Macrophages are activated through reverse costimulation through anti-CD80 and anti-CD86 to enhance the production of cytokines and nitric oxide, both of which play an important role in protection against pathogens (*Mtb*).

Lafferty et al.¹²³ suggested that T lymphocytes require two coordinate signals to be fully activated. The first signal confers specificity to the immune response and plays an important role in the recognition of the MHC class–peptide complex at the surface of APCs by T lymphocytes. However, this primary signal is not sufficient to completely activate these T cells. To become fully effective, a second, nonspecific costimulatory signal is often required by T cells. These signals are provided by interactions between APCs

and T cells through surface molecules expressed on T lymphocytes. APCs express costimulatory molecules such as CD80 (B7-1) and CD86 (B7-2) which belong to the B7 family. They provide modulation of T-cell function by ligation to their receptors, CD28 or CTLA-4. Engagement of CD28 or CTLA-4 has been observed to have opposite effects. CD28 promotes T-cell activation and survival while CTLA-4 inhibits T-cell responses and regulates peripheral T-cell tolerance.¹²⁴ This demonstration provides clear evidence that costimulatory molecules are involved in the fine-tuning of the T-cell response by mediating both stimulatory as well as inhibitory signals. In many recent studies, numerous new costimulatory molecules have been described leading to the recognition that costimulation pathways are more complex than the classical two-signal model. In general, costimulatory molecules are divided into two main families: molecules from the B7:CD28 family, such as CTLA-4 or PD-L1, and from the tumor necrosis factor receptor (TNFR) superfamily such as OX40 or CD27.¹²⁵ All these costimulatory molecules have particular effects on T-cell activation, function, and survival and are implicated in nearly all inflammatory diseases. Studies to better characterize the specific role of these molecules in allergy and asthma are still ongoing. In this chapter, we summarize current knowledge concerning the role of costimulatory molecules in allergy and analyze the potential functions of the emerging new subsets of costimulatory molecules recently described.

Involvement of the B7:CD28 Family Molecules in the Regulation of Allergic Diseases

The first costimulatory molecules described were the ligands of CD28: CD80 (B7-1) and CD86 (B7-2). The CD28 costimulation pathway is an important factor for the promotion of an effective antigen-specific immune response. However, CD28-deficient mice are still able to develop allergic airway inflammation showing that CD28 cannot be solely responsible for the development of an allergic response.¹²⁶ The expression of CD28 ligands (CD80 and CD86) has been extensively studied in clinical samples from asthmatic patients. Hofer et al.^{127a} reported that B lymphocytes from asthmatic patients exposed to allergens express higher levels of surface CD86, but not CD80, compared with those from asthmatic patients not exposed to allergen or with those from healthy individuals (Table 3.3). Another study demonstrated that CD80 and, to a lesser extent, CD86 were upregulated at the surface of alveolar macrophages from allergic patients compared with those from pulmonary sarcoidosis, extrinsic allergic alveolitis patients, or from normal subjects.^{127b} In contrast, Burastero et al.¹²⁸ observed that in allergic individuals, CD80, but not CD86, is highly expressed by alveolar macrophages. CD86 can also be expressed in a soluble form, where the transmembrane domain is deleted. This form is mostly produced by circulating monocytes and, like membrane-bound CD86, crosslinks CD28 or

TABLE 3.3 Important Roles Played by Various Costimulatory Molecules in Allergic Diseases and Asthma

Costimulatory Molecule Family	APC	Effector Cell	Functions and Characteristics
B7 Family	CD80	CTLA-4	Contributes to the suppressive activity of allergen-specific regulatory cells during sensitization
	CD86		Polymorphism in Ctla-4 promoter and gene favors allergic diseases
		CD28	CD80 and CD86 expression is upregulated on the surface of various cells of allergic patients. Soluble CD86 is increased in the sera of allergic patients
	ICOS-L	ICOS	Regulates Th2-effector cell function and their infiltration in the lungs, production of Th2 cytokine, promotes B-cell differentiation and IgE production
			Expression on iNKT cells contributes to airway hyperreactivity
			Contributes to the differentiation of regulatory cells in pulmonary lymph nodes
	PD-L1	PD-1	Drives the differentiation of Foxp3+ CD4+ T cells
			Downregulates contact hypersensitivity reaction
	PD-L2	PD-1	Regulates asthma by an IFN- γ -dependent mechanism
			Downregulates airway hyperreactivity, prevents eosinophil infiltration in the lungs, and prevents IgE production
B7-H3	Receptor unknown	Promotes Th2 differentiation, eosinophil infiltration, and development of airway hyperreactivity	
		Decreases the severity of allergic conjunctivitis	

TNFR Family	OX40L	OX40L	Promotes the development of Th2 cells
			Prevents differentiation of Treg cells
			Abrogates mast cell degranulation
	CD30-L	CD30	CD30 is expressed by Langerhans cells, CD4+ and T cells of atopic patients
			Soluble CD30 is increased in the sera of allergic patients
	4-1BB-L	4-1BB	Promotes airway hyperreactivity, eosinophil infiltration, and IgE production
			Upregulates Th2 cell proliferation and mast cell cytokine production
	Fas	FasL	Delays resolution of airway hyperresponsiveness
			Promotes eosinophil apoptosis in the lungs
	CD27	CD70	Increases production of IgE by B cells
	CD40	CD40L	Contributes to isotype class switching toward IgE
			Enhances development of airway inflammation
Increases production of Th2 cytokine and decreases number of Treg cells			
Others	CD2	CD58	Promotes differentiation of Th2 cells and the production of IgE
			Expressed on monocytes of allergic patients

CTLA-4 and activates T lymphocytes.¹²⁹ In patients with acute asthma, the level of soluble CD86 has been shown to be increased relative to that in patients with stable asthma or in healthy individuals. Monocytes from allergic patients produce more soluble CD86 compared with those of healthy individuals.¹³⁰ It was also observed that the level of soluble CD86 is correlated with the severity of the airway hyperreactivity (AHR). These results are consistent with studies which show that the concentrations of soluble CD80 and soluble CD86 are elevated in asthmatic patients. Interestingly, the administration of a glucocorticoid (e.g., prednisolone), used to reduce airway inflammation in allergic patients, reduces the level of circulating CD86.¹³¹ Recently, Ritprajak et al.¹³² demonstrated that the topical administration of a silencer RNA specific to the CD86 gene reduced local inflammation in a mouse model of atopic dermatitis by decreasing the recruitment of DCs into the skin, production of antigen-specific IL-4, and induction of serum IgE and IgG1. It has also been reported that pulmonary tolerogenic DCs stimulated by allergen exposure express high levels of both CD80 and CD86.^{133,134} Taken together, these studies suggest that overexpression of CD80 and/or CD86 is correlated with the development of allergic disease and asthma. Within the B7 family, CD28 and inducible costimulator (ICOS) are most homologous with respect to structure and function. Both are type I transmembrane receptors expressed as homodimers, with an extracellular (Ig)V-like domain, a hallmark of receptors of the B7-related family. ICOS expression is induced *in vitro* within 24–48 h of activation on all Th-primed cells.¹³⁵ ICOS has been shown to regulate the production of Th2 cytokines¹³⁶ and plays a critical role in lung mucosal inflammatory responses.¹³⁷ Further, this costimulatory molecule has been suggested not only to intensify some of the functions of CD28 during an already established immune response but also to induce additional T-effector cell functions.¹³⁸ Recent studies suggest that ICOS-mediated costimulation may regulate Th2-effector cell function without affecting Th2 differentiation. Another study showed that transfer of ICOS-enriched T cells followed by allergen airway challenge-induced infiltration of recipient T and B cells as well as local production of allergen-specific IgE by intrapulmonary plasma cells.¹³⁹ In contrast, transfer of the ICOS-depleted T-cell fraction resulted in the recruitment of significantly lower numbers of B cells with no local IgE production. These data indicate that expression of ICOS defines a subset of T-effector cells that are required for B-cell infiltration and local IgE production in lung tissue. According to Tesciuba et al.,¹³⁸ ICOS stimulation increases the migration of lymphocytes into draining lymph nodes by augmenting the expression of attractant chemokines CCL21 and CXCL13. In other reports, the increased production of IL-5, which is a main factor for the differentiation, maturation, and recruitment of eosinophils, is attributed to ICOS+ cells. ICOS+ T cells also promote the differentiation of B cells and IgE-producing plasma cells through the enhanced production of IL-4 and IL-10. ICOS-deficient mice are unable

to induce high IgE responses which demonstrates their role in the induction of IgE production. Some studies also suggest that intermediate ICOS expression is associated with high production of Th2 cytokines, whereas high levels of ICOS predominantly translate into high IL-10 production. Surprisingly, ICOS/ICOS-L interaction not only promotes the development of Th2-driven inflammation but also mediates mucosal tolerance, as studies have indicated that pulmonary DCs in the bronchial lymph nodes of mice exposed to respiratory allergen induced the costimulation of Tregs via the ICOS/ICOS-L pathway.^{133,134,140,141} These Treg cells produce IL-10, show inhibitory activity and, when adoptively transferred into sensitized mice, have the ability to inhibit the development of AHR. These reports suggest that both the development and inhibitory function of Treg cells depend upon the presence of IL-10 and ICOS/ICOS-L interaction.

CTLA-4 has been described to be an important regulator of T-cell activation. CTLA-4 is constitutively and exclusively expressed by T lymphocytes in both mice and humans.¹²⁴ CTLA-4 expression confers to T-lymphocyte regulatory functions. Indeed, the contribution of CTLA-4 in the regulation of the immune system is demonstrated by the development of multiple organ autoimmune pathologies and lymphoproliferative disease in CTLA-4-deficient mice. Blockade of CTLA-4 activity abolishes the suppressive function of CD4+ CD25+ T cells.¹⁴² In a mouse model of inflammatory bowel disease, the effect of transfer of a population of CD4+ CD45RB low Treg cells in decreasing intestinal inflammation is abrogated by the co-administration of a blocking anti-CTLA-4 antibody. These studies suggest that the engagement of CTLA-4 at the surface of Treg cells by its ligands CD80 or CD86 contributes to the regulation of suppressive functions of Treg cells. Polymorphism in CTLA-4 gene is also considered a risk factor for allergy and asthma. Howard et al.^{143a} characterized four single nucleotide polymorphisms which were related to allergic and asthma phenotypes. They demonstrated that these specific polymorphisms alone or in combinations are correlated with an elevated IgE titer or bronchial hyperresponsiveness in patients with asthma. Interestingly, in the same study, no correlation between allergic phenotype and single nucleotide polymorphisms for CD28 was observed. These findings confirm that CTLA-4 is indeed involved in the course of allergic diseases. Interestingly, CTLA-4 seems to play a more important role in the sensitization phase than in established allergy.

REGULATION OF THE IMMUNE RESPONSES TO PATHOGENS THROUGH COSTIMULATORY MOLECULES

Immune response to pathogens is regulated by stimulatory and inhibitory costimulatory signals. Positive costimulation is critical for the development of T-cell immune responses against foreign pathogens, while negative

regulation is critical for the termination of immune responses, for peripheral tolerance and to avoid inflammation-induced tissue damage.^{143b–145} Up to now four different families of costimulatory and coinhibitory molecules have been identified which are able to modulate TCR signaling in APC–T-cell interaction. These families include (1) B7-CD28 family including CD28, cytotoxic T-lymphocyte antigen-4 (CTLA-4, CD152), PD-1 (CD279), inducible costimulatory molecule (ICOS, CD278), and B- and T-lymphocyte attenuator (BTLA, CD272); (2) TNFR superfamily including CD27; (3) CD2/ signaling lymphocyte activation molecule (SLAM) family including SLAM (CD150), 2B4 (CD244), and CD48; and (4) Ig family including TIM-3, CD160, and Lag-3 (Figure 3.5).

In the presence of a pathogen, a specific and effective immune response must be induced and naive T cells undergo activation upon encounter with their specific antigens.^{146,147} This leads to antigen-specific T-cell proliferation,^{148,149} cytokines production, and induction of T-cell differentiation toward an effector phenotype¹⁵⁰ combined to survival signals.^{151,152} After clearance or control of the pathogen, the immune response must be terminated in order to avoid tissue damage and chronic inflammation.^{152,153} Two main mechanisms are involved in the contraction of the effector phase of immune responses, that is, either the inhibition of T-cell expansion¹⁵⁴ or the elimination of activated cells by apoptosis.¹⁵⁵ The latter is referred to as activation-induced cell death (AICD).^{155,156} Direct inhibition of T-cell proliferation is induced via signals through coinhibitory molecules such as CTLA-4 or PD-1, while 2B4 and SLAM are considered to be critical in the regulation of AICD.¹⁵⁷ The role of coinhibitory molecules in regulating the immune system is also evidenced by severe autoimmune and lymphoproliferative diseases resulting from the lack or aberrant expression of these molecules.¹⁵⁸

Expression of Coinhibitory Molecules on Effector T Cells

T cells play an important role in the defense against infectious agents. Upon recognition of their cognate antigen, naive T cells get activated and differentiate into effector cells.¹⁵⁹ This activation results in both phenotypic and functional changes that will determine the fate of effector T cells and the efficacy of the immune response.¹⁵⁰ Several studies have aimed to better define the profile of effector cells associated with the efficient control of infectious agents.¹⁶⁰ Viganò et al.^{161a} has recently performed and observed a comprehensive investigation of the expression of PD-1, 2B4, CD160, CTLA-4, TIM-3, Lag-3, and SLAM on CD4 and CD8 T-cell subsets identified according to their differentiation state (i.e., naive, memory, effector/memory). These analyses showed that whereas virtually none of the coinhibitory molecules tested was present on naive cells, they were present on memory T cells but at low levels, and more importantly, that effector/memory T cells expressed a significantly higher density of coinhibitory molecules simultaneously.

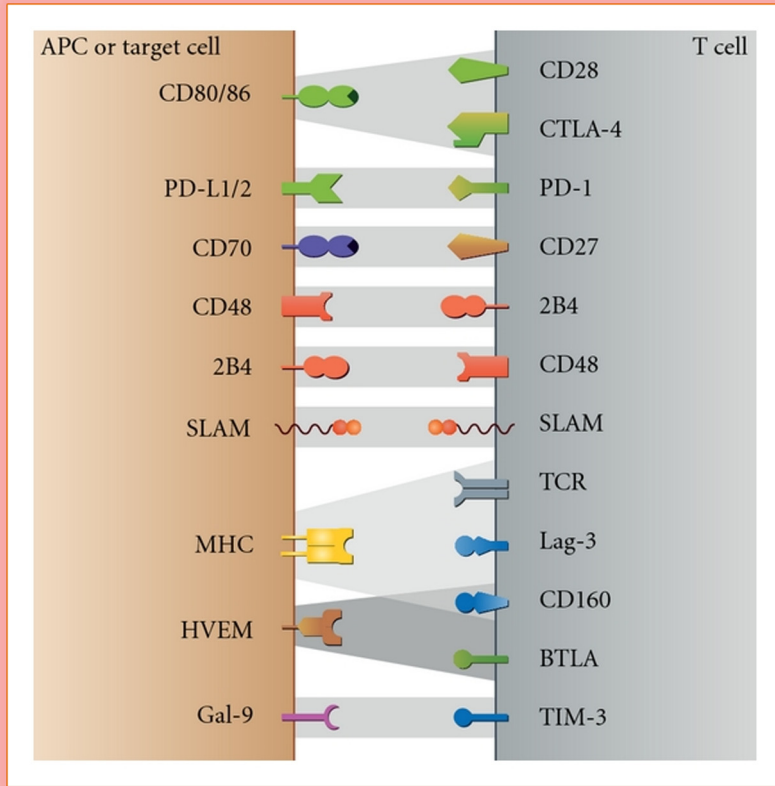


FIGURE 3.5 Regulatory molecules and their ligands. Schematic overview of the different costimulatory and coinhibitory molecules expressed by T cells (right panel) and association with their respective ligands expressed by APCs or target cells (left panel). Coinhibitory molecules are color coded according to their relevant families. The four families of regulatory molecules include (1) B7-CD28 family including CD28, cytotoxic T-lymphocyte antigen-4 (CTLA-4; CD152), programmed death-1 (PD-1; CD279), inducible costimulatory molecule (ICOS; CD278), and B- and T-lymphocyte attenuator (BTLA; CD272); (2) TNF-receptor superfamily including CD27 (3) CD2/signaling lymphocyte activation molecule (SLAM) family including SLAM (CD150), 2B4 (CD244), and CD48; and (4) Ig family including T-cell immunoglobulin mucin-3 (TIM-3), CD160, and lymphocyte-activation gene 3 (Lag-3).

ROLE OF COINHIBITORY MOLECULES IN THE CONTRACTION OF THE EFFECTOR PHASE OF IMMUNE RESPONSES

The immune system is able to mount strong and efficient immune responses against pathogens without damaging organs. This is notably achieved by the

induction of the contraction and termination of the immune response after control or elimination of the infectious agent. During the contraction, the majority of effector T cells die, while remaining cells survive as memory cells.^{152,153} The contraction of the immune response and the determination of T-cell fate depend on many transcription factors regulated during the course of the immune response.^{161b} These factors can be either induced or repressed by different signaling pathways provided, with different strength and kinetics, by costimulatory and coinhibitory molecules.^{162,163} One of the best-established mechanisms involved in the regulation of TCR signaling is the interaction between costimulatory (CD28) and coinhibitory (CTLA-4) molecules with CD80 or CD86 expressed by DCs. Crosslinking of CD28 on T cells synergizes with TCR signaling to induce activation. Conversely, crosslinking of CTLA-4 induces an inhibitory signal, which prevents T-cell activation.^{154,164} CTLA-4 is upregulated on activated T cells and, as a structural homologous to CD28 with higher affinity for CD80 or CD86, it competes with CD28 to inhibit TCR signaling.¹⁵⁴ PD-1 is another well-known regulatory molecule. It is expressed on activated CD4 and CD8 T cells, NKT, B cells, and activated monocytes,^{124,143b} and its expression is induced by TCR- and BCR-mediated signaling.¹⁶⁵ The two PD-1 ligands (i.e., PD-L1 and PD-L2) differ in their expression pattern.¹⁶⁶ PD-L1 (B7-H1, CD274) is expressed by a broad array of cells (e.g., vascular endothelial cells, epithelial cells, muscles cells, hepatocytes) whereas PD-L2 (B7-DC, CD273) expression is restricted to hematopoietic cell types (i.e., DC, macrophages, mast cells).¹⁶⁷ PD-1/PD-Ls pathway regulates the balance between stimulatory and inhibitory signals needed for effective immune responses against pathogens.^{168–171} Engagement of PD-1 by PD-L1 leads to the inhibition of CD28-mediated costimulation and thus of TCR-mediated lymphocyte proliferation and cytokines secretion. The relative levels of expression of inhibitory (PD-Ls) and stimulatory (CD80/CD86) ligands by APCs can determine the extent of T-cell activation while PD-L1 expression on nonlymphoid tissues may determine the extent of effector immune responses at sites of inflammation.¹⁶⁸ Also member of the B7-CD28 family, BTLA, is an inhibitory receptor able to recruit phosphatases to dampen TCR signaling¹⁷² through the interaction with herpes virus entry mediator (HVEM) expressed on naive T and B cells. HVEM-BTLA signaling was shown to limit T-cell activity *in vivo* and to negatively regulate homeostatic expansion of CD4 and CD8 T cells.¹⁷³ Finally, HVEM can also interact with CD160, resulting in an inhibitory signaling dampening T-cell activation.¹⁷⁴

Functional Exhaustion and Loss of Effector Functions

During chronic viral infections such as HIV and HCV, several inhibitory molecules are overexpressed on virus-specific CD4 and CD8 T cells, and this is associated with a state of functional deficiency also called functional

exhaustion. Exhaustion is characterized by the progressive loss of T-cell functionality, leading ultimately to the deletion of exhausted T cells. The loss of the distinct T-cell functions occurs sequentially.⁵² IL-2 production and T-cell proliferation potential are lost first. TNF- α production and cytotoxic capacity disappear later followed, ultimately, by the loss of IFN- γ production. Finally, deeply exhausted T cells are deleted via apoptosis.¹⁷⁵

The current hypothesis is that functional exhaustion occurs as a consequence of the attempt of the immune system to limit the magnitude of effector T-cell responses in order to safeguard against autoimmune responses and inflammatory damages. Nonetheless, this mechanism of protection may compromise effective immunity against persistent infectious agents and tumors.¹⁷⁶

More recently, the severity of lymphocytic choriomeningitis virus (LCMV) infection was associated to the number and the intensity of coinhibitory receptors expressed by virus-specific CD8 T cells.¹⁷⁷ Among these molecules, TIM-3, Lag-3, 2B4, CTLA-4, CD160, BTLA, KLRG1, CD305, and CD200R have been further investigated in the context of several human chronic virus infections and established tumors. In particular, the coexpression of TIM-3 and PD-1 was observed on both CD4 and CD8 T cells from patients with HIV¹⁷⁸ or HCV^{179–181} chronic infections and correlated with T-cell exhaustion and disease progression. In addition, TIM-3- and PD-1-expressing CD8 T cells represented a major population within tumor-infiltrating lymphocytes in several murine models of cancer and in the blood of patients with advanced melanoma.^{182,183} In all cases, TIM-3/PD-1-expressing cells represented the most impaired population of CD8 T cells. Of note, the blockade of both molecules could restore CD8 T-cell effector functions (proliferation potential and cytotoxic capacity) of antigen-specific CD8 T cells and was associated with the control of tumor growth.^{178–182}

CTLA-4 is another coinhibitory receptor upregulated in the context of chronic infections and tumors.¹⁸³ It has been shown that CTLA-4 was overexpressed on CD4, but not CD8, T cells of simian immunodeficiency virus (SIV)-infected macaques,¹⁸⁴ and HIV-infected patients.^{185,186} Furthermore, the combination of CTLA-4 blockade and 4-1BB (CD137) activation enhanced tumor rejection by increasing T-cell infiltration, proliferation capacity, and cytokines production.¹⁸⁷ Of interest, BTLA was reported to be persistently expressed by melanoma-specific CD8 T cells, thus inhibiting their antitumor function.¹⁸⁸ On the other hand, BTLA expression on CD4 and CD8 T cells decreased during HIV infection and this was associated with CD4 T-cell differentiation and activation.^{189,190} Enhancing BTLA pathway may therefore represent an alternative therapeutic strategy to overcome immune activation during chronic HIV infection.

Lag-3 is an activation-induced cell-surface molecule, whose overexpression during chronic virus infection is also commonly associated with T-cell exhaustion and functional impairment. Blocking of Lag-3 alone failed in rescuing

T-cell function or in decreasing plasma viremia during chronic LCMV infection, while blockade of both PD-1 and Lag-3 synergistically improved T-cell responses and decreased viral loads *in vivo*.¹⁷⁷ Elevated levels of Lag-3 and CTLA-4 were found in PD1+ CD4 T cells from HIV-infected patients¹⁹¹ and in tumor-derived NY-ESO-1-specific CD8 T cells.¹²⁴ Functionality of these T-cell subsets was more impaired than in Lag-3 – PD-1– or single Lag-3+ subsets.¹⁹²

SLAM family members are immunomodulatory receptors with a role in the regulation of costimulation, T-cell cytokines production, and cytotoxic activity. 2B4, which is a key molecule from this family, is involved in CD8 T- and NK-cell cytotoxicity. However, the proportion of 2B4+ CD8 T cells in HIV-infected patients correlated with immune activation of memory T cells and was increased in patients with progressive disease.¹⁹³ In addition, IFN- γ secretion and cytotoxic activity of 2B4+ CD8 T cells were significantly lower following stimulation with HIV as compared to influenza-derived antigens, respectively.¹⁹⁴ Furthermore, during infectious mononucleosis, the expression of SLAM and 2B4 on CD8 T cells correlated with severity of symptoms and viral loads.¹⁹⁵

The coexpression of molecules such as 2B4 and CD160, which have been related to potent cytolytic functions, was associated with exhaustion and regulation of virus-specific CD8 and CD4 T cells in the context of chronic virus infections.¹⁹⁶ A recent study showed a high frequency of CD8 T cells coexpressing PD-1, 2B4, CD160, KLRG1, LAG-3, and CTLA-4 in HCV infection. The coexpression of these molecules was associated with low levels of CD127 expression and correlated with impaired proliferation capacity.¹⁹⁶

The expression of another set of inhibitory molecules (i.e., PD-1, CTLA-4, CD305, and CD200R) has been investigated on CD4 T cells from HCV-infected patients. PD-1 and CTLA-4 were upregulated by HCV-specific CD4 T cells from patients with chronic infection, while CD305 and CD200R were upregulated in patients with cleared infection. Of note, the blockade of PD-Ls increased the expansion of CD4 T cells.¹⁹⁷

In the context of HIV infection, the presence of HIV-specific CD8 T cells coexpressing CD160, 2B4, and PD-1 but not Lag-3 was reported. The simultaneous expression of these molecules correlated with the level of virus replication and decreased cytokines production. The proliferative capacity was restored by blocking both PD-1/PD-L1 and 2B4/CD48 interactions.¹⁹⁸ Along the same line, another group showed that more than 30% of HIV-specific CD4 T cells expressed simultaneously PD-1, CTLA-4, and TIM-3, whereas less than 2% of simian immunodeficiency virus (CMV)- or varicella-zoster virus-specific CD4 T cells coexpressed all three receptors. The coexpression of these molecules on HIV-specific CD4 T cells was more strongly correlated with the viral load compared with the expression of each receptor individually.

Potential Therapeutic Applications of Costimulatory Molecules in Infections

The well-established immunosuppressive properties of coinhibitory molecules and the potential to revert exhausted or inactivated T-cell responses following selective blocking of their function made these markers interesting targets for therapeutic intervention in patients with persistent viral infections or cancer. To date, clinical and preclinical data are available for anti-CTLA-4 and anti-PD-1 blocking agents.^{199–202} Initial human clinical trials assessing the effects of a blocking anti-CTLA-4 antibody demonstrated not only a reduction in tumor mass and clinical benefit in a minority of treated subjects but also an increase in systemic inflammation.^{203,204} Improvement in safety of these antibodies resulted in the recent approval by the U.S. Food and Drug Administration of a human monoclonal antibody against CTLA-4 (Ipilimumab, MDX-010, Yervoy) for the treatment of metastatic melanoma. In both early and late phase trials, Ipilimumab has demonstrated consistent activity against melanoma. However, serious (grade 3–5) immune-related adverse events occurred in 10–15% of patients. Thus, while providing a clear survival benefit, Ipilimumab administration requires careful patient monitoring combined to, sometimes, treatment with immunosuppressive therapy.^{199,200} In contrast, anti-CTLA-4 blockade failed to show benefit in terms of plasma viral load or survival in acutely or chronically SIV-infected macaques.^{205,206} Since CTLA-4 is preferentially upregulated on CD4 T cells and not on CD8 T cells,¹⁸⁵ it might be possible that the blockade of anti-CTLA-4 induced an expansion and activation of CD4 T cells thus providing additional targets to HIV without significant improvement of CD8 T-cell functions.

Preclinical data showed how prevention of *in vivo* interactions between PD-1 and PD-L1 enhanced T-cell responses via the restoration of their ability to undergo proliferation, secrete cytokines, and lyse-infected cells and ultimately induce substantial reduction in viral loads. Of note, blockade of the PD-1/PD-L1 inhibitory pathway *in vivo* demonstrated a beneficial effect on CD8 T cells in mice that were lacking CD4 T-cell help.¹⁶⁶ This study identified a potentially effective immunotherapeutic strategy for chronic viral infections. This has then been further explored in nonhuman primates in a recent study evaluating the safety and immunomodulatory potential of an anti-PD-1 blocking antibody in SIV-infected macaques. The treatment was well tolerated and led to a rapid increase in virus-specific CD8 T-cell responses with improved functional quality, both in peripheral and in gut associated lymphoid tissue (GALT). PD-1 blockade also resulted in the expansion of virus-specific CD4 T cells, memory B cells, and higher titers of virus-specific antibodies. In contrast, one additional study showed an increase in CD4 T-cell activation and viral replication in mucosal sites.²⁰⁶ Furthermore, a humanized anti-PD-1 monoclonal antibody (ONO-4538) is

currently being tested in a Phase 1 study in patients with recurrent or treatment-refractory cancer. Preliminary data support the safety, tolerability, and pharmacokinetic profile of a single dose of the drug. In addition, preliminary evidences of antitumor activity were observed.^{201,202}

There is currently a strong interest in the potential for clinical interventions targeting immunoregulatory networks to enhance immunity against cancer cells and persistent viruses or to boost the efficacy of preventive and therapeutic vaccines. The studies discussed previously have yielded promising results but have also highlighted important safety issues. This strongly indicates the importance to better understand mechanisms of immune regulation in order to exploit them for potential therapeutic applications.

SUMMARY

Coinhibitory molecules are involved in maintaining the balance between the capacity to generate effector T cells able to control pathogens and the preservation of tolerance. During the development of immune responses, key coinhibitory molecules are upregulated with different kinetics and play a role in regulating the development and the fate of effector and memory T-cell responses. In most cases, pathogens replication is controlled by the immune system leading to the contraction of effector T cells. Many different coinhibitory molecules, that is, PD-1, CTLA-4, BTLA, SLAM, and 2B4, play a role during this phase. The remaining memory T cells express some coinhibitory molecules, which depend on the type and biology of the pathogens, and also on the level of differentiation. However, a hallmark of memory T cells is the lack of simultaneous expression of multiple coinhibitory molecules. Conversely, when pathogens replication is not controlled, continuous stimulation of T cells, due to antigen persistence, prevents the full contraction and leads to functional exhaustion of effector T cells. In contrast to memory T cells (see previous paragraph), a hallmark of exhausted effector cells is the simultaneous expression of several coinhibitory molecules. The simultaneous expression of these coinhibitory molecules is associated with their functional anergy, also called exhaustion. However, several findings both *in vitro* and *in vivo* suggest that this anergic state can be reverted by blocking the interactions between coinhibitory molecules and their ligands. For this reason, coinhibitory molecules are now targets of preclinical and clinical studies aimed to identify new therapeutic strategies in the context of chronic infections and tumors. To date, only two coinhibitory molecules have been investigated in clinical trials, that is, PD-1 and CTLA-4, but recent evidences have underlined the importance of targeting multiple pathways in order to improve functional restoration. It is very likely that in the close future many additional targets will be assessed in preclinical and clinical studies. In addition, while most studies focused their attention on the reversion of functional

exhaustion, additional parallel strategies may be envisioned, such as the prevention of exhaustion in the context of therapeutic immunization.

Moreover, preventing/reverting exhaustion as a therapy for chronic conditions might be difficult to achieve notably for safety issues. On the one hand, the prevention/reversion of exhaustion counteracts a physiological mechanism that is probably settled in order to avoid tissue damages and autoimmunity. On the other hand, restoration of functionality might not be sufficient since it will restore the functions of cells that failed to control the infection or to eliminate the pathogens. Therefore, it seems wise to plan to combine the functional restoration of T cells to other immunotherapeutic interventions.

Induction of the expression of costimulatory molecules on exposure to various inflammatory cytokines (IL-6, IL-12, TNF- α , IFN-c) or PAMPs/damage-associated molecular patterns (DAMPs) is regulated by transcription factors such as NF- κ B, IRF-3, AP-1, and NFAT.^{86,87} Activation of these transcription factors is tightly regulated by various kinases or phosphatases that include MAPKs, TNF-receptor-associated factor proteins (TRAF), IL-1 receptor-associated kinase 4 (IRAK4), phosphoinositide 3-kinase (PI3K), and Janus-kinase. For example, triggering of TLR-4 with LPSs elicits pathways dependent on myeloid differentiation primary response gene 88 (MyD88) and TIR-domain-containing adapter-inducing interferon- β (TRIF). Both the molecules induce downstream signaling to activate MAPKs, NF- κ B, and IRF family proteins, which are responsible for the enhanced expression of CD40 and CD86. Further, T-cell interaction with APC involving TCR and costimulatory molecules activates a plethora of downstream signaling molecules, which leads to the induction of the expression of CD40L, PD-1, and CD28. Intracellular pathogens utilize an array of mechanisms to manipulate costimulatory molecules. Upon infection, pathogens trigger IL-10 secretion by inhibiting p38MAP kinases and promoting ERK phosphorylation. IL-10 blocks the degradation of I κ B- α that inhibits the NF- κ B activation eventually inhibiting the expression of costimulatory molecules. Interference in TLRs signaling by the intracellular pathogens is considered to be a foremost event in the suppression of costimulatory molecules. ManLAM of *Mtb* binds to DC-SIGN and compromises the LPS-induced activation of DCs by interfering in the TLRs' signaling. Binding of *Mtb* ESAT-6 to TLR-2 activates AKT and prevents interaction between the MyD88 and IRAK4, consequently abrogating NF- κ B activation that suppresses CD80 expression. Similarly, HIV upregulates PDL-1 on DCs and monocytes by attenuating the signaling of TLR-7 and TLR-8. In addition, HIV-mediated PI3K activation upregulates PDL-1 on APCs and thus suppresses the activation of HIV-specific CD8+ T cells. Rapid endocytosis of costimulatory molecules upon infection is suggested as one of the mechanisms to interrupt the function of APCs. For instance, Nef protein of HIV modulates the actin-dependent trafficking mechanism to remove CD80/CD86 from the monocytes surface, making

them inefficient to activate T cells. TCR signaling is central for the optimum expression of costimulatory molecules and effector function of T cells. Therefore, it may serve as an important target for pathogens to paralyze T-cell activity. *Mtb* glycolipids interfere with TCR signaling and block the activation of CD4+ T cells. HIV gp120 interacts with CD4 (a coreceptor for TCR) that inhibits intracellular signal transduction through TCR, which leads to decreased hydrolysis of PI, Ca²⁺ influx, activation of PKC, and eventually failure of NFAT translocation. These events culminate in the inhibition of CD40L expression. Decline of CD40L on T cells abrogates the bidirectional signaling and reduces the exhibition of CD80 on APCs. In contrast, Nef activates NFAT and stimulates IL-2 release to overcome the exogenous requirement of IL-2 to promote T-cell proliferation, consequently disseminating the infection. However, Nef retards CD28 and CD4 expression, thereby making T cells incapable of promoting CMI. Corroborative with these observations, Nef-mutated HIV is unable to cause persistent infection. Similarly, core protein of HCV inhibits TCR signaling and upregulates PD-1. The above-mentioned mechanisms imply that pathogens can effectively utilize various costimulatory pathways to subvert immune response to persist in the host. The noncompliance with the relatively high dose and extended therapeutic regime reduces the effectiveness of current drugs, which leads to global emergence of MDR/XDR/TDR pathogenic strains. Interestingly, costimulatory molecules have been suggested for therapeutic intervention to treat lymphoma patients. To develop alternative or adjunct (with drugs) therapies, an intensive effort has been undertaken in the last decade to understand how intracellular pathogens exploit costimulatory molecules, which are the tour de force of the immune system. The potent role of costimulatory molecules is aptly established in the optimum activation of T cells and APCs; the cells that play a cardinal role in curbing the infections. Hence, immunotherapy involving costimulatory molecules can be a breakthrough strategy to treat various diseases, minimizing side effects inflicted by drug therapies and in restricting the emergence of drug resistance.

The prevalence of allergic diseases has increased rapidly in recent years. It is well established that the deleterious allergic response is initiated by T-cell recognition of MHC class II-peptide complexes at the surface of APCs. While this first signal gives antigen specificity to the adaptive immune response, a second nonspecific costimulatory signal is required by T cells to become fully activated. This signal is provided by interactions between APCs and T cells through molecules borne at the surfaces of the two cell types. Depending on the type of molecules involved, this secondary signal can promote the development of an inflammatory allergic reaction or may favor immune regulation. Several molecules of the B7 family (CD80, CD86, PD-1, ICOS, CTLA-4) and TNFR family (OX40, CD30, 4-1BB, Fas, CD27, CD40) play an important role in delivering costimulatory signals in early and late phases of allergic response. Therefore, costimulatory

molecules involved in the promotion or prevention of allergic immune responses are potential targets for the development of novel therapeutic approaches. Families of costimulatory molecules are involved in the regulation of most inflammatory diseases by finely controlling the intensity of the immune response. Costimulatory molecules are implicated in the development and control of allergic inflammation characterized by the establishment of an acute Th2 polarization. Elucidation of the role of costimulation pathways in the development of new subsets of Th cells has just begun, and most of the mechanisms underlying the regulation of atopic diseases by costimulatory molecules are unknown and require further investigation. With an increased understanding of these immunological mechanisms, new therapeutic strategies in the treatment of allergic airway diseases can be created by analyzing the role of costimulatory molecules that are critically involved in the induction and maintenance of allergen-induced airway diseases. Taken together, recent studies have begun to provide insight into the role of costimulatory molecules and give us new clues to design more efficient therapies to fight the increasing public health problem that allergies represent.

Some of the most successful pathogens of humans, such as *Mtb*, HIV, and *L. donovani* not only establish chronic infections but also remain a grave global threat. These pathogens have developed innovative strategies to evade immune responses such as antigenic shift and drift, interference with antigen processing/presentation, subversion of phagocytosis, induction of immune regulatory pathways, and manipulation of the costimulatory molecules. Costimulatory molecules expressed on the surface of various cells play a decisive role in the initiation and sustenance of immunity. Exploitation of the “code of conduct” of costimulation pathways provides evolutionary incentive to the pathogens and thereby abates the functioning of the immune system. Here we review how *Mtb*, HIV, *Leishmania* sp., and other pathogens manipulate costimulatory molecules to establish chronic infection. Impairment by pathogens in the signaling events delivered by costimulatory molecules may be responsible for defective T-cell responses; consequently organisms grow unhindered in the host cells. This review summarizes the convergent devices that pathogens employ to tune and tame the immune system using costimulatory molecules. Studying host–pathogen interaction in context with costimulatory signals may unveil the molecular mechanism that will help in understanding the survival/death of the pathogens. We emphasize that the very same pathways can potentially be exploited to develop immunotherapeutic strategies to eliminate intracellular pathogens.

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Chapter 4

Costimulation Immunotherapy in Allergies and Asthma

INTRODUCTION

Allergic diseases affect individuals of all ages from infancy to old age and have become one of the epidemics of the twenty-first century in developed countries. The prevalence and severity of allergic diseases and asthma have increased dramatically over the past decades, affecting an estimated 20% of the population in developed countries. Different hypotheses have been postulated to explain this increase in prevalence. The “hygiene hypothesis” is one of the main propositions that suggest improved hygiene, with improved public health measures and the use of vaccines and antibiotics, has reduced the incidence of infections that would normally stimulate the immune system. These infections would “mature” the immune system thereby protecting against the development of allergic responses against innocuous environmental substances¹. Several epidemiologic studies support the hygiene hypothesis and have clearly shown that modifications of the pattern of microbial exposure represent a critical factor underlying the rise in prevalence of atopic disorders but the immunological mechanism associated with the hygiene hypothesis has been controversial. Initial interpretations concluded that a lack of shifting from the T-helper-2 (Th2) to the Th1 phenotype (missing immune deviation) because of the reduced microbial burden was responsible.¹ A Th2 phenotype favors the development of asthma and allergy, whereas Th1 immune responses protect against asthma. However, several observations challenge the paradigm of missing immune deviation; therefore, an alternative interpretation of the hygiene hypothesis has been proposed that states that a reduced activity of regulatory T cells (Treg cells; reduced immune deviation) may be the explanatory mechanism for the hygiene hypothesis.² Treg cells are essential for the induction and maintenance of tolerance and tightly regulate both Th1 and Th2 immune responses. A high pathogen burden is instrumental for the development of Treg cells and thereby for a balanced development of the immune system.

Allergic diseases arise as a result of aberrant immune responsiveness against innocuous environmental proteins (antigens). Although we are all continuously being exposed to harmless antigens, only a certain percentage of the individuals experience adverse immunological reactions to these antigens. This is due to the fact that the normal immune response to allergens is associated with the induction of tolerance. Abrogation of tolerance or failure to induce tolerance may lead to the induction and perpetuation of active immune responses. Allergen-specific T-helper (Th) cells play a pivotal role in the pathogenesis of allergic hypersensitivity reactions and in the development of asthma.^{3,231} Activated allergen-specific T cells are present in high frequency in the peripheral blood of allergic patients. The allergen-specific CD4+ T cells involved in the initiation of allergic reactivity have a distinct cytokine profile, characterized by the enhanced production of interleukin (IL)-4, IL-5, IL-9, and IL-13 (Th2 cytokines).^{4,5} The presence of high levels of Th2 cytokines contributes to the clinical features of allergic pathophysiology. Although the most important determinant for the development of allergy and asthma is a genetic predisposition, environmental factors, such as exposure to allergens, infections, and air pollution, also play an important role in the development of allergic and asthmatic inflammatory responses. These Th cells activate a complex immune reaction that triggers the release of potent mediators and enhances the recruitment of inflammatory cells, which in turn elicit an inflammatory response that leads to the clinical symptoms of allergic disease.

The current therapies for allergic diseases focus primarily on control of symptoms and suppression of inflammation, without affecting the underlying cause. However, the knowledge about the pathophysiology of allergic diseases has substantially increased, offering new opportunities for therapeutic intervention. In this review, we will focus on current insights into the mechanism of allergic reactions. Advances in basic immunology research have enhanced our understanding of the cellular and molecular basis of the allergic response.

IMMUNOLOGY OF ALLERGY AND ASTHMA

Hypersensitivity reactions have been classified into four different types, characterized by different immunological mechanisms.^{6,231} Briefly type I hypersensitivity refers to immediate hypersensitivity responses against foreign proteins that are common (pollen, grass, animal dander, etc.). It can be observed in allergic rhinitis and allergic asthma. This type of hypersensitivity is characterized by immunoglobulin (Ig) E production during the sensitization phase, which will bind to high-affinity IgE receptor (FcεRI) on mast cells and basophils. Upon renewed contact, the allergen will bind cellular IgE. Crosslinking of FcγRI receptors will lead to degranulation of mast cells and basophils.^{6,7} Mediators released that way will lead to immediate and, sometimes, delayed immune responses. Type II hypersensitivity is a humoral

response mediated by IgG or IgM that are produced against surface antigens on body cells. An example of type II hypersensitivity is drug-induced cytopenia caused by penicillin, cephalosporin, or transfusion reactions. In drug allergic subjects, IgG or IgM will be formed in response to drugs or their metabolites, which accumulate in membrane structures. This binding will allow the killing of the target cells by three different ways.^{8–10} The first one is the activation of the classic complement pathway, which will lead to cytolysis. The second one is the antibody-dependent cell-mediated cytotoxicity, which will lead to lysis of target cells by natural killer cells. The last one is the recognition of Fc fragments of IgG and IgM by phagocytic cells, which will lead to opsonization of the target cells. Type III hypersensitivity involves the formation of immune complexes that are not well cleared by innate immune cells as in malaria, rheumatoid arthritis, or farmer's lung. This response can occur within 4–6 h. The accumulation of immune complexes (antigens bound to antibodies) in vessels and tissues can be caused by antigen excess for subjects with immunosuppression or insufficient antibody production. It can also be caused by repeated antigen exposure, which will lead to excess IgG antibody production. The presence of those persistent immune complexes will give rise to an inflammatory response due to leukocytes' activation.¹¹ Finally, type IV hypersensitivity is a delayed response principally mediated by T cells. The best-known example of type IV hypersensitivity is contact allergy. The sensitization phase lasts 10–15 days and is asymptomatic. In contact allergy, a hapten, a low molecular weight molecule, will interact with skin cells or proteins. After this contact, Langerhans and dendritic cells (DCs) will present antigen to naive T cells to stimulate their differentiation into CD4+ and CD8+ T cells and to induce production of memory T cells. A renewed contact between the skin and the antigen will stimulate sensitized memory T cells via antigen-presenting cells (APCs). Early arrival of CD4+ and CD8+ T cells and activation of keratinocytes, will all contribute to promote inflammatory reaction via cytokine secretion. Finally, this reaction will be controlled by Treg cells.¹² This classification of allergic reactions has been widely accepted.

One of the most studied allergic diseases is asthma. Asthmatic response is provoked by allergy in 75–80% of all asthmatic cases¹³ According to the Global Initiative for Asthma,¹⁴ asthma is defined as a chronic inflammatory disorder of the airways involving many cells and mediators. The principal associated symptoms are airway hyperresponsiveness and usually reversible airflow obstruction. These symptoms lead to recurrent episodes of wheezing, breathlessness, chest tightness, and coughing. Presently it is estimated that 300 million people suffer from asthma in the world and that this number will reach 400 million people by 2025. This represents a world prevalence of approximately 10% and a prevalence of 14.1% for Canada and of 10.9% for the United States.¹⁵

PATHOPHYSIOLOGY OF ASTHMA

Asthma is a complex trait that is influenced by several genes as well as by the environment; thus it presents heterogeneous clinical manifestations conventionally recognized as different subphenotypes.

Researchers agree that asthma is not a single disease but rather an array of disorders that share common characteristics.^{13,16} These characteristics are inflammation, intermittent bronchial obstruction, bronchial hyperreactivity, mucus hypersecretion, and hypertrophy and hyperplasia of smooth muscle.¹⁷ As mentioned above, in 75–80% of cases these phenotypes are caused by an allergic response, which triggers a Th2 immune response. It is a type I hypersensitivity reaction, that is an immediate exaggerated or harmful immune reaction.^{9,10} Interestingly, only 7% of allergic people develop asthma, which can lead us to believe that they present a unique phenotype that distinguishes them from other allergic, but nonasthmatic, individuals. The main immune cells involved in asthma are CD4+ T cells, mast cells, and eosinophils.¹⁸ When an allergen penetrates the airways, if not expelled by the mucociliary barrier, it comes in contact with DCs, which will internalize and then digest the allergen. Some DCs then migrate toward the lymph nodes to present the antigen to naive T cells.^{19,20} This presentation happens through the major histocompatibility complex (MHC) type II conjugated to the CD80 and CD86 costimulation molecules.²¹ The introduction of the antigen to naive T cells triggers the differentiation of CD4+ T cells into Th1 or Th2. This differentiation is modulated by the cytokines present during the introduction of the allergen, for example, IL-12 (Figure 4.1). Evidence shows, at least for mild-to-moderate asthma, that Th2 cells predominate. Thus, after the differentiation of naive T cells in Th2 cells, the latter produce cytokines, including IL-4, which induces the production of IgE by B lymphocytes (Figure 4.2).

Response to Allergens in Asthma

When an individual has already been in contact with an allergen, its presence and the presence of IgE can activate mast cells and stimulate their degranulation just minutes after the contact with the allergen. The activation happens when the allergen binds with more than one IgE, which are linked to their high-affinity receptor (Fc γ RI; Figure 4.2). Mast cells are key cells in the immediate response. In fact, mast cell granules contain pro-inflammatory molecules such as histamine; tryptase, and other proteases; tumor necrosis factor (TNF); and heparin. New molecules are also produced and then released. These molecules are leukotrienes and prostaglandins, as well as cytokines, chemokines, and matrix metalloproteinases.²² The releasing of these mediators triggers the development of immediate response symptoms such as coughing, bronchial spasms, smooth muscle contraction, edema, mucus secretion, and infiltration of immune cells. Mast cells, through these

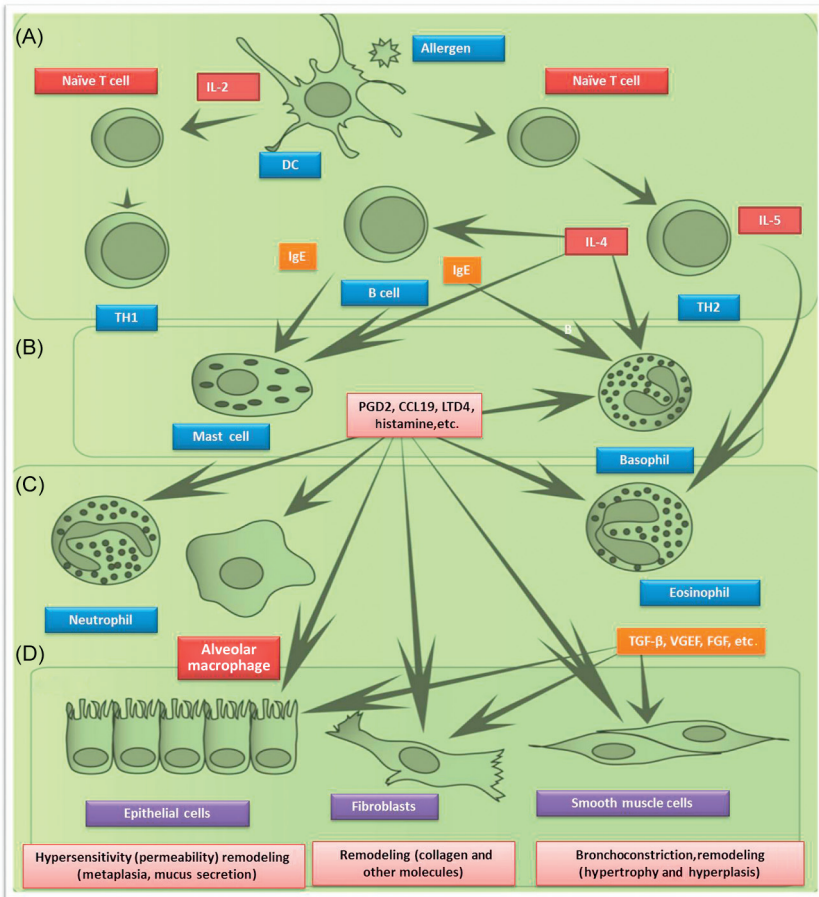


FIGURE 4.1 Innate and adaptive immune response in allergy and asthma. APCs bind with allergen, produce antigen (Ag), and display it on their surface on the MHC. This is then able to bind an antigen-specific TCR on the surface of the naïve T cell (also known as signal 1). To respond to this antigen, the T cell must also be costimulated through additional receptors (costimulation, also known as signal 2). This dual stimulation can lead to a T-helper (Th)1 or Th2 response, depending on the antigen. The innate immune system plays a critical role in determining the type of T-cell differentiation and, thus, the type of adaptive immune response. A newly described subset, Th17 cells, mediates neutrophil inflammation and may play a role in asthma. Treg cells play an important role in overall regulation of the allergic response.

mediators, contribute to the recruitment and activation of immune cells in the lungs, such as eosinophils, T cells, macrophages, basophils, neutrophils, structure cells (fibroblasts, smooth muscle, and epithelial cells), and other mast cells.^{23,24} Basophils, with the presence of Fc γ RI receptors on their surface and the expression of Th2 cytokines, histamine, and granules, can also

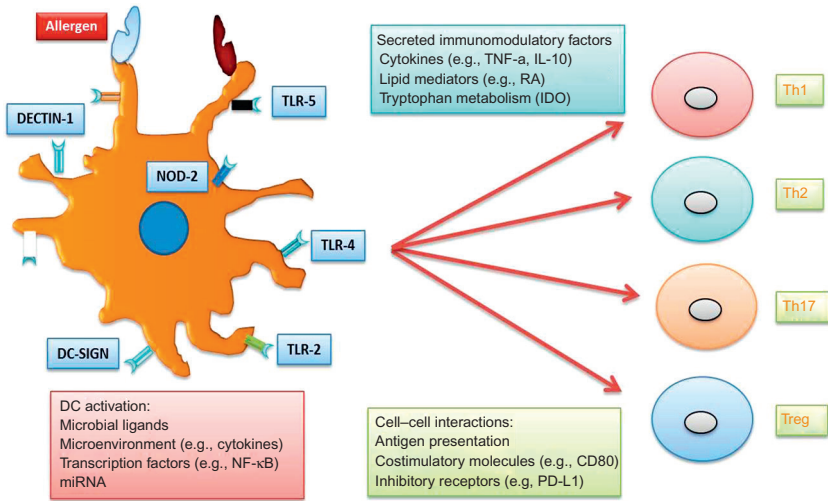


FIGURE 4.2 Cell–cell interactions in allergy and asthma. To respond to allergen, the T cell must be activated through TCR signal and costimulatory signal (costimulation, also known as signal 2). This dual stimulation can lead to a T-helper (Th)1 or Th2 response, depending on the antigen. A newly described subset, Th17 cells, mediates neutrophil inflammation and may play a role in asthma. Treg cells play an important role in overall regulation of the allergic response.

play a role in immediate response. However, their role in asthma is less documented than that of mast cells.²⁵

Delayed, or late response, is not present in all asthmatics. It is mainly modulated by immune cells recruited by mast cells⁹ (Figure 4.1). Among these cells, eosinophils are the main cells implicated in the development of this response. Eosinophils produce Th2 cytokines, leukotrienes, and proteins that cause damage to airway cells, such as major basic protein, eosinophil cationic protein, eosinophil-derived neurotoxin, and eosinophil peroxidase (EPO).^{25,26}

Chronic inflammation in airways in asthma is characterized by a persisting inflammation and the presence of structure alterations in the airways (Figure 4.1). Several immune cells play a role in this response. Mast cells participate in the chronic response through mediators that have an effect on bronchoconstriction and on airway remodeling, which causes contraction, hypertrophy, and hyperplasia of smooth muscles as well as fibrogenesis.^{27,28} DCs that remain in the airways after coming in contact with the allergen repeatedly present the antigen to CD4⁺ T cells.^{29,30} These cells, as well as the activated T cells, produce pro-inflammatory cytokines and maintain the chronicity of the inflammation and the eosinophilia. Eosinophils are indirectly involved in bronchial remodeling as regards collagen deposit, fibrogenesis, angiogenesis, and hyperplasia of smooth muscles in the peripheral airways via mediators secretion and other cells activation.^{31–33}

Phenotypic Variability in Asthma

To complete the topic asthmatic allergic response, the phenotypic heterogeneity has to be taken into consideration. Indeed, apart from allergic asthma, which is the most common form, other types of asthma have been described.^{9,34} However, as with allergic asthma, the scientific community recognizes that IgE could be implicated in an inflammatory cascade, as well as CD4+ and CD8+ T cells, eosinophils, and mast cells. A second type of asthma is characterized by sensitivity to aspirin. It is estimated that approximately 10–20% of adult asthmatics suffer from this particular type.³⁵ Aspirin sensitization is thought to be non-IgE mediated so could be considered as a nonallergic subphenotype of asthma. A dysfunction of the eicosanoid metabolism is responsible for this type of asthma. It is estimated that 9–15% of all cases of asthma in adults are linked to the workplace and that up to 25% of new cases of adult asthma fall in the occupational asthma category. The two major hypotheses that could explain the induction of asthma after an effort are related to the augmentation of ventilation in the lungs. These two hypotheses are the osmotic hypothesis (dehydration by evaporation caused by the increase in ventilation) and the thermal hypothesis (airway cooling during exercise and rewarming after exercise).³⁶ These two hypotheses could better explain the phenotype rather than each one separately.

Asthma can also be classified according to inflammatory patterns. The three principal types are eosinophilic asthma, neutrophilic asthma, and paucigranulocytic asthma.¹⁶ Paucigranulocytic asthma is defined as an asthma response without eosinophils or neutrophils. Another way to classify asthma is according to the severity of the phenotype. This classification is used in the clinical treatment, but is also used in research in order to document the molecular and cellular biology variation related to the severity of the disease. Intermittent, mild, moderate, or severe asthma are part of this classification. It is also interesting to note that a recruitment of Th1 and CD8+ T cells (cytotoxic T cells) has also been observed in the case of severe and chronic asthma and during exacerbation. Thus, other cell types involved in the Th1 response or in immune response regulation, such as alveolar macrophages (they inactivate DCs and they can be activated or inhibited through different pathways), could play a role in the inflammatory response observed in asthma.^{37,38}

A prominent characteristic of the allergic response is a persistently elevated level of IgE antibodies against specific antigens (allergens) to which the affected individual is regularly exposed by inhalation, ingestion, or contact with the skin.³⁹ Allergic sensitization involves processing of the antigen by an APC and presentation, in association with a class II MHC protein, to a T-cell receptor (TCR). While DCs (and macrophages) predominantly act as APCs in primary immunization, B cells may also participate in secondary immune responses to allergens. Activation of T cells requires the signal from

this trimolecular interaction and an additional costimulatory signal resulting from the binding of B7 (CD80/86) on an APC to CD28 or CTLA-4 on a T cell.⁴⁰ When stimulated by antigens, Th cells produce specific cytokines that have been designated as either Th1 cytokines [IL-2, interferon (IFN)- γ , and TNF- β] or Th2 cytokines (IL-4, IL-5, IL-9, IL-10, and IL-13). The state of T-cell activation depends on several factors.^{40,41} The first of these factors is the strength of the interaction between the antigen and the T cell. The site of the antigen recognized by the T cell is termed the *epitope*. The affinity of a T cell for a specific epitope depends on the concentration of the antigen, the type of APC,⁴² and the cytokine milieu of the T cell during antigen interaction. Thus, IFN- γ and IL-12 promote a Th1-like response, whereas IL-4 promotes a Th2-like response.⁴³ Additionally, host immune system genes may bias the overall immune responsiveness of an individual to favor a Th1- or Th2-like phenotype. A Th2-like cytokine profile is associated with the induction of IgE antibody (Ab) production *in vitro* and *in vivo*.⁴⁴ Specifically, IL-4 favors the development of Th2-like cells from uncommitted T cells, and both IL-4 and IL-13 play a role in IgE antibody production. Manifestation of an allergic reaction depends on the specific IgE levels and the amount of exposure at the time of the reaction. Although an allergic condition is a risk factor for asthma, 20–30% of asthmatics do not show positive skin tests to allergens. In general terms, asthma is an inflammatory disease in which not only lymphocytes but mast cells, basophils, eosinophils, and epithelial cells play a role. Studies to date suggest that Th2-like cytokines, such as IL-4 and IL-5, also play an important role in nonatopic asthma^{45,46} (Figure 4.2).

ROLE OF MAST CELLS IN ALLERGY AND ASTHMA

Mast cells arise from progenitor cells in the bone marrow and full maturation of mast cells occurs in the peripheral tissue under the influence of various cytokines and other factors in the extracellular milieu. Mast cells express a great variety of stimulatory and inhibitory receptors. Antigen-specific mast cell activation can occur via IgE, IgG1, IgG2a, and IgG2b through crosslinking of the Fc ϵ RI and the Fc γ RI receptor.⁸ Antigen-specific mast cell degranulation can also be induced by crosslinking of Ig-free light chains.⁴⁷ Further, antigen-independent mast cell activation can be established by various other receptors such as neurokinin receptors, c-kit, Toll-like receptors (TLRs), complement receptors, CD200 receptors, gp49a, or via direct interaction with G-proteins.^{8,48–51} Activated mast cells release their granule content via exocytosis (anaphylactic degranulation) or via differential release of mediators without degranulation (piecemeal degranulation).⁵² Furthermore mast cell activation results in the synthesis and release of lipid mediators and production of various chemokines and cytokines.⁵³

Mast cells play a significant role in allergy and asthma and their role in allergic diseases has extensively been studied. Mast cell triggering plays an

important role in the elicitation of the immediate phase of an allergic response leading to acute local responses such as edema formation, tissue swelling, or bronchoconstriction. By the release of chemotactic and proinflammatory mediators, mast cells can also have an effect on the late-phase responses of an allergic response.⁴⁹ Mast cells have been demonstrated to be of importance in the elicitation of a full-blown inflammatory response. Because of their close anatomical association with sensory nerves, bidirectional communication between mast cells and nerves can further influence the inflammatory responses.

In chronic asthma it has been demonstrated that multiple exposures to antigens resulted in FcR γ -dependent (via IgE and/or IgG1) and FcR γ -independent mast cell activation. This mast cell activation contributes to the induction of enhanced airway responsiveness and chronic inflammation including eosinophil and lymphocyte infiltration, goblet cell hyperplasia, and increased levels of lung collagen.⁵⁴

In human asthmatic lungs, mast cells migrate into the airway epithelium, the airway mucous glands, and the airway smooth muscle, while virtually no mast cells were found in these regions in normal human airways. The number of mast cells infiltrated in the airway smooth muscle bundles was shown to correlate significantly with bronchial hyperresponsiveness in asthmatics, implicating their importance for the pathophysiology of asthma.⁵⁵

IMMUNOTHERAPY TO ALLERGENS

The term “allergy” was first introduced by Clemens Freiherr von Pirquet, an Australian pediatrician, who believed that it was a pathological state of altered immune reactivity. However in 1911 Leonard Noon and John Freeman proposed the concept of allergen immunotherapy (IT). They hypothesized that toxins from grass pollen somehow accounted for symptoms seen in patients suffering from hay fever. By inoculating the patients with gradually increasing doses of the toxin itself to stimulate the immune system against the toxin, the symptoms could be reduced or even abolished.

Since then allergen IT has become a proven approach for treating allergic rhinitis and allergic asthma and has undergone significant development in the last two decades. Current allergen-specific IT (SIT) involves administering increasing doses of the causative allergen in order to reduce the clinical signs and symptoms associated with exposure to the allergen and thereby produce tolerance. As currently practiced, IT involves subcutaneous or sublingual administration of allergens, both methods of which have been extensively investigated. In addition to allergen IT, a number of additional non-SIT approaches are being used or are in phase II/phase III clinical trials. Such therapies include anti-IgE antibodies and the soluble IL-4 receptor (sIL-4R). Other experimental IT approaches are at the preclinical research stage and may also proceed to clinical trials and the clinic within the next

5–10 years. Here in this chapter we will discuss the pros and cons of recent developments in both currently practiced and experimental IT approaches with a focus on costimulation-based IT for allergy and asthma.

The allergens are administered by two different routes: parenteral or subcutaneous IT (SCIT) and sublingual IT (SLIT), which was introduced relatively recently. The current state of the art for each of these immunotherapeutic approaches is discussed below.

Subcutaneous IT

As currently practiced, SCIT has proven effective in allergic rhinitis and asthma and is Food and Drug Administration (FDA) approved and reimbursable. SCIT has been extensively studied in double-blind trials to determine effective doses, establish duration, define the mechanisms, and to investigate the persistence of efficacy after treatment ends. However, the need for multiple visits to the clinic for shots is inconvenient, only a few allergens have been standardized for SCIT, and the potential for systemic anaphylactic reactions is a serious limitation. Although allergen IT has been around for about a century, little is known about the absorption and fate of subcutaneously administered allergen. The pharmacokinetics of SCIT has been studied using leukocytes labeled with ^{99m}Tc -HMPAO on allergic patients injected intravenously in contralateral arms. Local inflammatory activity was noted in the first hour and the increase was time-dependent. The immune system responded quickly as the activity was traced in the lymphoid tissue of the upper mediastinum and anterior region of the neck. Thoracic and bowel focalization was also noted for the subcutaneous route. Factors important in SCIT include the dose of allergen being administered, the quality of the allergen extract, and the duration over which it is given.^{56,57} The effects of duration of allergen dosing appear to depend on individual factors according to a review of studies performed between 1976 and 2006 in which the rate of relapse ranged from 0% to 50%.⁵⁸ The immune mechanism of SCIT has been extensively studied. SIT alters allergen-specific T-cell responses from Th2-like to Th1-like and reduces inflammatory cells and mediators in the target nasal mucosa and the airways.⁵⁹ For example, IT with house dust-mite extract induces antigen-specific suppressive activity by CD4+ CD25+ T cells in allergic subjects, which causes the regulatory/suppressor T cells to secrete IL-10 and transforming growth factor- β (TGF- β). This in turn increases IgG4 and IgA antibody production and suppression of IgE antibodies by B cells, thus mimicking a healthy immune response to environmental allergens.

Interestingly, a controlled study of traditional allergen vaccine IT to prevent asthma in atopic children demonstrated that more children in the non-treated group developed asthma than in the IT group. Moreover, it has been shown that IT in children sensitive to a single aeroallergen prevented new sensitization to other allergens.⁶⁰ Currently, only 2–3 million people of the

55 million with allergic diseases are on SCIT. The main reasons for this are that the target populations dislike injections or cannot afford the time and expense of frequent visits to a doctor's office. There is also a concern about the safety of SCIT, and it cannot be used for young children with allergy or asthma. In an effort to decrease the frequency of allergen administrations, intralymphatic allergen administration was attempted in a randomized (but not blinded) controlled trial.⁶¹ While the results of this study are promising, it was not a blinded study and the technology of ultrasound-guided intralymphatic injection may not be available in all clinics.

Sublingual IT

SLIT consists of placement of an aqueous allergen extract (50% glycerin and 0.4% phenol) in single allergen tablets or capsules under the tongue. It has been extensively practiced in Europe,⁶² and tested in clinical trials in the United States but is not yet FDA approved. Presently, SLIT is considered to have about one-tenth the risk of SCIT and has been shown to prevent new sensitization and progression from rhinitis to asthma. In addition, SLIT provides effective protection that persists for a long time after treatment is stopped. The disadvantages are that SLIT is less effective than SCIT and has not been tested for multiple allergen mixes. The pharmacokinetics of local IT by SLIT has been studied in allergic volunteers using radiolabeled allergen given in the form of soluble tablets. Plasma radioactivity was measured at different intervals using early and late sequential scintigraphic acquisition. No absorption through the oral mucosa was observed, and the plasma radioactivity, which peaks at 2 h, was increased only after swallowing. No allergen was detected in the blood stream.⁶³ Another study, in which the biodistribution of ¹²³I-radiolabeled Der p2 was assessed in allergic volunteers, confirmed these findings. Plasma radioactivity was only noted after the tablets were swallowed and the effect peaked at 1–2 h.⁶⁴ It was noted that in order for the IT to be successful, the allergen had to be swallowed after the tablet had dissolved sublingually. This confirms that the contact with oral mucosa is an important factor, although no allergen was absorbed through the oral mucosa. It was also found that duodenal enzymes were necessary for hydrolyzing the allergen.⁶⁵ Treatment with SLIT is initiated with a build-up phase lasting 4–6 weeks starting with the lowest concentration. The dosage is then gradually increased up to a maintenance dose. The recommended duration of the treatment is 3–4 years for optimal results.⁶⁶ SLIT can be administered coseasonally, preseasonally, or continuously.⁶⁷ The majority of SLIT studies have shown clinical efficacy in the form of improvement in symptom scores or medication scores, or both for allergic rhinitis and allergic asthma. A meta-analysis of five randomized double-blind, placebo-controlled studies on patients with allergic asthma showed significantly improved symptom scores and decreased use of rescue medication.⁶⁸

Experimental IT Approaches

The availability of recombinant allergens makes it possible to treat allergic patients with identical, consistent vaccines. Recombinant allergens have other advantages which include (1) administration of allergens in optimal doses; (2) immunization with vaccines tailored to the major allergens to which the patient is allergic; (3) more accurate dosing; and (4) the possibility of modifying the structure of IgE-binding allergen epitopes, thereby increasing the safety of allergen IT. Recombinant allergens are not yet used in medicine since like other recombinant proteins; they must undergo rigorous clinical trials before they are available for clinical use.

Vaccination with a recombinant allergen before sensitization resulted in a state of immune deviation—a shift from a Th2, IgE response to a Th1, IgG2a response in a murine model.⁶⁹ Because the immune system of neonates and adult mice is similar in terms of their ability to develop immune deviation, and because allergic sensitization occurs predominantly in the first 2 years of human life,^{70,71} using recombinant allergens for IT holds promise and needs further investigation. The recombinant allergens for IT may be used as aqueous, enteric-coated, or liposome-packaged vaccines or polymerized as recombinant allergoids using formaldehyde. Polymerized recombinant allergens are antigenic but nonallergenic and therefore may improve allergen-SIT. The problem with the available polymerized allergen vaccines is that they are not standardized and considerable batch-to-batch variation exists, making it uncertain whether all allergens have been uniformly polymerized or if some have been denatured. Recombinant allergens may also be conjugated with *n*-formyl-methionyl-leucyl-phenylalanine before being used to treat allergic patients.⁷² These chemically modified recombinant allergen vaccines should provide a safer and more effective vaccine with fewer injections than are needed with current allergen vaccines. Furthermore, once the epitopes have been identified, an allergen-cDNA can be altered specifically to reduce the IgE-binding ability of the corresponding recombinant allergen without compromising its capacity to stimulate T cells. Genetically modified allergens not only appear to be safe, but also may prove to be effective agents for patient-tailored IT.

Allergen genes expressed in an appropriate host may be used as live vaccines while recombinant allergens and their corresponding cDNAs can be used to prevent allergen-specific IgE responses. Genes can be introduced via live bacteria, or viruses, or using plasmid vectors. Two models are noteworthy. First is oral immunization with *Salmonella typhimurium* expressing the cDNA coding for the major birch tree allergen, Bet v 1. This live vaccine promoted IgG2a instead of the default IgE antibody response in mice;⁷³ however, the Bet v 1-specific IgG2a response could be detected in only 10% of the immunized mice. Second, a live vaccine to treat allergic diseases using recombinant Bacillus Calmette-Guerin-expressing allergens has been proposed.⁷⁴

These experimental approaches appear promising and may lead to effective prophylactic allergy vaccines. About one-quarter of the population is genetically predisposed to develop allergic disease. With advances in the identification of genes associated with asthma, it may be possible to develop methods for predicting atopic predisposition, which may then allow for the vaccination of predisposed individuals against the dominant allergens in their environments.

Alternatively, plasmids expressing allergens can be used for therapeutic vaccination. The immunization of mice with an allergen-cDNA cloned in a plasmid vehicle resulted in an allergen-specific IgG2a and Th1-like response, with no detectable IgE. Furthermore, the gene immunization induced a Th1-like response by reversing the ongoing allergen-specific Th2-like response. These studies suggest that immunization with allergen-cDNAs may provide a novel type of IT for allergic diseases; however, the application of DNA vaccines as a prophylactic may be more feasible because of the possibility of inducing anti-DNA antibodies and autoimmunity.

Peptide-Based Therapies

Because T cells play a dominant role in IgE synthesis, T-cell peptides may be useful as immunotherapeutic vaccines. T-cell peptide vaccines induce energy in allergen-specific T cells or interfere with the formation of the trimolecular complexes involving the interaction of MHC–peptide duplexes with the appropriate TCR.^{75,76} Peptide vaccines have several advantages: a defined chemical structure, simplicity of preparation, and prolonged shelf life. They also appear to be safe because, at least in mice, T-cell peptides do not bind to IgE;⁷⁷ however, human immune responses to allergens are more complex than murine immune responses. Various theoretical constraints are predictable: (1) some major allergens, particularly pollen allergens, contain several T-cell epitopes that are recognized by allergic individuals; (2) certain major allergens appear to have various isoforms containing crossreacting and noncrossreacting epitopes, which increase the repertoire of allergenic epitopes;⁷⁸ (3) allergic individuals differ with respect to their recognition of these epitopes; (4) B-cell and T-cell epitopes may be present on the same peptide, which increases the risk of systemic reactions; and (5) excessive doses of some peptides may induce autoimmune reactions. Thus, in individuals allergic to certain complex aeroallergens, treatment with peptides may not be effective or appropriate. In experimental animals, synthetic Fel d1 (cat major allergen) or Der p1 (house dust-mite major allergen) T-cell peptides induced peripheral T-cell tolerance.^{79,80} The activation of allergen-specific T cells and IgE antibody synthesis was inhibited by *in vivo* administration of peptides by intranasal, oral, and subcutaneous routes.

The largest clinical study with allergen peptides was conducted using a total of four injections of 750 μ g each of two peptides from the cat-allergen Fel d1 (ALLERVAX-CAT; ImmuLogic, Boston, MA) at 2-week intervals.

In a multicenter, randomized, double-blind, placebo-controlled study, 133 cat-allergic patients chronically exposed to cats or who had failed previous conventional cat IT, were given ALLERVAX-CAT as a 750 μg dose, and patients with reduced baseline forced expiratory volume in 1 s (FEV1) showed improved pulmonary function. Side effects of the treatment were considered to be of mild-to-moderate severity and were reported by 77 patients who had at least one respiratory system event including chest tightness, dyspnea, coughing, throat irritation, wheezing, and asthma aggravation. The most common dermatologic reaction presented was pruritus. Although severe adverse effects (AEs) did occur, there was no statistical difference in the frequency of their appearance between the treated and placebo group.⁸¹ Therefore, although peptide therapy in severely cat-allergic patients was associated with some adverse effects, the therapy was found to be effective.

Allergen-Specific Therapy with Th1-Stimulating Adjuvants

Immune deviation from a Th2-like to Th1-like response may be achieved by designing vaccines with allergens or recombinant allergens in conjunction with adjuvants. The adjuvants may be cytokines or synthetic compounds, such as immunostimulatory DNA sequences containing a cytosine-phosphoguanosine (CpG) motif that induces a Th1-like cytokine response.^{82,83} The adjuvant may be injected with natural allergens or genetically linked with allergen-cDNA. Among the cytokines, IFN- α and IL-12 both induce a strong Th1-like cytokine profile and therefore have the potential to convert allergen-specific Th2-like responses to Th1-like responses. IL-12 has also been suggested as an adjuvant for vaccination against diseases in which the Th2 profile predominates. IFN- τ , a type I IFN that lacks the toxicity associated with type I IFNs, inhibited IgE production in a murine allergy model and in an IgE-producing human myeloma cell line. Administration of a recombinant allergen (ovalbumin) vaccine with the IL-12 (subunit p40) fusion protein downregulated ovalbumin-specific IgE responses *in vivo*;⁸⁴ however, IL-12 may cause side effects, and its effectiveness requires IFN- α production by the target cells. Because IL-12's ability to convert an established Th2 to a Th1 response remains uncertain, the use of IL-12 as an adjuvant for allergen vaccines is unproven. Also of note are the studies involving TLR antagonists, which are based on the observation that LPS, the ligand of TLR-4, inhibits Th2 cytokine production in nasal explants of children.⁸⁵ MPL, a chemically modified derivative of LPS, was still a TLR-4 agonist and decreased combined symptom and medication scores compared to placebo in clinical trials.⁸⁶ A ragweed-TLR-9 agonist vaccine for IT was also tested in clinical trials in allergic rhinitis patients. While the results for the primary endpoint were not statistically significant, the secondary endpoints were significantly reduced. A similar study with this TLR-9 ligand in asthmatics showed a clear reduction of FEV1 and sputum eosinophils after

allergen challenge.⁸⁷ Another potential form of IT involves the Fc γ 1-linker-major cat-allergen, Fel d1, fusion protein, which was effective in inhibiting allergic responses in mice *in vivo* and human cells *in vitro*.⁸⁸ Gamma Fel D inhibits cat-allergen-induced degranulation of human basophils and cord blood mast cells in culture. Since results of mouse studies may not be translatable to humans, the proof-of-concept needs to be obtained in humans and the potential adverse effects need to be examined. Also, IT with mite allergen Der p1 on virus-like particles was reported to be safe and highly immunogenic in healthy adults.⁸⁹ The IgG response to dust mite allergens, particularly IgG1 and IgG3, but not IgG2 and IgG4, was significantly induced by these particles. However, this study was limited by lack of comparisons between Der p1 on virus and Der p1 without virus, lack of cellular studies and appropriate clinical endpoints relevant to human allergy and asthma.

Novel Approaches to Allergen-SIT

Although excellent drugs to control the symptoms of allergy and asthma are available, the only treatment able to cure allergy remains allergen-SIT. SIT is, however, rarely chosen as a therapeutic option by allergic patients mainly because it requires a treatment period lasting between 3 and 5 years to reach protection against the offending allergen. This unsatisfactory situation has strongly stimulated innovative research activities within the allergy field aimed at making SIT faster and safer in order to augment patient's compliance. Two main problems need to be reduced or eliminated to render SIT more attractive, and these are side effects and treatment time. Several exciting approaches to solve these issues are currently under clinical investigation. Therapy-related SIT side effects derive from the intrinsic property of allergens to crosslink high-affinity receptors for IgE on effector cells resulting in their degranulation and release of premade toxic substances, like histamine and leukotrienes, leading to local or generalized anaphylactic reactions. Allergoids, hypoallergens, and immunodominant allergen peptides are allergen surrogates with markedly reduced IgE-binding capacity, while conserving the whole and/or the immunodominant T-cell epitopes required for eliciting protective (IgG-mediated) immune responses. SIT with allergoids reached the market some time ago and has yielded results comparable to those obtained with allergen extracts with a reduced side effect profile, but it still requires a long treatment time. Peptide IT showed promising results in clinical trials with bee venom allergic and cat-allergic individuals.

Many genetically engineered hypoallergens have shown promising results in mouse models of allergy. Perhaps the more advanced form of IT with allergens showing a favorable safety profile is SLIT. SLIT still suffers from a long treatment time comparable to those of classical SCIT but is increasingly used as a therapeutic option, especially in Europe. These approaches have significant potential for improvement, particularly in combination with

novel adjuvants and modified treatment regimens. Obviously, a long treatment time strongly reduces patient's compliance, and therefore many efforts have been undertaken to reduce the number of injections (currently 50–80 are required) in order to achieve protection. Four different strategies have entered phase I/IIA clinical trials and yielded promising results: peptides covalently coupled to virus-like particles,⁸⁹ A-type CpG oligodeoxynucleotides as an adjuvant,⁹⁰ direct injection of allergens into lymph nodes,⁶¹ and modular antigen translocating vaccines, which directly target the MHC class II antigen presentation pathway.^{91,92} All these therapeutic innovations have, if confirmed in larger multicenter clinical trials, the potential to turn IT into a true vaccination approach because only a few injections are required to achieve long-lasting protection. These approaches indicate that the problems related to the long treatment time of classical IT, which is mechanistically based on immunomodulation, which requires a long time to establish, can be reduced using strong adjuvants, by changing the route of application, or by direct targeting of the antigen presentation pathways. We conclude that the limited efficacy associated with classical SIT is not closely related to the potential of the immune system to mount protective responses against allergens. There is clear evidence that low amounts of antigen, such as those encountered during natural exposure to most of the environmental allergens, favor the switch of B cells toward IgE production in predisposed individuals, whereas higher doses of antigen favor the development of protective IgE antibody responses. Therefore, the most promising approaches to faster and more successful SIT treatment regimens will most likely derive from strategies aimed at increasing the therapeutic dose of the allergen in lymph nodes, while avoiding effector cell degranulation.

Current Immunomodulatory Strategies for Asthma Under Investigation

A number of anti-cytokine therapies are currently being examined. Daclizumab (an IL-2 receptor antagonist) has been examined in a phase II study in patients with moderate-to-severe asthma. Patients treated with daclizumab showed improved pulmonary function; reduced asthma symptoms and medication use; increased interval between severe exacerbations; and reduced blood eosinophils and serum eosinophil cationic protein levels.^{93,94} Two humanized IL-5 monoclonal antibodies (mAbs)—reslizumab and mepolizumab—have been studied in humans. Mepolizumab decreased blood and airway eosinophils, but no significant sustained changes in FEV1, asthma symptom scores, sputum eosinophils, or physician-evaluated overall condition were noted with mepolizumab or reslizumab compared with placebo.^{95,96} In order to neutralize the activity of IL-4, recombinant human sIL-4R has been examined.⁹⁷ Patients with moderate, persistent asthma had significant improvement in asthma symptom scores, rescue β -agonist use, and exhaled

nitric oxide levels compared with those receiving placebo.⁶³ However, a follow-up study revealed that only asthma patients receiving the highest dose of sIL-4R maintained their lung function.⁹⁸ Blocking TNF- α using neutralizing antibodies improved lung function and quality of life and reduced exacerbation frequency in patients with asthma in some studies but not in all studies.^{99–101} It is likely that additional cytokines (e.g., chemokines) will also be targeted in the near future for patients with allergy and asthma. Several humanized mAb that bind to the Fc portion of the IgE molecule have been used in clinical studies, and one (omalizumab) is currently available for treatment of allergic asthma. Anti-IgE in combination with allergen-SIT resulted in a greater reduction in seasonal allergic rhinitis symptoms and rescue medication scores compared to allergen-SIT alone.¹⁰² Importantly, the addition of omalizumab to a rush SIT protocol significantly reduced the risk of anaphylaxis.¹⁰³

Anti-histamine compounds that target the histamine receptor 1 (H1R) have been used for some time. However, the discovery of a new histamine receptor, H4R, has provided a new target for drug development. Thus far, only animal model data are available that demonstrate proof of the efficacy for H4R antagonists in models of asthma, allergic rhinitis, and pruritis.¹⁰⁴ Candidate molecules have been identified, which are expected to enter clinical trials shortly.

Monoclonal Anti-IgE Antibody Therapy

The development of an allergic state is a gradual process, and as a genetically predisposed individual undergoes the “allergic march,” he or she may develop allergies to several antigens.¹⁰⁵ Because IgE crosslinking on the mast cell membrane by allergens is an important first step in an allergic reaction, IgE has been an attractive therapeutic target. Although the idea of anti-IgE as a therapeutic agent was conceived during the 1970s, the use of anti-IgE as a therapeutic strategy failed because the antibodies initially employed caused mast cell degranulation. With the advances in recombinant DNA technology, it has become possible to engineer mAbs that reduce unbound IgE levels in the serum by binding to circulating serum IgE antibodies but not to IgE antibodies bound to Fc ϵ RI or to Fc ϵ RII.¹⁰⁶

Two mouse monoclonal anti-IgE antibodies, TES-C21 and MAE11, have been humanized¹⁰⁷ and MAE11 (rhuMAb E25) has been evaluated in clinical studies. In humanizing MAE11, a framework derived from consensus sequences of human VL and VH chains was used, and the critical amino acids responsible for binding IgE were engrafted onto a consensus human IgG1 framework.⁸⁹ Several clinical studies have been conducted to evaluate the safety and effectiveness of using anti-human IgE antibodies to treat allergic asthma and allergic rhinitis. The results of these studies show that anti-IgE therapy reduces allergic symptoms, is well tolerated by patients, and

does not cause any severe AEs.¹⁰⁷ The two monoclonal anti-IgE antibodies, rhuMAb E25 and CGP51901, have been shown in a number of clinical trials including subjects with asthma or allergic rhinitis to be safe and effective in generating tolerance. The anti-IgE antibody CGP51901, which is a chimeric version of the mouse monoclonal anti-IgE TES-C21, has been evaluated in patients with seasonal allergic rhinitis.¹⁰⁸ These studies have shown a decrease in free serum IgE but an increase in total IgE levels. The slow clearance of IgE–anti-IgE complexes explains the increase, with complexed IgE having a half-life of 11–13 days. The reduction in free serum IgE levels was dependent upon anti-IgE doses and was reversible. It was necessary to sustain 85% or greater reduction in free serum IgE levels to achieve improved clinical symptoms, and an 85% reduction in IgE required a serum CGP51901 concentration of 5000 ng/mL. The treatment was safe, with no serum sickness and only one case of urticarial among 153 evaluated patients.¹⁰⁹

The anti-IgE antibody rhuMAb E25 has been evaluated in both allergic rhinitis and asthma. It decreased free serum IgE in a dose-dependent fashion in patients with seasonal allergic rhinitis;^{232,233} however, in this study, it was not possible to evaluate its clinical efficacy because only 11 subjects had undetectable IgE levels. To achieve undetectable IgE concentrations, the dose of rhuMAb E25 was 0.005 mg/kg/week for each international unit per milliliter of baseline IgE. There were no AEs related to this treatment. The rhuMAb E25 antibody has also been studied in asthma, but asthma symptoms, rescue medication use, or FEV1 did not improve; however, there was improvement in the allergen early- and late-asthmatic response, methacholine reactivity, and some inflammatory parameters in induced sputum. The evaluated patients had mild asthma, which may have limited the potential for improvement. The administration of rhuMAb E25 significantly increased the allergen doses required to cause a 15% decrease in FEV1, with a correlation between the decrease in free serum IgE and the protection against inhaled allergen.^{110,111} The same study also showed that a higher dose of methacholine was necessary to reduce FEV1 by 20% after administration of rhuMAb E25. Another study showed rhuMAb E25 reduced maximal bronchoconstriction by 60% during the late asthmatic response following allergen challenge. Reduction of the early response after bronchial allergen challenge was less affected.⁷¹ The late asthmatic response following allergen challenge correlates with airway hyper-responsiveness, airway inflammation, and improvement of asthma symptoms, suggesting rhuMAb E25 may have a clinical benefit. Additional studies support this possibility with rhuMAb E25 administration reducing the number of asthmatic exacerbations and facilitating a 50% dose reduction in use of inhaled and oral steroids.^{112,113}

Among all of these reports, only minimal toxicity was observed with one case of urticaria. Recently, however, more side effects have been noted including local injection site irritation, upper respiratory infection, headache, and urticaria.¹¹⁴ Among the total of 118 cases of anaphylaxis found by the

FDA AE reporting system, 19 had anaphylaxis within the first hour, 32 had reaction after receiving the first dose, while 14 reacted after the second dose, and 77 patients required hospitalization. These patients were treated with epinephrine or corticosteroids, and had anti-IgE withheld or discontinued. The vaccine was given subcutaneously every 2 or 4 weeks and the dose was determined by body weight and total pretreatment serum IgE levels. Absorption is slow from the inoculation site and the mean half-life of the drug is 26 days,¹¹⁵ while the evidence shows that anti-IgE is a safe and effective immunotherapeutic treatment for asthma, it is noteworthy that anti-IgE antibodies also aid with allergen IT and reduce the risk of adverse reactions. Casale et al.¹⁰³ reported that anti-IgE reduces acute allergic reactions with SCIT rush IT, which reduces the frequency of hospital visits inherent with classical SCIT. In addition to the new treatments described above, several other potential approaches are being explored. Most of these are aimed at modulating the allergic immune response and inflammation in murine models using either soluble receptors or antibodies that alter cytokines, immunocyte coreceptors, IgE receptor binding, and cellular adhesion.

IgE-Binding Receptors

The IgE antibody binds to the FcεRI and CD23 on mast cells and basophils and to CD23 on monocytes and eosinophils. Both receptor and ligand are potential therapeutic targets. Antibodies attached to CD23 may be able to facilitate antigen presentation to T cells, but IgE regulation was not affected. Transgenic mice that overexpress CD23 on T and B cells exhibit decreased IgE production,¹¹⁶ which suggests that enhancing CD23 expression on B cells before activation can effectively inhibit IgE production. Conversely, a decrease in CD23 levels with a metalloprotease resulted in an increase in IgE.¹¹⁷ The administration of anti-CD23 antibodies inhibits eosinophil airway recruitment and reduces airway hyperresponsiveness.^{118,119} This observation suggests that anti-CD23 antibodies may be used to block the interaction between IgE and CD23, reducing eosinophil recruitment and inflammation. More studies are necessary before IgE receptors can be viewed as important therapeutic targets for allergic disease.

Anti-IgE Monoclonal Antibodies

The anti-IgE antibody omalizumab has so far only been studied as a treatment for severe allergic asthma. There are well-documented effects on risk of asthma exacerbation.¹²⁰ The drug is expensive and requires injections at monthly or biweekly intervals. A reduction in free IgE levels following the anti-IgE therapy could lead to reduction in FcεRI expression on mast cells, basophils, and DCs.¹²¹ One could speculate that treatment with anti-IgE in high-risk children could prevent allergic sensitization and delay or prevent development of

allergic asthma. On this premise, an observational study has shown that after stopping prolonged treatment with omalizumab, asthma control can persist as a residual effect, possibly indicating a modifying effect on the natural history of asthma.¹²² The results of the XPORT trial (NTC01125748), a double-blind placebo-controlled withdrawal study of moderate-to-severe asthmatic adult patients who received omalizumab for ≥ 5 years, have been recently disclosed. Patients were randomized to either continue receiving omalizumab, or switch to placebo for one additional year. At the end of that period, whereas the proportion of patients receiving the active treatment who had not suffered an asthma exacerbation was 67%, this only happened in 48% of patients in the placebo group. However, this would suggest that in about half of patients the effect of omalizumab would persist at least for 1 year after it was stopped.¹²³ Furthermore, omalizumab reduces airway remodeling by modulating bronchial reticular basement membrane thickness and eosinophil infiltration.¹²⁴ New immune modulator interventions are being developed to improve the immune response to viral infections in children with impaired innate immunity, as shown by decreased levels of IFN- γ or impaired function of TLRs.¹²⁵ Examples are inhaled IFN- β and a TLR agonist. They are being evaluated for treatment, but they may potentially have a preventive activity.

In summary, there are still no drugs to be used for primary prevention of asthma. There are, however, treatment possibilities for tertiary prevention of deterioration and exacerbations although effects on the long-term prognosis are still uncertain. Prevention of rhinovirus infections may provide a major step forward in primary prevention; immune modulators improve the innate immunity and response to virus infections; and there are other potential approaches.

sIL-4R Therapy

IL-4 is one of the most important cytokines in a variety of allergic diseases, including asthma.¹²⁶ IL-4 is produced primarily by CD4 Th2 cells but also by CD8 T cells, eosinophils, mast cells, and basophils. The role of IL-4 in human asthma has been established in several studies. The proof of the importance of IL-4 includes the following: (1) characterization of T-cell clones producing IL-4 in allergic and nonallergic individuals;¹²⁷ (2) *in situ* localization of IL-4-producing cells in bronchial biopsies;²³⁴ (3) expression of IL-4R mRNA and protein in epithelium, subepithelium, and endothelial cell layers in bronchial biopsies of atopic asthmatics compared with controls;¹²⁸ and (4) analysis of mutations of the IL-4R and their association with asthma.¹²⁹ The importance of IL-4 in the regulation of the allergic response, coupled with the identification of the soluble receptor, suggested that the sIL-4R might be effective as a therapeutic agent. Soluble IL-4R inhibited the IL-4-induced proliferation of B cells; the expression of low-affinity IgER and MHC class II; and the secretion of both IgE and IgG1 antibodies in mice.¹³⁰ In a murine model of allergen sensitization, soluble murine IL-4R

inhibited polyclonal and particularly antigen-specific IgE and IgG1 production following restimulation with these allergens.¹³¹ In addition to inhibiting Ig class switching and IgE production, murine sIL-4R reduces allergen-induced airway reactivity, vascular cell adhesion molecule-1 (VCAM-1) expression; allergen-induced eosinophilia in bronchoalveolar lavage (BAL); and allergen-induced pulmonary infiltration and airway occlusion by inflammatory cells. The safety of sIL-4R therapy is demonstrated by transgenic mice that produce a 100-fold greater sIL-4R concentration than nontransgenic littermates. Mice with increased sIL-4R compared with controls have similar numbers of B and T lymphocytes, lymphocyte surface marker expression, and antigen-specific antibody responses.¹³²

In clinical studies, soluble IL-4R blocked CD8 T-cell-mediated IgE production in allergic patients, suggesting that sIL-4R may be useful for treating allergic diseases.¹³³ Studies on the effects of recombinant humanized sIL-4R (rhusIL-4R) on IL-4/staphylococcal enterotoxin B-stimulated peripheral blood mononuclear cells from patients with eczema suggested that rhusIL-4R may be an immunomodulatory drug for atopic eczema.¹³⁴ In another study of human asthmatics, rhusIL-4R, was nebulized and given to 62 patients with moderate asthma at doses of 0.75, 1.5, or 3 mg. The study was double-blinded and placebo-controlled, and corticosteroid therapy was discontinued at entry. The results showed that sIL-4R was well tolerated, and the group treated with 3 mg sIL-4R demonstrated less labile FEV1 and improved asthma symptom scores compared to the placebo group.⁶³ In an open-label, randomized, dose-ranging study, the safety and tolerability of sIL-4R was evaluated in 16 adult patients with mild atopic asthma (FEV1 >70% of predicted). A single nebulized dose (50–100 µg) was generally well tolerated and pulmonary function improved.¹³⁵ In a phase I/II randomized, placebo-controlled trial,⁶³ 25 patients with moderate, inhaled corticosteroid-dependent asthma were randomly assigned to receive a single nebulized dose of IL-4R or placebo after discontinuing inhaled corticosteroids. The results indicate that sIL-4R is effective with once-weekly inhalation. The treatment is generally well tolerated and prevents a decline in FEV1, improves asthma symptom scores, and reduces β 2-agonist rescue following discontinuation of inhaled corticosteroids. Pulmonary inflammation, as assessed by exhaled nitric oxide, was significantly lower in the sIL-4R treated group compared to the placebo group. The treated group had nonstatistically decreased levels of VCAM-1, intercellular adhesion molecule-1, and EPO and increased CD23 expression compared to the placebo group. These results support the potential use of sIL-4R as a therapy for asthma and other allergic diseases, but additional studies are needed.

Targeting Tregs for IT of Allergy and Asthma

The fundamental role of Treg cells in maintaining immune tolerance has been demonstrated in a wide range of animal models, in which the adoptive

transfer or deliberate expansion of Treg cells was shown to prevent or cure several T-cell-mediated diseases, which include allergy, asthmatic lung inflammation, autoimmune diseases, and allograft rejection, by restoring immune tolerance to allergens, self-antigens, or alloantigens. Multiple molecular mechanisms for Treg-mediated immunosuppression have been described with secretion of IL-10 being of particular importance.¹³⁶ Absence or defective function of Treg cells has also been correlated with hyper-IgE syndrome, hypereosinophilia, and autoimmunity in humans, whereas their presence has been associated with immune tolerance.¹³⁷

Studies on the mechanisms by which immune responses to nonpathogenic environmental antigens lead to either allergy or nonharmful immunity have demonstrated that allergen-specific IL-10 producing Tregs (TR1 cells) are the dominant T cell subset in healthy individuals.^{138,139} Repeated exposure of nonallergic healthy beekeepers to bee venom antigens during the beekeeping season represents a valuable *in vivo* model to ascertain mechanisms of immune tolerance to venom antigens.¹⁴⁰ After multiple bee stings, venom antigen-specific Th1 and Th2 cells switch toward IL-10-secreting TR1 cells. This occurs in parallel to the suppression of cutaneous late-phase responses to allergens and inhibition of allergen-specific Th1 and Th2 cells. The response is observed as long as venom exposure persists and returns to initial levels within 2–3 months after the end of the beekeeping season. In the same model, the upregulation of H2R on specific Th2 cells suppressed allergen-stimulated T cells and increased IL-10 production. Various strategies, which are designed to enhance Treg function *in vivo*, are currently under investigation. These include the adoptive transfer of inducible or constitutive Treg cells and their induction by specific adjuvants or immunomodulators. These approaches are attractive compared to conventional treatments, as the antigen-specific suppressor capacity of Treg cells does not result in general immunosuppression and may actually lead to long-lasting antigen-specific regulation *in vivo*. Moreover, individual patient-specific treatments are possible with limited side effects. Many immunomodulators that have been developed or are under development, such as rapamycin, the CD80/CD86:CD28 costimulation blocker abatacept (Orencia; Bristol–Myers Squibb, Co., Uxbridge, Middlesex, UK), nonmitogenic anti-CD3 mAbs, and T-cell depletion and anti-TNF- α mAbs, display direct or indirect effects on Treg cells, which may enhance or suppress their function.^{141–143}

There is a selective advantage to expand a population of Treg cells that can target the organ (or the lymph nodes that drain the organ) by recognition of an allergen or an autoantigen expressed in inflamed organs in mouse models.¹⁴⁴ Thus, transfer of organ-specific Treg cells can be effective at suppressing ongoing disease, although those Treg cells do not necessarily need to recognize exactly the same autoantigen as the autoaggressive effector T cells.¹⁴⁵ This observation has implications for therapeutic strategies aimed at targeting the Treg cell arm of immune tolerance against allergens,

autoantigens, or transplantation antigens. Possibilities of adoptive transfer of Treg cells or small molecular compounds that induce Treg cells in the tissue are being investigated,¹⁴⁴ but no double-blind, placebo-controlled studies have been reported so far. To date, allergen-SIT is the only antigen-specific approach that induces Treg cell production and activation in humans. Allergen-SIT induces Treg and IL-10-secreting TR1-like cells, and treatment with glucocorticoids and β 2-adrenergic agonists seems to promote the number and activity of these cells.^{146–148} The essential transcriptional elements regulating expression of the Foxp3 promoter have been recently reported, and these will provide new targets for the development of novel therapeutics.¹⁴⁹

COSTIMULATION IN ALLERGY AND ASTHMA

The activation of T lymphocytes in the airways, which leads to production of the inflammatory cytokines and mediators characteristic of asthma, is clearly dependent on the presence of inhaled allergen *in situ*. However, allergen by itself is not sufficient to initiate this cascade of events. It is well established that the deleterious allergic response is initiated by T-cell recognition of MHC class II–peptide complexes at the surface of APCs. While this first signal gives antigen specificity to the adaptive immune response, a second non-specific costimulatory signal is required by T cells to become fully activated. This signal is provided by interactions between APCs and T cells through molecules borne at the surfaces of the two cell types. Depending on the type of molecules involved, this secondary signal can promote the development of an inflammatory allergic reaction or may favor immune regulation. Several molecules of the B7 family [CD80, CD86, programmed death-1 (PD-1), inducible costimulatory antigen (ICOS), CTLA-4] and TNF receptor (TNFR) family (OX40, CD30, 4-1BB, Fas, CD27, CD40) play an important role in delivering costimulatory signals in early and late phases of allergic response (Figure 4.3). Therefore, costimulatory molecules involved in promotion or prevention of allergic immune responses are potential targets for the development of novel therapeutic approaches. In order to become fully activated, T lymphocytes also require a so-called “second signal,” or “costimulatory” signal, which is provided to the T lymphocytes through interaction with cells specialized in antigen capture, or APCs, found in lymphoid organs as well as at mucosal surfaces such as the airway and lung.¹⁵⁰ In the absence of costimulatory signals, the T-dependent immune response is greatly diminished, or even eliminated altogether. Two costimulatory molecules expressed on the surface of APCs have so far been identified, termed CD80 and CD86. Both CD80 and CD86 mediate their function by binding a receptor on T lymphocytes termed CD28. Constitutive and induced expression of CD80 and CD86 varies on different APC types, and the reasons why APCs have two different versions of the same CD28 binding molecule are unclear. However, there have been some intriguing observations that atopic Th2-type immune

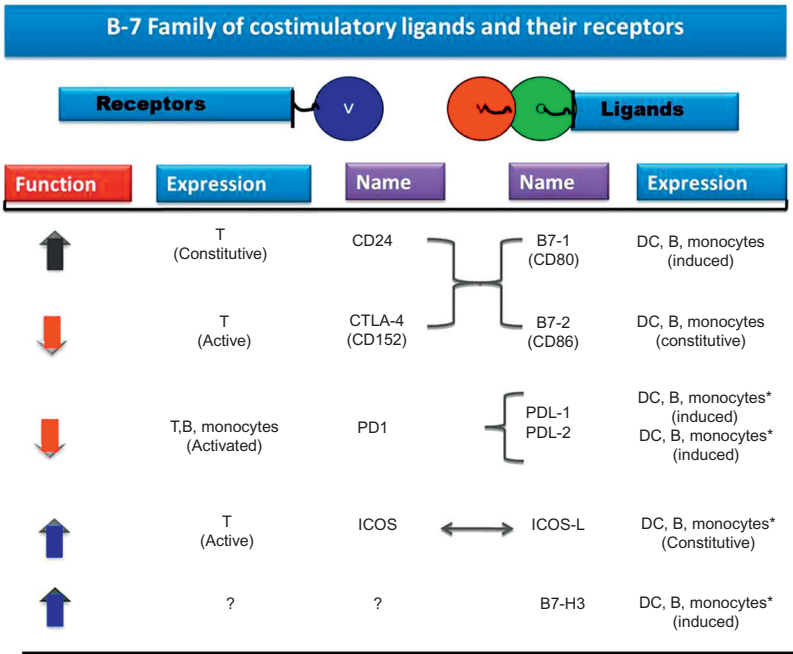


FIGURE 4.3 Summary of B7 family ligands and their receptors. The names of receptors and ligands are indicated, as well as a brief summary of predominant expression patterns for each. The conserved structure of a single IgV extracellular domain for receptors and IgV and IgC extracellular domains for ligands is depicted at the top. Function arrows indicate whether the pathway is thought predominantly to costimulate or inhibit the response of the receptor-bearing cell. Integration of signals through this family of costimulatory and inhibitory receptors and their ligands is critical for activation of immune responses and tolerance.

responses may be preferentially activated by CD86 while Th1-type immune responses are regulated by CD80.¹⁵¹ DCs in the lung and airway express both CD80 and CD86, as do pulmonary macrophages, although at much reduced levels.¹⁵² These cells therefore provide an effective source of costimulatory signals for the optimal activation of T lymphocytes locally.

Costimulation in Asthma

T-cell activation and cytokine secretion are important mediators of inflammation in allergic asthma.^{153,154} Recent studies indicate that the costimulatory pathway CD28/CD80/CD86 plays an important role in allergic asthma. The effect of other costimulatory molecules on allergic responses (e.g., OX40L) is under scrutiny. The roles of CD80 and CD86 have been examined by selective inhibition of either CD80 or CD86 in a murine model of allergic pulmonary inflammation.¹⁵⁵ Inhibition of costimulation by antibodies to either CD80 or

CD86 showed decreased allergic responses. Specifically, blockade of either CD80 or CD86 in allergen ovalbumin-sensitized and challenged mice resulted in a decrease in airway eosinophilia serum IgE production, airway hyperresponsiveness (AHR), and IL-4 and IL-2 production, with an enhanced IFN- γ production. Studies using mice with germline deletions of the CD80 and/or CD86 molecules showed that mice lacking both CD80 and CD86 had no elevation of serum IgE, airway eosinophilia, or AHR.¹⁵⁶ These same allergic parameters were also reduced in mice lacking either CD80 or CD86. Absence of CD80 and/or CD86 resulted in increased Th1 cytokine production. In humans, costimulatory molecule expression (CD28, CD80/CD86) is increased in allergic asthmatics.¹⁵⁷ These observations support a role for both CD80- and CD86-mediated costimulation in allergic pulmonary inflammation.

The costimulatory molecule OX40L, a member of the TNF superfamily, has been shown to be important in T-cell priming and cytokine production. In a murine model of allergic inflammation with OX40L-deficient mice, total serum IgE, pulmonary eosinophils, cytokines, and pulmonary inflammation were significantly decreased compared with wild-type controls.¹⁵⁸ Levels of eotaxin messenger RNA, an eosinophil-specific chemoattractant, were also markedly reduced, paralleling the significant reduction in pulmonary eosinophils. Levels of allergen-induced Th1 as well as Th2 cytokines were also significantly reduced. Together, the data support a critical role for OX40L signals in allergic responses.¹⁵⁹

Families of costimulatory molecules are involved in the regulation of most inflammatory diseases by finely controlling the intensity of the immune response. Costimulatory molecules are implicated in the development and control of allergic inflammation characterized by the establishment of an acute Th2 polarization. Elucidation of the role of costimulation pathways in the development of new subsets of Th cells has just begun, and most of the mechanisms underlying the regulation of atopic diseases by costimulatory molecules are unknown and require further investigation. With an increased understanding of these immunological mechanisms, new therapeutic strategies in the treatment of allergic airway diseases can be created by analyzing the role of costimulatory molecules that are critically involved in the induction and maintenance of allergen-induced airway diseases. Taken together, recent studies have begun to provide insight into the role of costimulatory molecules and give us new clues to design more efficient therapies to fight the increasing public health problem that allergies represent.

Recent studies have addressed the role of the costimulatory signal in animal models of asthmatic inflammation.¹⁶⁰ The simultaneous blockade of the signals delivered by CD80 and CD86 to T lymphocytes has dramatic effects on lung immune responses: infiltration of T lymphocytes is significantly diminished, and migration of eosinophils to the lung tissue and into the BAL fluid is also eliminated. Importantly, blockade of CD80 and CD86 only at the time of intranasal allergen challenge is sufficient to completely eliminate the

lung inflammatory response, which indicates this approach could be effective even after prolonged exposure to a specific allergen has taken place.

Unfortunately, as is the case for many other immunosuppressive treatments, the effect of blockade of CD80 and CD86 is not restricted to the lung, but also has systemic effects on antibody production and other unrelated immune responses in separate organs, which essentially induces a state of generalized immune suppression. The usefulness of this approach would therefore be limited to only the most severe cases of atopic asthma. In an attempt to address this concern, recent studies in our laboratory have examined the effect of selective blockade of the CD80 costimulatory molecule by using specific reagents that inhibit CD80 but leave CD86-dependent costimulation completely functional.¹⁶⁰ Rather surprisingly, selective blockade of CD80 costimulation has a dramatic effect on the lung inflammatory response and appears to almost completely block the local infiltration of eosinophils that is characteristic of asthmatic inflammation.¹⁶¹ At the same time, anti-CD80 treatment has no apparent effects on systemic immune responses, which leaves specific antibody responses intact, and also fails to inhibit the elevation of blood eosinophilia that is associated with allergen exposure. However, these eosinophils fail to migrate into the lung tissue upon antigen challenge, which implies that local mechanisms have been severely impaired due to unavailability of the CD80-dependent costimulatory pathway. The molecular mechanism of this finding has not yet been established, and is currently being addressed in our laboratory. We hypothesize that CD80 may provide additional costimulation, which is critical for the expression of mucosal immune responses, thereby acting as a safeguard against inappropriate immune activation. In this scenario, lack of CD80-dependent costimulation would result in incomplete activation of T cells in the lung, diminished production of IL-5 and/or other chemokines involved in the chemotaxis and migration of eosinophils into the tissues, and in turn, failure to activate a local inflammatory response.

Costimulation and T-Cell Reactivity to Allergens

The activation of T cells first requires a signal from the antigen in the context of the human leukocyte antigen (HLA) molecule and then an additional costimulatory signal involving the binding of B7 on APCs (CD80/86) to CD28 or CTLA-4 on T cells. Without costimulation, T cells become anergic. Thus, the costimulatory molecules have become targets for developing drugs that will modulate an allergic immune response. The chimeric CTLA-4-Ig molecule, which consists of the extracellular domain of CTLA-4 and the hinge, CH2, and CH3 regions of IgG, blocks the costimulatory signal that T cells receive through the interaction of CD28 with the counter-receptors CD80 or CD86 expressed on the APC membrane. CTLA-4-Ig is able to downregulate or suppress the mucosal immune response in allergic

rhinitis.¹⁶² CTLA-4-Ig inhibits airway eosinophilia and hyperresponsiveness by enhancing Th1-like cell activity. IgE formation is enhanced by the Th2 cytokine IL-4. Therefore, it is expected that downregulating Th2 activity will decrease IgE formation, as has been demonstrated in CTLA-4-Ig-treated mice.¹⁶³ The combination of antibodies to CD80/CD86 with the ligand for CTLA-4, blocks the development of allergic airway inflammation, whereas only a partial reduction occurs using either of these antibodies alone. However, only the anti-CD86 antibody inhibits the production of IgE.¹⁶⁴ Therefore, the CD28–CD80/CD86 complex is a potential target for developing new strategies of allergy treatment.

The first costimulatory molecules described were the ligands of CD28: CD80 (B7-1) and CD86 (B7-2). The CD28 costimulation pathway is an important factor for the promotion of an effective antigen-specific immune response. Costimulatory molecules are involved in the fine tuning of the T-cell response by mediating both stimulatory as well as inhibitory signals. In many recent studies, numerous new costimulatory molecules have been described leading to the recognition that costimulation pathways are more complex than the classical two-signal model. In general, costimulatory molecules are divided into two main families: molecules from the B7:CD28 family, such as CTLA-4 or PD-L1, and from the TNFR superfamily, such as OX40 or CD27.¹⁶⁵ All these costimulatory molecules have particular effects on T-cell activation, function, and survival (Table 4.1) and are implicated in nearly all inflammatory diseases. Studies to better characterize the specific role of these molecules in allergy and asthma are still ongoing. Recent investigations have uncovered the importance of a second costimulatory molecule on T cells termed CTLA-4 or CD152 that is also activated by CD80 and CD86 expressed on APCs.¹⁶⁵ In contrast to CD28, CTLA-4 delivers a negative signal to T cells, but because CTLA-4 is not normally expressed on the surface of T cells until 48 h after activation it is seen as a late-acting component that serves to shut off an immune response. Intriguingly, in experiments when this negative CTLA-4 signal was neutralized during an *in vivo* Th2 immune response, the level of Th2 cytokine production was greatly enhanced and the timing of production was advanced.¹⁶⁶ Preliminary investigations looking at T cells from atopic patients have also indicated that CTLA-4 is not expressed as highly as in healthy controls, as well as being delayed in its appearance after T-cell activation.¹⁶⁷ Such differences in CTLA-4 expression, between atopic and normal, may indicate an underlying genetic abnormality. These observations could indicate that T cells from atopic patients are not as sensitive to the normal inhibitory controls, which leads to prolonged immune responses with the potential for provoking disease. Even if the association between individual variation of CTLA-4 expression and atopy does not hold, it would certainly be worthwhile to explore the development of agonists of CTLA-4 activity attempt to terminate unwelcome atopic T-cell immune responses.

TABLE 4.1 The Role of Various Costimulatory Molecules in Allergic Diseases and Asthma

Costimulatory Family	APC	Effector Cell	Functions and Characteristics
B7 family	CD80 CD86	CTLA-4	Contributes to the suppressive activity of allergen-specific regulatory cells during sensitization
			Polymorphism in CTLA-4 promoter and gene favors allergic diseases
		CD28	CD80 and CD86 expression is upregulated on the surface of various cells of allergic patients
			Soluble CD86 is increased in the sera of allergic patients
	ICOS-L	ICOS	Regulates Th2-effector cell function and their infiltration in the lungs, production of Th2 cytokine
			Promotes B-cell differentiation and IgE production
			Expression on iNKT cells contributes to airway hyperreactivity
			Contributes to the differentiation of regulatory cells in pulmonary lymph nodes
	PD-L1	PD-1	Drives the differentiation of Foxp3+ CD4+ T cells
			Downregulates contact hypersensitivity reaction
	PD-L2	PD-1	Regulates asthma by an IFN-dependent mechanism
			Downregulates airway hyperreactivity, prevents eosinophil infiltration in the lungs, and prevents IgE production
B7-H3	Unknown receptor	Promotes Th2 differentiation, eosinophil infiltration, and development of airway hyperreactivity	
		Decreases the severity of allergic conjunctivitis	

Costimulatory Family	APC	Effector Cell	Functions and Characteristics
TNFR family	OX40L	OX40	Promotes the development of Th2 cells
			Prevents differentiation of Treg cells
			Abrogates mast cell degranulation
	CD30L	CD30	CD30 is expressed by Langerhans cells, CD4+, and CD8+ T cells of atopic patients
			Soluble CD30 is increased in the sera of allergic patients
	4-1BB-L	4-1BB	Promotes airway hyperreactivity, eosinophil infiltration, and IgE production
			Upregulates Th2 cell proliferation and mast cell cytokine production
	Fas	FasL	Delays resolution of airway hyperresponsiveness
			Promotes eosinophil apoptosis in the lungs
	CD27	CD70	Increases production of IgE by B cells
CD40	CD40L	Contributes to isotype class switching toward IgE	
		Enhances development of airway inflammation	
		Increases production of Th2 cytokine and decreases number of Treg cells	
Other costimulators	CD2	CD58	Promotes differentiation of Th2 cells and the production of IgE
			Expressed on monocytes of allergic patients

The expression of CD28 ligands (CD80 and CD86) has been extensively studied in clinical samples from asthmatic patients. It is well known that B lymphocytes from asthmatic patients exposed to allergens express higher levels of surface CD86, but not CD80, compared with those from asthmatic patients not exposed to allergen or with those from healthy individuals.¹⁶⁸ Another study demonstrated that CD80 and, to a lesser extent, CD86 were upregulated at the surface of alveolar macrophages from allergic patients compared with those from pulmonary sarcoidosis, extrinsic allergic alveolitis patients, or from normal subjects.¹⁶⁹ CD86 can also be expressed in a soluble form, where the transmembrane domain is deleted. This form is mostly produced by circulating monocytes and, like membrane-bound CD86, crosslinks CD28 or CTLA-4 and activates T lymphocytes.¹⁷⁰ In patients with acute asthma, the level of soluble CD86 has been shown to be increased relative to that in patients with stable asthma or in healthy individuals. Monocytes from allergic patients produce more soluble CD86 compared with those of healthy individuals.¹⁷¹ It was also observed that the level of soluble CD86 is correlated with the severity of the airway hyperresponsiveness (AHR). These results are consistent with studies that show that the concentrations of soluble CD80 and soluble CD86 are elevated in asthmatic patients. Interestingly, the administration of a glucocorticoid (e.g., prednisolone), used to reduce airway inflammation in allergic patients, reduces the level of circulating CD86.¹⁷² Recently, it has been shown that the topical administration of a silencer RNA specific to the CD86 gene reduced local inflammation in a mouse model of atopic dermatitis by decreasing the recruitment of DCs into the skin, production of antigen-specific IL-4, and induction of serum IgE and IgG1.¹⁷³ It has also been reported that pulmonary tolerogenic DCs stimulated by allergen exposure express high levels of both CD80 and CD86.¹⁷⁴ Taken together, these studies suggest that overexpression of CD80 and/or CD86 is correlated with the development of allergic disease and asthma.

WHAT MAKES COSTIMULATORY MOLECULES A PROMISING TARGET IN ALLERGY AND ASTHMA?

From studies in murine models of allergen-mediated airway inflammation, as well as from experiences of other immune modulators in allergy and asthmatic patients, some baseline conclusions may be drawn that characterize specific costimulatory molecules as optimal targets for therapeutic intervention (Table 4.2).

T-Cell Specificity

When targeting T cells, T-cell specificity is a major criterion, as simultaneous expression of the target molecule on anything other than T cells may lead to unforeseen side effects. An example of this was experienced in early

TABLE 4.2 Criteria for Optimal Targets in Allergic Diseases Through Costimulatory Molecules

Criteria	BTLA	CD27	CD28	CD30	CTLA-4	HVEM	ICOS	OX40	PD-1	SLAM	4-1BB
T-cell specificity	–	–	+	–	+	–	+	+	–	–	–
Nonconstitutive expression	–	–	–	+	+	–	+	+	+	+	+
Predominance for secondary immune reaction	–	–	–	+	–	–	+	+	+	–	–
Positive regulatory role	–	+	+	+	–	+	+	+	–	+	+
Th2 basis	–	–	–	+	–	ND	+	+	–	–	–

Note: BTLA: B- and T-lymphocyte attenuator; CTLA: cytotoxic T-lymphocyte-associated antigen; HVEM: herpes virus entry mediator; ICOS: inducible costimulatory antigen; OX40: CD134; PD: programmed death; SLAM: signaling lymphocyte activation molecule; Th2: type-2 T-helper cell; ND: not determined.

clinical trials blocking the CD40 ligand (CD40L) with mAbs. Blocking of CD40L in murine models was shown to attenuate IgE production,¹⁷⁵ although it had no effect on cellular infiltration of the airways. In patients, a mAb directed against CD40L (Ruplizumab) was first used for the treatment of systemic lupus erythematosus. However, the trials had to be discontinued because of life-threatening prothrombotic side effects due to expression of CD40L on activated platelets.¹⁷⁶ Therefore, T-cell restricted expression should be considered critical when identifying targets with which to modulate T-cell function. To date, this criterion is not met by the costimulatory molecules CD27, CD30, B- and T-lymphocyte attenuator (BTLA), and PD-1.

Expression on T Cells

Targeting a molecule that is expressed constitutively on T cells implies targeting all T cells unselectively. Nonselective depletion of T cells bears the risk of long-lasting lymphopenia and immune suppression, as experienced in clinical trials using anti-CD3 mAbs for immune suppression after organ transplantation.¹⁷⁷ Nonselective and strong stimulation of all T cells via a constitutively expressed costimulatory molecule can lead to cytokine release syndrome as seen recently with the anti-CD28 antibody TGN1412.¹⁷⁸ As these are unacceptable side effects in the treatment of allergic inflammation, the constitutively expressed costimulatory molecules CD27, CD28, BTLA, and herpes virus entry mediator do not qualify as optimal targets.

Predominant Involvement in Secondary Immune Reactions

In patients with bronchial asthma, allergic airway disease is an ongoing secondary immune response triggered by reexposure to specific allergens. A therapeutic intervention against asthma would preferably interfere with these secondary inflammatory reactions, while leaving primary immune reactions—critical in mounting a host defense against pathogens—unaffected. CD28, for instance, is the major costimulatory factor for the activation of naive T cells, and is therefore critical for the initiation of primary immune responses. In murine models, CD28 blockade was most effective when administered during allergen sensitization. This would render CD28 blockade an improper treatment strategy for human allergies because the time-point of sensitization is impossible to determine in the course of human allergic disease. However, blocking CD28 during secondary immune responses has shown some anti-inflammatory effects in mice. Moreover, clinical trials for the treatment of rheumatoid arthritis, employing the blockade of the CD28/B7 pathway with a CTLA-4-Ig fusion protein, have shown little apparent toxicity and good anti-inflammatory properties.^{179,180} In fact, this CTLA-4-Ig fusion protein, abatacept, is the first approved anti-costimulatory drug on the market.¹⁸¹ Two costimulatory molecules predominantly involved in secondary

immune reactions are ICOS and OX40. In contrast to CD28, ICOS expression is limited to currently activated T cells.^{182,183} In murine studies, ICOS-blocking reagents were most effective during secondary immune responses. Targeting OX40 in animal models of allergic airway disease was revealed to be effective during sensitization, as well as during rechallenge with allergen.

Positive Regulatory Signal

One major concern, besides compromising primary physiological immune reactions, is that negative signals delivered by costimulatory molecules would also be compromised by this approach. Blocking negative signals bears the risk of autoreactive immune responses and increased development of autoimmune diseases as a consequence of a disrupted negative feedback mechanism of immune balance. Therefore, this may disqualify negative costimulatory factors, such as BTLA and CTLA-4, as optimal treatment targets.

Th2 T-Cell Bias

The allergic airway inflammation is a Th2-dominated process, although recently, evidence for the involvement of Th1 cytokines, especially in established airway disease, has arisen. The predominant involvement of Th2-cytokines in airway inflammation makes these molecules a preferable target in the therapy of allergic reaction. This criterion is only met by the costimulatory molecules CD30, ICOS, and OX40. CD30, however, does not meet the criteria of T-cell specificity. ICOS is expressed on Th1 as well as on Th2 T cells, but only Th2-mediated airway inflammation is affected by ICOS-blockade. However, during an established inflammatory response like allergic asthma, this situation might be more complex than it appears in murine models of acute airway inflammation. ICOS is also expressed on Treg cells and may be important for the production of the immune suppressive cytokine, IL-10.¹⁸⁴ Even though a further study using a murine colitis model demonstrated that ICOS blockade had minimal effect on Treg function,¹⁸⁵ it remains to be determined how the immune balance is affected by long-term blockade of ICOS. A complete lack of ICOS expression in humans is associated with a loss of B-cell memory,¹⁸⁶ a side effect that may be desirable concerning local memory for IgE, but is definitely intolerable for any other systemic immune response. Therefore, the route of application and the place of deposition are important factors that need to be considered when targeting costimulatory molecules.

TARGETING COSTIMULATORY MOLECULES FOR TREATMENT OF ALLERGEN-INDUCED AIRWAY INFLAMMATION

Keeping in view the role played by costimulatory molecules in the regulation of immune response to allergens, costimulatory molecules have become

targets for developing drugs that will modulate an allergic immune response. The chimeric CTLA-4-Ig molecule, which consists of the extracellular domain of CTLA-4 and the hinge, CH2, and CH3 regions of IgG, blocks the costimulatory signal that T cells receive through the interaction of CD28 with the counter-receptors CD80 or CD86 expressed on the APC membrane. CTLA-4-Ig is able to downregulate or suppress the mucosal immune response in allergic rhinitis. CTLA-4-Ig inhibits airway eosinophilia and hyperresponsiveness by enhancing Th1-like cell activity. IgE formation is enhanced by the Th2 cytokine IL-4. Therefore, it is expected that downregulating Th2 activity will decrease IgE formation, as has been demonstrated in CTLA-4-Ig-treated mice. The combination of antibodies to CD80/CD86 with the ligand for CTLA-4, blocks the development of allergic airway inflammation, whereas only a partial reduction occurs using either of these antibodies alone. However, only the anti-CD86 antibody inhibits the production of IgE. Therefore, the CD28–CD80/CD86 complex is a potential target for developing new strategies of allergy treatment.

Even though not fully applicable to human disease, animal models provide valuable insight into the processes involved and are the basis for further research in humans. Due to the lack of human studies on costimulatory molecules in allergic airway disease, the present discussion is based on current knowledge as gained from animal models. In general, it is obvious from the experimental data that early blocking of allergic sensitization effectively inhibits the development of specific IgE, cytokine production by type-2 T-helper cells (Th2; mostly determined in BAL fluids), airway hyperresponsiveness (AHR), and inflammation (Table 4.3). The role of costimulatory molecules in allergen-induced airway inflammation may be classified according to the time of intervention as (1) primary intervention that targets the induction phase of the allergic sensitization; (2) secondary prevention, which targets the induction of airway inflammation in already sensitized animals; or (3) treatment of animals after sensitization and airway allergen challenges. The most intensively studied molecules in this context are CD28, ICOS, and CD134 (OX40), and because only preliminary data is available regarding other costimulatory pathways, these molecules are discussed in detail. See Table 4.3, which provides an overview of the available data regarding the role of costimulatory molecules in allergen-induced airway disease.

Targeting B7-CD28 Costimulatory Pathway

The role of CD28 in models of allergic airway inflammation has been extensively studied in allergen-induced airway inflammation murine models (Table 4.3). Although the primary goal of these studies was to analyze the impact of costimulatory molecules on different aspects of the inflammatory process, for example, eosinophilic airway inflammation, Th2 cytokine production, and AHR and systemic IgE synthesis, these data can also be used to

TABLE 4.3 Experimental Evidence from Blocking of Costimulatory Signaling in Allergen-Induced Airway Disease in Murine Models

Approach	Target	Outcome			
		Eosinophilia in BAL	Pulmonary Infiltration	AHR	IgE
Prevention of sensitization	CD80/CD86	↓	ND	↓	↓
	CD80/CD86	ND	↓	↓	ND
	CTLA-4	↑	ND	↑	↑
	ICOSL	ND	↓	ND	ND
	ICOS	ND	↓	ND	↑
	ICOS	(↓)	↔	↔	(↓)
	OX40L	↓	↓	↓	↓
	CD30L	ND	↓	↓	↓
Prevention of airway inflammation	CD80/CD86	↓	↓	ND	↓
	CD80/CD86	↓	ND	↓	↓
	CD80/CD86	↓	ND	↓	↓
	CD80/CD86	↓	ND	↓	↓
	CD80/CD86	↓	ND	↓	
	CD80/CD86	↔	ND	↔	↔
	CD80/CD86	↓	↓	ND	↔
	CTLA-4	(↓)	ND	↔	↓
	ICOS-L	↓	↓	↓	ND
	ICOS-L	↓	↓	ND	ND
	ICOS	↓	↓	↓	↓
	OX40L	↔	↔	↔	ND
	OX40L	↓	↓	↓	↓
	Treatment of established airway inflammation	CD80/CD86	↓	ND	ND
ICOS-L		ND	↔	ND	↔
CD30		ND	↔	↔	↔
CD30-L		ND	↔	↔	↔

Note: BAL: bronchoalveolar lavage; AHR: airway hyperresponsiveness; Ig: immunoglobulin; ND: not determined; CTLA: cytotoxic T-lymphocyte-associated antigen; ICOS: inducible costimulator antigen; ICOS-L: ICOS ligand; OX40: CD134; OX40L: OX40 ligand; ↓: decreased; ↑: increased; (↓): weak increase; ↔: unchanged.

assess the suitability of CD28 as a therapeutic target in the treatment of allergic airway inflammation. Systemic administration of the fusion protein cytotoxic T-lymphocyte-associated antigen-4 (CTLA-4)-Ig inhibits the ligation of CD28 with its ligands, CD80 and CD86. When given during the course of allergen sensitization, a drastic effect on airway inflammation and AHR was observed; however, the production of the Th1 cytokine interferon (IFN- γ) was unaltered.¹⁸⁷ Some of these effects may be explained by the complete inhibition of the germinal center reaction after blockade of the CD28/CD86 signal during the primary immune response.¹⁸⁸ When CTLA-4-IgG was applied after the initial T-cell sensitization, that is, during the phase of allergen rechallenge, this treatment also led to a marked reduction in the inflammatory response.¹⁸⁹ Only one report has shown contrasting results with no effect on AHR and cellular infiltration following the administration of CTLA-4-Ig at the time of allergen challenge.¹⁹⁰ Thus, interruption of the CD28 costimulatory signal also inhibits secondary immune responses, in at least some protocols of murine allergen-induced airway inflammation, but has the strongest impact on primary immune responses upon the first exposure to allergen.

In other studies, mAbs against the two ligands of CD28, CD80, and CD86 were administered independently to delineate the role of these molecules. Overall analysis of these studies reveals that CD86 appears to be the major ligand responsible for CD28-dependent immune responses observed during allergen-induced airway inflammation (Table 4.3). This is also implied by the observation that a combination of antibodies against both ligands was not superior to anti-CD86 treatment alone in two out of three studies. The question of a differentiated role of the two CD28 ligands was further addressed by the use of a fusion protein, Y100F. This molecule blocks CD28–CD80 interactions, leaving CD28 signaling via CD86 intact. Similar to CTLA-4-Ig, application of Y100F reduced eosinophilic infiltration into the lungs, mainly due to decreased IL-5 productions by allergen-specific T cells. In contrast to blocking all CD28 signals, Y100F had no effect on the number of eosinophils in peripheral blood or on systemic IgE production, indicating a major role of CD86 signals in local immune responses. In somewhat different experimental settings, ligation of CD28 by the CD28 ligand CD80 has been shown to have either no, an increasing, or a decreasing effect on allergen-induced airway responses.^{160,191–193} Thus, the precise role of this CD28 ligand in the T-cell differentiation remains an unanswered question.

CTLA-4 has been described to be an important regulator of T-cell activation and is constitutively and exclusively expressed by T lymphocytes in both mice and humans. Contribution of CTLA-4 in the regulation of the immune system is demonstrated by the development of multiple organ autoimmune pathologies and lympho-proliferative disease in CTLA-4-deficient mice.^{194,195} Blockade of CTLA-4 activity abolishes the suppressive function of CD4+ CD25+ T cells.¹⁹⁶ In inflammatory bowel disease, the effect of

transfer of a population of CD4⁺ CD45RB^{low} Treg cells in decreasing intestinal inflammation is abrogated by the coadministration of a blocking anti-CTLA-4 antibody.¹⁹⁷ This suggests that the engagement of CTLA-4 at the surface of Treg cells by its ligands CD80 or CD86 contributes to the regulation of suppressive functions of Treg cells. Polymorphism in CTLA-4 gene is also considered a risk factor for allergy and asthma. Four single nucleotide polymorphisms have been related to allergic and asthma phenotypes.¹⁹⁸ They demonstrated that these specific polymorphisms alone or in combinations are correlated with an elevated IgE titer or bronchial hyperresponsiveness in patients with asthma. Interestingly, in the same study, no correlation between allergic phenotype and single nucleotide polymorphisms for CD28 was observed. The polymorphism at the level of the promoter was correlated to asthma severity while the +49 C/G polymorphism is associated with airway hyperresponsiveness.^{199,200} Interestingly, CTLA-4 seems to play a more important role in the sensitization phase than in established allergy. In a model of mice sensitized with grass pollen, the administration of a blocking anti-CTLA-4 or a blocking anti-CD154 (anti-CD40L) antibody during the sensitization phase prevents the production of allergen-specific antibody. These findings confirm that CTLA-4 is indeed involved in the course of allergic diseases.

Targeting PD-1, PD-L1/PD-L2 Costimulatory Pathway

The other B7:CD28 family of receptors includes the PD-1 receptor and its ligands PD-L1 (B7-H1) and PD-L2 (B7-DC). The PD-1 receptor was initially discovered in T cells undergoing cell death. The inhibitory signal provided by engagement of PD-1 was demonstrated by the development of autoimmune diseases in PD-1-deficient mice.²⁰¹ Since then several groups have reported that engagement of PD-1 by PD-L1 or PD-L2 results in inhibition of proliferation and polarized or altered cytokine production.^{202,203} In asthma it has been demonstrated that PD-L2 is highly expressed on pulmonary DCs and macrophages of sensitized mice. Moreover, administration of blocking antibody against PD-L2, but not PD-1 or PD-L1, during challenge enhances the airway hyperresponsiveness and production of Th2 cytokines. This effect is mediated by IFN- γ , given that no improvement is observed in IFN- γ -deficient mice following treatment with anti-PD-L2.^{204,205} The administration of the sHIgM12 antibody (an antibody inducing reverse signaling through PD-L2) in a mouse model of allergic asthma blocks the development of AHR.²⁰⁶ An additional study demonstrates that administration of PD-L2-Fc in a mouse model of allergic asthma resulted in elevated levels of serum IgE and increased eosinophilic and lymphocytic infiltration into the BAL fluid.²⁰⁷ These studies emphasize the pivotal role of PD-L2 in the development of allergic asthma. In addition, in a model of experimental allergic conjunctivitis, treatment with anti-PD-L2 blocking antibody during the effector phase enhanced

infiltration of eosinophils into the conjunctiva without change in the systemic response.²⁰⁸ Finally, using PD-L2-deficient mice, it was reported that this molecule is dispensable for Th2 differentiation and required for the induction of mucosal tolerance.²⁰⁹ Taken together, these reports strongly suggest that PD-L2 is involved in the downregulation of Th2-allergic immune response.

Only preliminary data exist on manipulation of the negative regulatory molecule PD-1: an analysis of the distinctive contribution of its two ligands, PD-L1 and PD-L2, in the development of allergen-induced murine airway inflammation showed that a substantial number of pulmonary DCs, macrophages, and B cells isolated from the lungs of naive mice expressed PD-L1, which was further enhanced after allergen sensitization and airway challenge.²⁰⁵ In contrast, PD-L2 expression was detectable at very low levels in lymphocytes of unchallenged mice and was only moderately enhanced in DCs and macrophages after allergen challenge of sensitized mice. Although PD-L1 was abundantly expressed on various APCs in the inflamed lung tissues, treatment with a blocking antibody did not alter the allergic airway response, whereas treatment with anti-PD-L2 mAb (TY25) during allergen challenge significantly increased the development of AHR, and lung eosinophilia increased levels of IL-5 and IL-13 in BAL fluid.²⁰⁷ When PD-L2 was only blocked during sensitization, no effect on the inflammatory response was demonstrated, indicating that the inhibitory function of PD-L2 was limited to the effector phase (airway challenge) of the airway response.

Anti-PD-L1 blocking antibody enhanced contact hypersensitivity reaction, possibly by increasing the proliferative response of T cells in response to hapten-pulsed APCs. This suggests a unique role of PD-L1 in the regulation of inflammatory responses.²¹⁰ During allergen-SIT, the expression of PD-L1 on both monocytes and B lymphocytes is increased relative to that in the untreated control group. So PD-L1 could be used as a marker to monitor the effect of allergen-SIT and could also be targeted to enhance immunosuppression.²¹¹ Upon stimulation by a TLR-4 ligand, PD-L1-positive tolerogenic Langerhans cells in the sublingual mucosa in humans release a higher level of IL-10 compared with untreated control cells. These Langerhans cells have decreased capacity to stimulate T cells and are able to support the differentiation of Treg cells expressing Foxp3, producing IL-10 and TGF- β .^{212,213} These sublingual tolerogenic Langerhans cells stimulated via their TLR-4 expressed higher levels of the coinhibitory molecules PD-L1 and B7-H3, while CD86 expression is lowered. Consequently, the expression of these molecules by APCs seems to be linked with tolerogenic properties. These studies represent a body of evidence that suggests that PD-L1 is involved in the maintenance of the peripheral tolerance and may contribute to the induction of allergy. Clearly, further work is required to understand the role of PD-1 and its ligands in allergic diseases and asthma.

Targeting ICOS

In contrast to blocking CD28, blockade of ICOS by the application of ICOS-Ig or anti-ICOS mAbs at the time of allergen sensitization, showed only little effect on the development of airway inflammation (Table 4.3). However, blockade of ICOS in sensitized mice significantly reduced signs of allergic airway inflammation, such as increased IgE and Th2 cytokine production.¹⁹⁰ Using a model of adoptive transfer, it was further demonstrated that blockade of ICOS distinctively inhibited Th2-mediated lung eosinophilia and AHR, but did not abrogate Th1-mediated neutrophilic airway inflammation.²¹⁴ Similarly, the Th2-associated airway inflammatory response to *Schistosoma mansoni* eggs was attenuated by ICOS blockade, whereas the priming of T cells toward the Th2 cell direction was abolished by this approach.²¹⁵ The differential role of ICOS in Th1- versus Th2-biased immune reactions was further analyzed in a model of allergen-induced primary immune reactions induced by the local application of Th1 and Th2 cytokines. In this model, blockade of ICOS inhibited the early inflammatory cell influx and IL-5 production in the lungs in the Th2, but not in the Th1, model.²¹⁶ Taken together these results render ICOS a promising target to interfere with secondary immune responses, and suggest a predominant involvement of ICOS in Th2-type responses.

Targeting OX40

Utilizing OX40 knock-out mice, it was demonstrated that OX40 plays an important role in effector cell expansion and the formation and reactivation of memory T cells. Blocking the OX40 signal with mAb against OX40L in a model of *Leishmania major* infection resulted in decreased synthesis of Th2 cytokines (IL-4, IL-10 and IL-13) as well as reduced production of IgE.²¹⁷ In mouse models of allergen-induced airway inflammation there are somewhat controversial results regarding the time-point of OX40 action. Blockade of OX40L abolished the development of airway inflammation only when the anti-OX40L mAb was applied during allergen sensitization but not when given to already sensitized mice prior to allergen airway challenge.²¹⁸ In contrast, another study demonstrated that OX40L blockade of sensitized mice during allergen airway challenges abolished inflammatory responses, also at very late time-points of the experimental protocol.²¹⁹

Targeting BTLA-4

Very limited information is available on the effects of blocking other costimulatory molecules in murine models of allergen-induced airway disease. To date, no studies investigating the blockade of BTLA or B7-H3 have been published. Recently, it was shown that a single injection of mAb against the costimulatory factor CD137 (4-1BB) prevented the development of AHR,

eosinophilic airway inflammation, and elevated IgE production. This treatment was also able to reverse previously established airway disease. The inhibitory effect was most likely due to reduced Th2 cytokine production and increased secretion of IFN- γ by CD8 $^{+}$ T cells. Recently, blockade of CD30 and its ligand CD153 with mAbs was shown to significantly reduce airway inflammation, AHR, and the production of allergen-specific IgE when the blocking antibody was administered before and after sensitization.^{220,221} However, neither blocking reagent had any effect when administered during established allergic airway disease.

TOOLS FOR MODULATION OF COSTIMULATORY SIGNALS

Not only is the identification of the optimal target molecule a prerequisite for effective treatment by interfering with costimulatory molecules, but also careful considerations need to be made about the tools, administration, and the immunological aim of successful intervention into an ongoing inflammatory process.

Blocking costimulatory signals can either be achieved using mAbs or fusion proteins containing the counter-receptor. Half-life and tissue distribution of the blocking reagent within the human body are critical variables that will determine the duration of the blocking effect and the frequency required for application. The first insights in the human system have been gained by the use of anti-CD3 mAbs in organ transplantation and the CTLA-4 fusion protein for rheumatoid arthritis. Data from these and other studies show that using humanized mAbs is, in general, safe and does not bear the risk of anaphylactic side effects. Due to the complexity of the allergic immune-response and the asthmatic airway-response it is highly likely that targeting a single molecule may not be sufficient to gain complete control of allergen-mediated inflammation. For murine models of allograft rejection, it was shown that dual blockade with mAbs was more effective than blockade of a single molecule.²²² In this respect, a feasible approach may be to target not only one, but two (e.g., ICOS and OX40), or even more costimulatory molecules to treat asthma (Figure 4.2). However, such studies are still outstanding for allergic disease models. The consequences of using agonistic mAbs may be far more difficult to predict than those with blocking reagents. While dosing antagonists is stoichiometric, the action of agonists is a function of the strength of the agonist, the cascade it induces and the consequences of that cascade.²²³ In-depth knowledge of receptor distribution and action, as well as intensive animal testing, are critical, as demonstrated by the TeGenero fiasco with an agonistic anti-CD28 antibody.

Administration of Immune Modulators

Besides systemic administration of an immune modulator, local deposition of a blocking reagent in the lungs should be considered a feasible route of

application. During allergic airway inflammation, the number of T cells producing Th2 cytokines is increased in the pulmonary compartment.²²⁴ Therefore, it seems reasonable to directly target this process in the lung to avoid systemic side effects. Indeed, in mouse models of allergen-induced airway disease, local application of mAbs or fusion proteins into the airways was shown to control allergic airway inflammation, as demonstrated for the IL-13 receptor, a fusion protein, and anti-IL5 mAbs.^{225,226} In humans, one study showed that administration of an inhaled monoclonal anti-IgE antibody (E25) was generally well tolerated and led to detectable levels of the antibody in both BAL and serum.²²⁷ However, the aerosolized antibody did not attenuate the airway response to inhaled allergen as it did when applied intravenously. The authors speculated that the aerosol route of application did not lead to sufficient concentrations of the antibody in the critical tissue compartments surrounding IgE effector cells. Furthermore, one subject developed IgG and IgA antibodies against E25, suggesting that lung deposition of the antibody may be more immunogenic than the parental route. Therefore, local deposition, tissue penetration, serum levels and half-life, as well as immunogenicity, are critical variables that need to be considered when evaluating this route of application.

Blockade Versus Elimination of Allergen-Specific T Cells

An interesting alternative to continuous application of a blocking agent would be the permanent elimination of allergen-specific T cells from the immune system.²²⁸ This would require a target that is T-cell specific and not expressed on naive or quiescent memory T cells, but rather is expressed on recently activated T-effector or reactivated memory cells. From the group of costimulatory molecules, ICOS and OX40 meet these criteria. Elimination of all ICOS- and OX40-positive T cells during an ongoing allergic reaction may remove the allergen-specific T-effector and reactivated memory T cells, subsequently leading to downregulation of humoral responses and inflammatory cell infiltration, which both depend on T-cell help. The aspired benefits of such an approach would be a long-lasting therapeutic effect achieved by only a short-term treatment. Furthermore, this sort of treatment would be available and effective for allergic sensitization against all possible allergens, in contrast to SIT. However, the feasibility of this approach remains to be tested in appropriate animal models.

The underlying immune mechanisms of allergic airway disease have been untangled via the use of mouse models and clinical studies. The time has now come to widen the repertoire of therapeutic tools and to initiate immunological and pathogenesis-related approaches to treatment. One of the most promising new strategies is based on the blockade of costimulation, a critical and early event in the induction and maintenance of allergen-induced airway diseases such as bronchial asthma. So developing costimulatory molecules for the intervention in allergic and asthmatic diseases holds the promise for treatment in the future.

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Costimulation in Lymphomas and Cancers

INTRODUCTION

Cancer is a major health problem worldwide and is one of the most prominent causes of morbidity and mortality in children and adults causing about 9 million deaths annually. The transformation of normal cells to cancer cells may arise due to dysregulation of oncogenes, tumor suppressors, and/or stability genes. These transformed cells are sensed by the cells of the immune system, especially T cells, through specific receptors for an effective immune response. But unfortunately even after the interaction with T cells, an effective immune response is not generated. The success of novel cancer therapies depends on the identification of functional targets that play an essential role in tumor growth and metastasis, survival, and evasion from immunosurveillance. Cancer and the immune system are fundamentally interrelated. Cancer cells express tumor-specific aberrant antigens and must therefore evade immune detection to survive, either by inducing immunosuppression or deriving survival signals from tumor-infiltrating immune cells. T-cell-mediated antitumor immunity requires recognition of cancer-associated antigen by the major histocompatibility complex (MHC), and appropriate costimulatory and repressive secondary signals arising from complex interactions with other immune, stromal, and tumor cells. Dysfunction of costimulation pathways may contribute to failed antitumor immunity.

Although many costimulatory molecules like CD80, CD86, CD40, heat stable antigen, intercellular adhesion molecule (ICAM-1), vascular cell adhesion molecule (VCAM-1), B7-DC/programmed death ligand-2 (PDL-2), B7-H1/programmed death ligand-1 (PDL-1), etc., are known to be expressed on the surface of antigen-presenting cells (APCs). But the best-defined costimulatory molecules are two structurally related proteins known as CD80 (B7-1) and CD86 (B7-2).¹ Both CD80 and CD86 have their ligands, CD28 and cytotoxic T-lymphocyte associated antigen (CTLA-4), expressed on T cells. The delivery

of costimulatory signals is necessary to ensure an effective immune response against the tumors. It has been established recently that CD80 and CD86 play a role in the activation of APCs. Stimulation via CD86 in B cells can modulate their proliferation; IgG secretion and expression of proapoptotic and antiapoptotic molecules; nuclear localization of NF- κ B (p50) subunit; phosphorylation of v-rel reticulo endotheliosis viral oncogene homolog A (Rel A) (p65) and I κ B-alpha; and increased octamer-binding transcription factor 2 (oct-2) expression.^{2,3} It has been now established that CD80/CD86 contact with CD28 and CTLA-4 delivers signals that activate and inhibit T-cell activity, respectively. Furthermore, in the case of dendritic cells (DCs), it has been shown recently that it enhances the production of IL-6 and IFN- γ which in turn, up-regulates the expression of the immunoregulatory enzyme indolamine 2,3-dioxygenase (IDO) that results in tryptophan catabolism, suppresses T-cell responses, and promotes tumor resistance. Reverse signaling by CTLA-4 through CD80/CD86 in DCs causes the activation of the transcription factor NF- κ B followed by suppressor of cytokine signaling 3 (SOCS3) down-regulation and IFN- γ release. This in turn causes the induction of IDO and tryptophan catabolism, hence its depletion, and kynurinine production, which produces the biological effect.⁴⁻⁶ It has also been shown that silencing of SOCS3 by small interfering RNA (siRNA) makes CD28-Ig capable of activating IDO hence in the absence of SOCS3 CD28-Ig becomes immunosuppressive and mimics the action of CTLA-4-Ig on tryptophan catabolism. These effects appear to result from a combination of unopposed IFN- γ signaling and the occurrence of IFN- γ like actions by IL-6. However, in the case of CD80, it has been illustrated that signaling through this molecule can arrest the growth of lymphomas by upregulating expression of proapoptotic molecules and downregulating antiapoptotic molecules (see [Figure 5.3](#) and [Figure 5.4](#)).⁷⁻¹¹

Major Costimulatory and Coinhibitory Pathways in Antitumor Immunity

Costimulatory molecules belong to two major families: B7/CD28 family and tumor necrosis factor (TNF)/TNF receptor (TNFR) family (see [Table 5.1](#)). All molecules belonging to B7/CD28 family are members of the larger immunoglobulin superfamily (IGSF) and are involved in the triggering of cell-mediated immune response. Instead, the TNF/TNFR family members are involved in the later phases of T-cell activation and are induced from hours to days following the T-cell receptor (TCR) engagement. Currently the B7 family has seven members: B7.1, B7.2, ICOS-L, PD-L1, PD-L2, B7-H3, and B7-H4. While as TNFR/TNF family has six members herpes virus entry mediator (HVEM), B- and T-lymphocyte attenuator (BTLA), CD70, CD30, 4-IBBL, and OX40L.^{9,10} Presence of an efficient costimulation is crucial for improving antitumor immunity. In fact, one of the mechanisms through which a tumor is able to evade immune surveillance is the lower expression

TABLE 5.1 Members of B7 and TNFR Family of Costimulatory Molecules and Ligands

Costimulatory Families	Superfamily	Expression	Costimulatory Molecule	Ligand	Signal Type
TNF/TNFR Superfamily	TNF/TNFR	Constitutive	CD27	CD70	Positive
	TNF/TNFR	Inducible	OX40 (CD134)	OX40L	Positive
	TNF/TNFR	Inducible	4-1BB (CD 137)	4-1BBL	Positive
	TNF/TNFR	Inducible	CD30	CD30L (CD153)	Positive
TNF/TNFR, CD28/B7 Superfamily	TNF/TNFR, CD28/B7	Constitutive	HVEM	LIGHT, BTLA	Positive
	TNF/TNFR, CD28/B7	Constitutive	BTLA	HVEM	Negative
CD28/B7 Superfamily	CD28/B7	Constitutive	CD28	B7-1, B7-2 (CD80, CD86)	Positive
	CD28/B7	Inducible	CTLA-4 (CD152)	B7-1, B7-2 (CD80, CD86)	Negative
	CD28/B7	Inducible	ICOS	ICOS-L	Positive
	CD28/B7	Inducible	PD-1	PD-L1, PDL-2	Negative
	CD28/B7	Inducible	Unknown	B7-H4	Negative
	CD28/B7	Inducible	Unknown	B7-H3	Obscure
CD2 Family	CD2	Inducible	SLAM (CD150)	SLAM (CD150)	Positive

Note: TNF, tumor necrosis factor; R, receptor; HVEM, herpes virus entry mediator; LIGHT, homologous to lymphotoxin inducible expression, competing for GpD of herpes virus, expressed on activated T lymphocytes; BTLA, B- and T-lymphocyte attenuator; ICOS, inducible costimulator; L, ligand; SLAM, signaling lymphocyte activation molecule; CTLA, cytotoxic T-lymphocyte antigen; PD, programmed death; B7-H4 and H3, B7 homologues 4 and 3.

of costimulatory molecules or the upregulation of coinhibitory molecules. The lack of costimulation in the tumor microenvironment could be responsible for the generation of anergic T cells and, consequently, the absence of an appropriate antitumor immune response.¹⁰

The B7 system of costimulatory molecules is one of the most important secondary signaling mechanisms and is essential in maintaining the delicate balance between immune potency and suppression of autoimmunity (summarized in [Table 5.2](#)). Potential therapeutic applications include immune-boosting adjuvants to conventional anticancer therapy, hematopoietic stem cell transplantation (HSCT), antitumor vaccines, bioengineered T cells as well as attenuation of graft-versus-host disease (GVHD). The B7 family is only one aspect of a complex signaling network (explained in [Figures 5.1 and 5.2](#)) that comprises other IGSF members, the TNF superfamily (TNFRSF), chemokines, cytokines, and adhesion molecules. However, based on a substantial evidence base and a growing therapeutic armory, the B7 family requires particular attention, as summarized in [Table 5.3](#). Therefore manipulation of signals delivered by B7 ligands shows a great potential in treatment of cancers. So interfering with the costimulatory/inhibitory ligand-receptor network that regulates the immune response has become a promising approach in cancer immunotherapy. Thus, antagonistic antibodies directed against inhibitory receptors of the immunoglobulin-like (Ig) superfamily (e.g., CTLA-4, PD-1) and agonistic antibodies directed against costimulatory receptors of the TNFR superfamily (e.g., 4-1BB, OX40, GITR, CD127, CD40) are being evaluated in clinical trials.¹⁰

Targeting T Cell Costimulatory Molecules to Improve Antitumor Immunity

The tumor microenvironment is often characterized by the presence of anergic T cells, due to the lack of positive costimulatory molecules, mainly B7-1 and B7-2, on the surface of cancer cells.²⁷ One strategy to reverse this scenario is to force B7 expression on tumor cells, rendering them able to activate T-cell immune response. Several studies showed that the induction of B7-1 on tumor cells was sufficient to stimulate the CD8+ T-cell-mediated rejection in several tumor models, as well as a memory response, but was insufficient to mediate rejection of poorly immunogenic tumors.¹⁰ Several phase I clinical trials evaluated the efficacy of B7-1 transfected tumor cell vaccines, with or without IL-2, with encouraging preliminary results in patients affected by metastatic renal carcinoma and nonsmall cell lung cancer^{28,29} ([Table 5.3](#)). In a phase II trial, 39 patients with metastatic renal carcinoma were vaccinated with B7-1 transfected autologous tumor cells in combination with systemic IL-2. The authors observed a decrease in the cancer.³⁰

TABLE 5.2 B7 Family Members Update and Their Nomenclature

B7 Designation	CD Designation	Molecule/Eponyms	Receptor	Expression Pattern	Functions
B7-1	CD80	B7-1 or B7	CD28/CTLA-4 (CD152)	DCs, activated macrophages and B cells	Activation of naive T cells, T-cell proliferation, IL-2 production, T-cell survival/inhibition of T-cell responses
B7-2	CD86	B7-2 or B7	CD28/CTLA-4 (CD152)	DCs, activated macrophages and B cells	Activation of naive T cells, T-cell proliferation, IL-2 production, T-cell survival/inhibition of T-cell responses
B7-H1	CD274	Programmed cell death Ligand-1 PDL-1	CD279 (PD-1) ?	DCs, monocytes, macrophages, B cells, activated T cells, endothelial cells and tumor cells	Effector T-cell functions, T-cell proliferation, cytokines (IL-10, IFN- γ) production, apoptosis of CTLs DCs and monocytes, inhibition of T-cell responses
B7-H2	CD275	Inducible costimulator Ligand (B7-h, B7RP-1, LICOS/ICOS-L, GL50)	ICOS (CD278)	DCs, monocytes, Langerhans cells, B cells, fibroblasts, endothelial cells, etc.	Effector T-cell functions, T-cell proliferation, Ig isotype class switching, cytokine (IFN- γ , IL-10, IL-4) production
B7-H3	CD276	B7-Homologue 3	?? ILT-2 (orphan ligand)	Induced on DCs and monocytes	T-cell proliferation, IFN- γ production, enhancement of CTL activity

(Continued)

TABLE 5.2 (Continued)

B7 Designation	CD Designation	Molecule/Eponyms	Receptor	Expression Pattern	Functions
B7-H4	CD276	B7-Homologue 4	BTLA (??)	Induced on T cells, B cells, DCs, Macrophages	Inhibition of T-cell proliferation, cytokine production, cell cycle arrest
B7-H6	CD276	B7-Homologue 6	NKp30	NK T cells	Costimulation, NK T-cell activation
B7-DC	CD273	Programmed cell death Ligand-2 (PDL-2)	PD-1 (CD279)	DCs, monocytes, macrophages, B cells, activated T cells, endothelial cells, and tumor cells	Effector T-cell functions, T-cell proliferation, cytokines (IL-10, IFN- γ) production, apoptosis of CTLs, DCs and monocytes, inhibition of T-cell responses, regulations of DC biology
–	CD277	BT3.1, Butyrophilin SF3 A1, BTF5	?	T, B, NK, mono, DC, endothelial, CD34 + cells, tumor cell lines	T-cell activation

Expression of selected antigens expressed on the cell surface of APC or tumor cells and their costimulatory or inhibitory ligands on the surface of T cells or NK cells.
Note: ? means unknown; and ?? means evidence is contested or based on limited data.

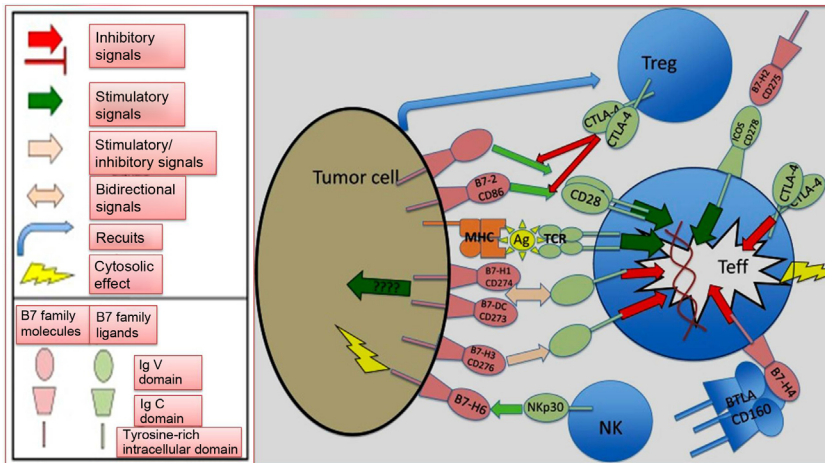


FIGURE 5.1 Expression of B7 family members and their receptors have complex interactions with the tumor immune environment.

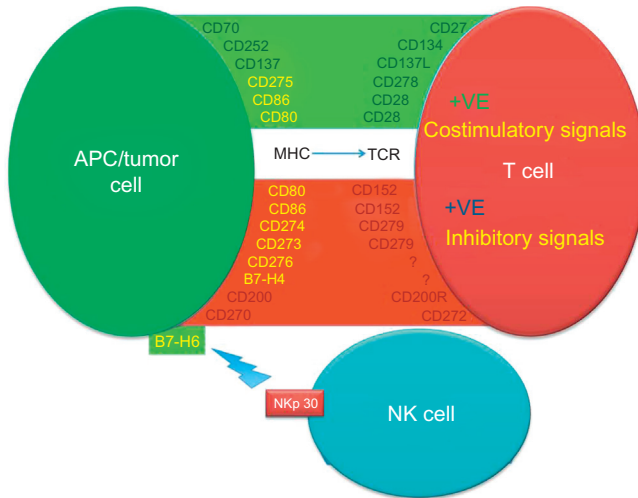


FIGURE 5.2 Expression of selected antigens expressed on the cell surface of APC or tumor cells and their costimulatory or inhibitory ligands.

B7-1 was also included in vaccine along with specific antigens, such as carcinoembryonic antigen (CEA) (Table 5.4). Recently, in a phase II trial for metastatic colorectal cancer, Kaufman et al. used a nonreplicating canarypox virus vector (ALVAC) expressing CEA and B7-1 in combination with chemotherapy. The observed objective response rate of 40.4% was similar to

TABLE 5.3 Therapeutic Potential of the B7 Family Molecules in Hematological Malignancies

B7 Target	Costimulatory Pathway	Preclinical Evidence Reference	Phase I/II Trial Agent
Augmentation of systemic antitumor immunity	CTLA-4 blockade	[13]	Ipilimumab
	PD-1/PD-L1 blockade	[14,15]	BMS936558, BMS936559, MK-3475
	ICOS pathway modulation	[16]	NA
Tumor-directed therapy against B7 targets	CD80	[13]	Galiximab
	PD-L1	[17,18]	NA
	B7-H6		NA
Augmentation of GVL	CTLA-4 blockade	[19]	Ipilimumab
	PD-1/PD-L1 blockade	[20]	NA
	ICOS pathway modulation		NA
Attenuation of GVHD	CTLA-4 stimulation		CTLA-4.Ig
	CD80/86 blockade		CD80/86 mAb
	ICOS pathway modulation	[21,22]	NA
Bioengineered T cells	CD28 CAR	[23]	Modified T cell
	CD80 CAR	[24]	Modified T cell
	NKp30 CAR		Modified NK cell
Antitumor vaccine augmentation	CTLA-4 blockade	[13]	Ipilimumab
	PD-1/PD-L1 blockade	[25,26]	NA
Treg depletion and inactivation	CTLA-4 blockade		NA

Note: mAb, monoclonal antibody; NK, natural killer; CTLA, cytolytic T-cell-associated sequence-4; PD, programmed cell death; ICOS, inducible costimulator; CAR, chimeric antigen receptor; and NA, there are currently no therapeutic agent available.

TABLE 5.4 Tumor Models and Clinical Trials of B7-CD28 Costimulatory Molecules

Tumor Model	Costimulatory Molecule	Therapeutic Strategy	Reference
Metastatic renal carcinoma	B7-1 (CD80)	Vaccination with B7-1-transfected autologous tumor cells in combination with systemic IL-2	[28–30]
Nonsmall cell lung cancer	B7-1 (CD80)	Vaccination with an adenocarcinoma cell line expressing B7-1 and human leukocyte antigen A1 (HLA-A1) or A2	[29]
Metastatic colorectal cancer	B7-1 (CD80)	Vaccination with ALVAC vector expressing CEA and B7-1 in combination with chemotherapy	
CEA-expressing carcinoma, metastatic carcinoma, prostate cancer	B7-1 (CD80)	Vaccination with TRICOM vector	[31–33]
Metastatic melanoma, metastatic renal cancer, nonsmall cell lung cancer	CTLA-4	MDX-010 Ab blockade of CTLA-4 alone or in combination with vaccine, IL-2, and chemotherapy	[34,35]
Melanoma, metastatic colorectal cancer, advanced gastric cancer, and esophageal adenocarcinoma	CTLA-4	CP-675,206 Ab blockade of CTLA-4 alone or in combination with chemotherapy	[36–39]
Hematological malignancies	PD-1	CT-011 Ab blockade of PD-1	[40]
Advanced solid cancer	PD-1	MDX-1106 Ab blockade of PD-1	[41]
Solid tumors	PD-1	MK-3475	[42]
Cancers, multiple indications	PD-1	MDX-1105-01	[42]

Note: CTLA, cytotoxic T-lymphocyte antigen; PD-1, programmed death 1; B7-1 means CD80.

that reported for chemotherapy alone.³¹ Improvements on new vaccine strategies led to the generation of viral vectors expressing a triad of costimulatory molecules (TRICOMs), such as B7-1, ICAM-1, and lymphocyte function-associated antigen 3 (LFA-3), along with CEA, mucin-cell-surface associated 1 (MUC-1), and prostate-specific antigen (PSA) antigens, with promising results in preclinical studies and clinical trials, both in terms of efficacy and safety^{32–34} (Table 5.4). In this regard, a phase II randomized controlled trial of poxiviral-based PSA-targeted immunotherapy in patients with metastatic castration-resistant prostate cancer showed that the treatment was well tolerated and associated with 44% reduction in the death rate.⁴³ Recently, multitarget vaccine approaches were tested *in vitro*, resulting in enhanced antitumor immune response against hepatocellular carcinoma and glioma cell lines.^{44,45} Several other preclinical evidences proved the efficacy of B7-1-based therapeutic strategy in the induction of tumor antigen-specific T-cell response.

B7 SUPERFAMILY OF COSTIMULATORY MOLECULES IN ANTITUMOR IMMUNOTHERAPY

B7-1/B7-2:CD28/CTLA-4

The B7-1/B7-2:CD28/CTLA-4 pathway is the best characterized pathway of T-cell costimulation and coinhibition and symbolizes the classical way where the ligand can bind two receptors for regulating both T-cell activation and tolerance. B7-1 (CD80) and B7-2 (CD86) are two ligands for both CD28 and CTLA-4. The balance between the activating and inhibitory signals derived from the engagement of CD28 and CTLA-4, respectively, is crucial to assure protective immunity, without falling into undesired autoimmunity. The expression of B7-1 and B7-2 is restricted to professional APCs, such as DCs, macrophages, and B cells. CD28 is constitutively expressed on the surface of T cells, whereas CTLA-4 expression is induced 24–48 h after T-cell activation, due to the action of lymphocyte cell kinase (Lck), Fyn, and resting lymphocyte kinase that phosphorylates CTLA-4, thus increasing its transport to the cell surface and preventing its internalization. CTLA-4 was shown to have higher affinity for both B7-1 and B7-2 than CD28 receptor.¹⁰

The B7-1/B7-2:CD28 pathway is the strongest costimulatory signal delivered by APCs to provide a full activation of T cells, promoting their proliferation and IL-2 secretion. The intracellular signaling of B7-1/B7-2:CD28 pathway occurs through the activation of phosphatidylinositol-3-kinase (PI3K)/Akt/nuclear factor- κ B (NF- κ B) and the mitogen-activated protein kinases (MAPKs) pathway, which support cell survival, memory development, proliferation, and cytokines production.⁴⁶ In contrast to the costimulatory signal derived from the binding of CD28 to B7-1 and B7-2, the engagement of CTLA-4 by these ligands provides a negative regulation of the immune response, as proved by the characterization of CTLA-4 deficient mice.

The tumor microenvironment is often characterized by the presence of anergic T cells, due to the lack of positive costimulatory molecules, mainly B7-1 and B7-2, on the surface of cancer cells.²⁷ One strategy to revert this scenario is to force B7 expression on tumor cells, rendering them able to activate T-cell immune response. Several studies showed that the induction of B7-1 on tumor cells was sufficient to stimulate the CD8 + T-cell-mediated rejection in several tumor models, as well as a memory response, but was insufficient to mediate rejection of poorly immunogenic tumors.¹⁰ Several phase I clinical trials evaluated the efficacy of B7-1 transfected tumor cell vaccines, with or without IL-2, with encouraging preliminary results in patients affected by metastatic renal carcinoma and nonsmall cell lung cancer (Table 5.3).

Although effective as monotherapy in the treatment of small and immunogenic tumors, a combination of CTLA-4 blockade with other immunotherapeutic strategies is needed to treat large and poorly immunogenic tumors. Combination of CTLA-4 blockade and irradiated tumor vaccine expressing granulocyte/macrophage colony-stimulating factor (GM-CSF) results in tumor rejection and reduction of tumor growth in the B16 melanoma model. Similar results have been reported for the poorly immunogenic SM1 mammary carcinoma line and a transgenic model of prostate carcinoma.^{10,47} Moreover, the combination of anti-CTLA-4 with DNA vaccine increased T-cell immune response against melanoma-associated antigens and induced B16 tumor rejection.⁴⁸ In the same tumor model, the CTLA-4 blockade combined with CD25 + Treg depletion and vaccination was reported to be effective in inducing tyrosinase-related-protein-2-(TRP-2-) specific CD8 + T cells and in rejecting larger tumor loads.⁴⁹ An increased antitumor immunity in B16 melanoma model was reported following CTLA-4 blockade with peptide vaccine and a synthetic oligodeoxynucleotide as adjuvant.⁵⁰ The use of anti-CTLA-4 along with radiotherapy led to the improvement of survival rate in a mouse model of metastatic breast cancer, whereas CTLA-4 blockade in combination with chemotherapy provided clinical benefits in the murine myeloma model MOPC-315.^{51,52}

CD80 in Lymphoma Growth Inhibition

The ability of the B cells to deliver the costimulatory signal to T cells by B7 molecules is very well established.^{53,54} In contrast, whether the engagement of CD80 and CD86 molecules by CD28 and CTLA-4 affect the function of the B cells is very poorly documented.^{55,56} Moreover, nothing is known precisely in the case of B-cell proliferation and differentiation by the engagement of CD80 and CD86 molecules. CD80 and CD86 serve as counter-receptors that transduce distinct signal to the APCs upon engagement by CD28 or CTLA-4. The intracellular domains of CD80 and CD86 are quite distinct and could mediate differential signal transduction. Such signaling could alter the B cells' ability to function as effector cells. However, there is

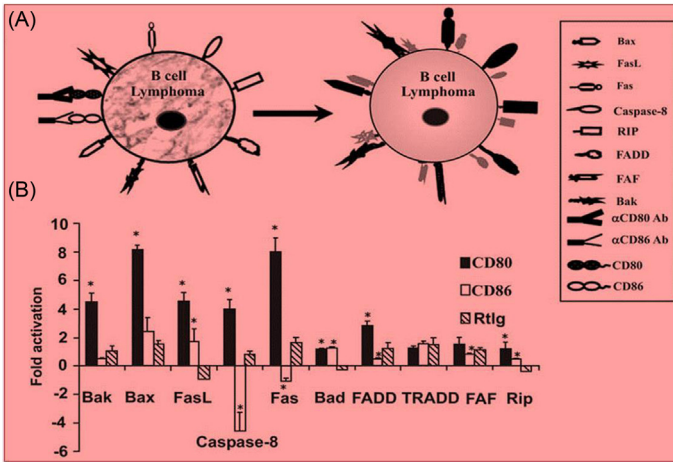


FIGURE 5.3 Modulation of the expression of proapoptotic molecules by CD80 and CD86 signaling.

indirect evidence reported earlier that CD28-CTLA-4/B7-signaling pathways may affect B-cell responses and the regulation of immunoglobulin (Ig) synthesis.^{57,58} Our study was aimed to evaluate for the first time the role of CD80 and CD86 mediated signal transduction in the costimulation of B cells. The results show that triggering through CD80 specifically inhibits the proliferation and IgG secretion by LPS1-stimulated B cells. By contrast, CD86 chiefly augments the B-cell activity. Further, the growth of B-cell lymphomas could also be retarded by CD80 cross-linking. Signaling through CD80 inhibited the growth of the B-cell lymphoma by upregulating the expression of proapoptotic molecules caspase-3, caspase-8, Fas, FasL, Bak, and Bax and downregulating the levels of antiapoptotic molecule Bcl-x[L]. In contrast, triggering through CD86 augmented the level of antiapoptotic molecules Bcl-w and Bcl-x[L] and decreased the levels of caspase-8. Thus, the signals delivered by anti-B7-1 and -B7-2 antibodies can differentially regulate the activity of B cell and its lymphoma (see [Figures 5.3 and 5.4](#)).^{1,2}

Costimulation signaling through CD80 in B-cell lymphomas can retard their proliferation by upregulating expression of proapoptotic molecules and downregulating antiapoptotic molecules and can therefore induce apoptosis. On the basis of our results recently, a phase I/II study of treatment with CD80-specific antibody has shown it to be quite effective in relapsed and refractory follicular lymphoma patients. Considering the importance of costimulation in the regulation of immune responses against relapsed cancer, the manipulation of this pathway to increase immunity represents a potential therapeutic approach. Furthermore, signaling by either anti-CD80 antibodies or through CD28-bearing T cells regresses the growth, augments the

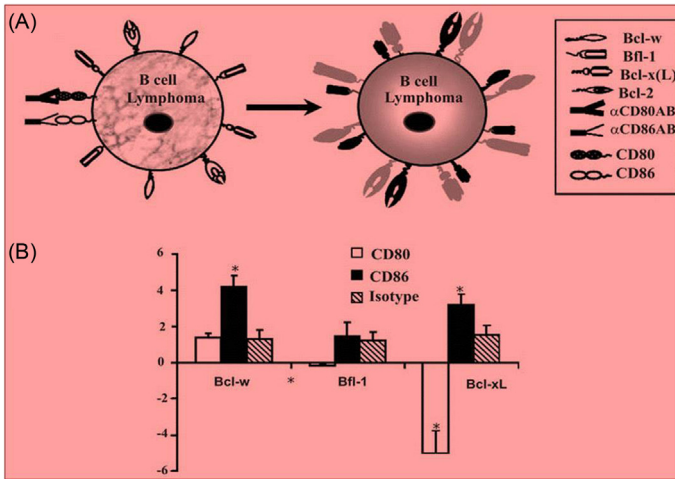


FIGURE 5.4 Modulation of the expression of antiapoptotic molecules by CD80 and CD86 signaling.

expression of proapoptotic molecules and induces apoptosis of CD80+ lymphomas. Therefore, immunotherapy utilizing anti-CD80 antibodies is a promising future treatment especially in the case of relapse and refractory lymphomas.^{59,60}

The exact mechanism of sequence of signals provided by different costimulatory molecules in the stimulation and inhibition of B cells is largely unknown. We for the first time have attempted to analyze the distinct role of CD80 and CD86 costimulatory molecules in the activation and differentiation of B cells. From our study five major findings emerged which are as follows: (1) cross-linking by anti-B7-1 mAb (16-10A1) inhibited the proliferation and differentiation of B cells; (2) anti-B7-1 and CD40 mAbs acted synergistically and declined the growth of B-cell lymphomas; (3) induction of secretion of IgG isotypes by WEHI-279 cells was triggered with anti-B7-2 mAb; (4) anti-B7-2 mAb (GL1) promoted the growth and differentiation of B cells; and (5) anti-B7-1 mAbs induced apoptosis in B-cell lymphoma by enhancing the expression of caspase-3, caspase-8, Fas, FasL, Bak, and Bax and decreasing the expression of antiapoptotic molecules Bcl-x[L].^{1,2,9} Our findings suggest that CD80 and CD86 costimulatory molecules are responsible for delivering the signals essential for the inhibition and expansion of B cells, respectively. To best of our knowledge, this was the first report regarding the role of CD80 and CD86 molecules in influencing the B-cell activity. To further provide weight to our findings of the regulatory capacity of anti-B7-1 and -B7-2 Abs in the activation of B cells, we utilized B-cell lymphomas, WEHI-279, which express CD80, CD86, and

CD40 molecules. More surprisingly, anti-CD80 and CD40 Abs could significantly arrest the proliferation of WEHI-279.² Thus it may be concluded that the signaling via CD80 and CD40 molecules may be responsible for inhibiting the growth of B-cell lymphomas. In contrast, costimulation through CD86 may help lymphomas to proliferate and secrete antibodies that may help lymphomas to evade immune surveillance. Further, triggering through CD80 and CD40 may help the regression of the lymphomas in the early stages. Although signaling through CD80 retards the growth of B-cell lymphomas by inducing the expression of Fas and FasL and in turn induces apoptosis, it fails to induce Fas and FasL or apoptosis in normal B cells.

At present, it is difficult to explain why ligation of B7-1 with anti-B7-1 has a differential effect on normal versus tumor B cells. Apart from demonstrating the expression of Fas and FasL by FACScan and their involvement in inducing apoptosis, we also monitored by ribonuclease protection assay the involvement of CD80 and CD86 molecules in the regulation of the activation of several pro- and antiapoptotic molecules in WEHI-279 B-cell lymphoma.² We have observed that signaling through CD80 molecule augmented the levels of proapoptotic molecules, that is, caspase-3, caspase-8, Fas, FasL, Bak, and Bax. This suggests that CD80 signaling induces apoptosis in WEHI-279 via a mechanism involving proapoptotic molecules caspase-3, caspase-8, Fas, FasL, Bak, and Bax, thus rendering WEHI-279 cells more vulnerable to apoptosis and therefore restricting the progression of the lymphoma. The T-cell costimulatory molecule CD28, which has a high and low binding affinity for CD86 and CD80, respectively, promotes T-cell survival by upregulating Bcl-x[L] and downregulating FasL expression.^{2,61} Based on these results and our findings, it may be postulated that there may exist a probability that, when CD28 is ligated with CD86 it may deliver a protective signal necessary for the clonal expansion of B cells. In contrast to this, when CTLA-4 gets engaged with CD80, it may deliver a lethal signal responsible for controlling the clonal expansion of B cell and B-cell lymphomas. In contrast, signaling through CD86 increased the expression of antiapoptotic molecules Bcl-w and Bcl-x[L]. Thus, there may be a possibility that ligation of B7-2 on WEHI-279 may promote their survival by increasing the expression of antiapoptotic proteins Bcl-w and Bcl-x[L]. It has also been reported earlier that increased expression of Bcl-x[L] but not Bcl-2 could prevent the apoptosis in B-cell lymphoma WEHI-231.⁶² Further, it has been reported that signaling through CD40 upregulated Bcl-x[L] and Bfl-1 and protected B-cell lymphoma from apoptosis.⁶³ Many studies have highlighted the lack of Costimulatory molecules as a predominant reason for the inefficient tumor rejection. Moreover, if the lymphomas express CD80, signaling through this molecule may impede their growth by inducing the expression of proapoptotic and downregulating the levels of antiapoptotic molecules and thereby facilitating apoptosis. It is important to mention here that CD80-mediated delivery of inhibitory signals may not only be

responsible for apoptosis of CD80-bearing tumor cells, but may also be responsible in the regulation of immune responses and maintaining self-tolerance in B cells. Thus, in the case of tumors, ligation of B7-1 by anti-CD80 Ab may also deliver effective antitumor immunity. In conclusion, we report here that selective inhibition of B-cell proliferation and differentiation via CD80 signaling may be viewed as a novel strategy to restrict the undesired stimulation of B cell and progression of B-cell lymphomas. Moreover, costimulation through CD80 and CD86 can modulate the humoral response by transducing positive and negative signals in B cells and may control the progression of B-cell lymphomas.

Costimulation Through CD80 in Immunity Against Cancer

CD80 is a costimulatory molecule known for its role in T-cell activation and also in regulating normal and malignant B cells activity.² Surface CD80 is expressed transiently on activated B cells, macrophages, and DCs. Surprisingly, CD80 is downregulated on most of the cancer cells and the loss of CD80 alone is sufficient to allow them to escape the attack of the immune system and to impart energy and apoptosis in tumor-infiltrating T cells.⁶⁴ In the absence of costimulation, recognition of antigens by T cells may not cause any response, even if tumor cells express MHC molecules and tumor-specific antigens (Figure 5.5). Hence most human malignancies that lack CD80 expression have been suggested to evade immune surveillance and therefore contribute for failure of immune recognition.⁶⁵ However, follicular lymphomas express CD80 and therefore could potentially be targeted by CD80 immunotherapies. Lack of either CD80 or MHC-I must be sufficient to allow tumor cells to evade immune response but it has been shown in some tumors that high expression of MHC-I and absence of costimulatory

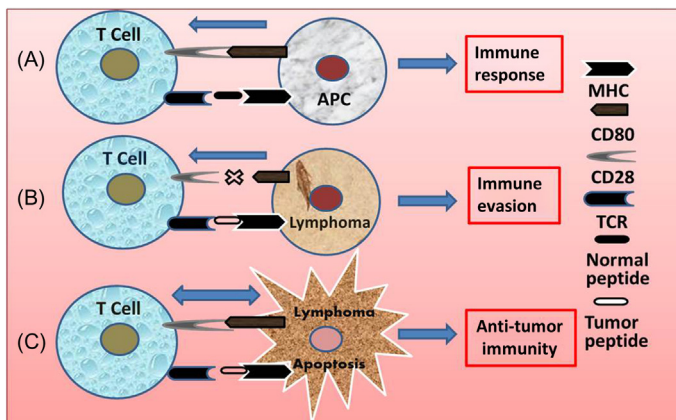


FIGURE 5.5 Reverse costimulation through CD80 in modulating the activity of cancer cells.

molecules renders them resistant to lysis by cytotoxic T lymphocytes (CTLs) (Figure 5.5A). It has been shown that high expression of MHC-I but lack of CD80 does not allow cytolysis of target cells by CTLs *ex vivo*. In contrast, CD80-transfected tumor cells become susceptible to lysis by CTLs *ex vivo*. Although priming of antitumor CTLs can also take place in absence of CD80, but the effector T cells are not produced without the expression of CD80 on tumor cells.⁶⁶ Interestingly, CD80 has also been known to enhance the memory response in CTLs.⁶⁷ This suggests that CD80 expression on a tumor is important for antitumor CTL effector function. Expression of CD80 on tumor cells confers direct presentation capacity to prime naive CTLs *in vivo*.⁶⁸ Transfection of tumor cells with CD80 will help in generating long-lasting immunity. Therefore, this approach may lead to the development of a successful vaccine against cancer.

CD80 expression is needed for the sustained predominance of CD8 + T cells within a tumor (Figure 5.5C). Transfection of CD80 cDNA into human erythro leukemia DR+ (HEL-DR+) cells induce the allogeneic response of purified T cells from both cord blood and peripheral blood of adult donors, demonstrating that CD80 expression could lead to accessory-cell-independent activation of naive T cells. Furthermore, in the P815 tumor system, CD86 was substituted for CD80, interestingly, no specific CTL activity was observed. Even though in the tumors that express CD86 there is a need for CD80 expression by host APCs for efficient eradication. Thus, it is apparent that transfection of CD80 in tumor cell serves as a co-factor for IL-2 production and in prevention of anergy by TCR ligation.⁶⁹ This suggests that expression of CD80 on cancer cells contributes towards the activation of T cells responsible for imparting anticancer immunity. The expression of CD80 on tumor cells also enhances natural killer (NK) cell recognition and lysis of tumors, which plays an important role in tumor immunity.⁷⁰ CD80 costimulation can direct the CD8 T cells to produce IL-2, proliferate and acquire cytolytic activity.⁷¹ This helper independent generation of CTLs may have practical application in the development of tumor-specific immunotherapy.⁷² So it is clear that downregulation of CD80 costimulatory molecule by various cancer cells acts as an effective immune evasion mechanism (see Figure 5.5B).

This has been reported in multiple myelomas (MM), carcinomas, leukemias, and transplantable malignancies, etc. Some of the colon carcinomas such as MC38, and melanomas such as B16, also lack the expression of CD80. This is evident from the fact that silencing of CD80 expression results in tumorigenicity and transfection of CD80 leads to decreased tumorigenicity.⁷³ CD80 expression triggers *in vitro* NK-cell-mediated cytotoxicity. The CD80 gene product functions as a triggering signal for NK-cell-mediated cytotoxicity and the strength of this response is such that it overrides the inhibitory signals mediated by MHC-I molecules.⁷⁴ The efficient control of solid allogeneic tumors by NK cells depends on co-delivery of both CD80 and MHC-I on the tumor cells. The co-delivery is required for optimal

expansion and effector function of NK cells in response to both melanoma and plasmocytoma that express allogeneic MHC-I. The two signals required for T-cell function can also regulate NK cell immunity and reveal an important similarity between the innate NK cell response and the adaptive T-cell response. Furthermore, it has been reported that NK cells and CD8⁺ T cells can eliminate CD80 expressing tumors that are resistant to IL-12 gene therapy.^{13,75} Hence expression of CD80 on tumor cells may also have an important bearing on NK cell effector function.

CD80 Gene Transfection of Tumor Cells to Generate Vaccine and Elicitation of Antitumor Response

The fact that CD80 plays an important role in immunity against cancer cells is now well established from the above discussion. Thus, it is apparent that expression of CD80 on cancer cells can enhance the immunity substantially (see [Figure 5.5B,C](#)). Expression of costimulatory molecule CD80 via somatic gene transfer represents one approach to improve the immunogenicity of tumor cells. Transfection of CD80 in colon cancer cells results in increased immunogenicity and tumor rejection. In contrast, silencing of CD80 expression in colon cancer cells resulted in lack of tumorigenicity.⁷³ The CD80 transfection into EL-4 cells, T-cell thymoma cell line, showed that CD80 plays an essential role in mediating *in vivo* antitumor rejection. Transfection of CD80 gene into various cancer cells, RMA, EL-4, P815, E6B2, B16, K1735, etc., has reduced or eliminated subsequent tumor growth in syngenic mice and can lead to the establishment of protective immunity against challenges with CD80-negative tumor cells.^{68,76} The positive effects of CD80 gene transduction have also been thoroughly explored in a rapidly lethal hematopoietic leukemia cell line, which is nonimmunogenic by itself. A tumor cell engineered to provide optimal costimulation would then directly induce T-cell activation, proliferation, and differentiation in to effector cells, following recognition of the antigen. CD80 gene transfection resulted in loss of tumorigenicity and induction of protective immunity against subsequent challenges with the parental CD80^{-/-} cell line in immunocompetent syngenic mice. CD80 transfection has been shown to inhibit lymph node metastasis by the mechanism of enhanced immunogenicity.⁷⁷ Moreover, not only the presence but also the levels of CD80 expression can affect the tumor growth.⁷⁸ The cells generated by introduction of the CD80 gene can work as prophylactic or therapeutic vaccines against tumors that lack CD80 by means of eliciting a strong CTL and NK cell response, which is desired in antitumor immunity.⁷⁹

The anticancer activity of the modalities, like γ -irradiation, mitomycin-C, melphalan, etc., is quite interesting that these exert their immuno-potentiating effect in tumor bearers by upregulating CD80 gene expression.⁸⁰ However, the mechanism by which surface expression of CD80 on tumor cells enhances the antitumor immunity seems to be the same as that which depends on the

maturation state of DCs regulating immune response by CD80 expression, as the low expression of CD80 on immature DCs suppresses T-cell immunity in a CD80-dependent fashion and high CD80 expression by mature DCs promotes immunity.⁸¹ Antitumor effects of CD80 expressing tumor cells can be mostly attributed to costimulatory activity resulting in activation, proliferation, and generation of CTLs. CD80 expression on tumor cells enhances their recognition by NK cells, hence causing enhanced lysis of tumor cells, which in turn makes the antigen available for the cross priming pathway, thereby generating systemic immunity to parental tumor. CD80 expression may protect T cells from activation-induced cell death upon recognition of tumor cell targets.⁸⁰ CD8+ T-cell precursors are directly primed by CD80-transfected tumors, which provide signal 1 and 2 to the T cells.⁷⁹ Indicating that, CD80-transfected cells serve directly as APCs *in vivo* for the priming of naive CTLs.

Keeping in view the importance of the CD80 costimulatory molecule in antitumor immunity against cancer, the manipulation of this molecule to increase immunity represents a promising therapeutic and prophylactic approach. CD80 has been co-administered with various cytokine genes to elicit antitumor activity. A single intratumoral co-administration of CD80 with the IL-12 gene, by electroporation, elicited specific antitumor CTL response and eradicated the tumor from mice. This co-administration also maintains the high level of endogenous CD80 of tumor cells through STAT-1 expression.⁸² The coexpression of CD80 and IL-12, administered by intratumoral injection with an adenovirus vector, effectively elicited antitumor immunity in a nonimmunogenic, therapy resistant, mouse pancreatic cancer model.⁸³ Recently it has been shown that primary tumor cells resected from cancer patients can be transfected effectively with CD80-SA (streptavidin) and that such transfected cells serve as APCs to induce autologous antitumor T-cell responses.⁸⁴ A dual modified tumor cell vaccine using CD80 and super antigen Staphylococcus enterotoxin A (a potent inducer of CTL activity and cytokine production) elicited significantly stronger antitumor immune responses *in vivo*.⁸⁵ When IFN- γ transfectant immunotherapy was used for cancer treatment; it promoted tumor progression. However interestingly, when a combination of IFN- γ and CD80 transfection immunotherapy was used, the cancer-promoting effect of cellular vaccination was completely abolished and instead the cancer vaccine potency to eradicate tumors increased.⁸⁶ Tumor cells transfected with CD80 and IL-7 genes induced CD28 and CD25 on tumor-infiltrating T cells and triggered strong antitumor responses. CD80 expression on tumor cells has been shown to eliminate resistance to IL-12 gene therapy. CD80 + MHC-II + -transfected tumor cells were used as a potent vaccine for stimulating tumor rejection in tumor bearing mice.⁸⁷ Transfection of CD80 gene into human hematopoietic malignant cell lines that are deficient in CD80 expression empowers their antigen presentation potency for activation of antitumor T cells. The combined expression of CD80 and IL-2 in acute myeloid leukemia blasts show increased

stimulation of both allogeneic and autologous T cells. The stimulated lymphocytes secrete higher levels of Th1 cytokines and show specificity.⁸⁸ Repairing the deficiency in the CD80 costimulatory pathway in tumor cells can be a novel immunotherapeutic approach for human hematopoietic malignancies.

Use of CD80 alone or in combination with other cytokines etc. has shown healthy results in controlling the growth of tumors or their recurrence once they are removed from the body. In patients with metastatic renal cell carcinoma, the administration of CD80 gene modified autologous tumor cell vaccine in combination with systemic IL-2 in phase I clinical trials have shown promising results.²⁸ Apoptotic uveal melanoma cells expressing CD80 were efficient at inducing an immune response and served as potent immunogens. A tri-cistronic viral vector co-expressing IL-12 (IL-12p40 plus IL-12p35) and CD80 showed enhanced CTL and proliferative response in a myeloma tumor model.⁸⁹ This strategy could prove to be a novel adjuvant immunotherapy for the disease.⁹⁰ CD80 on tumor cells can also interact directly with CD28 on NK cells and activate them to inhibit tumor growth.⁷⁹ It has also been shown that CD80 vaccination after removal of a tumor is useful for the prevention of tumor recurrence. The model of CD80 enhancing tumor immunity is that exogenous CD80 acts as a costimulatory molecule to directly interact with CD28 on T cells, thus resulting in the activation of CD8+ T cells against tumor cells.

Costimulation through CD80 induces apoptosis via a mechanism involving proapoptotic molecules, consequently rendering the cells more vulnerable to apoptosis and therefore restricting their proliferation. It has been shown that overexpression of caspases is sufficient to cause apoptosis. The proapoptotic molecule (Bax) has been correlated with disease regression and shorter survival of B cells in chronic lymphocytic leukemia (CLL).^{91–93} Resistance to apoptosis is also one of the mechanisms employed by tumor cells for evasion and it is not only relevant for tumorigenesis and resistance to chemotherapy but also influences immunosurveillance and immunotherapy. It is important to mention here that costimulatory molecule-mediated delivery of inhibitory signals may not only be responsible for the regulation of immune responses but also for apoptosis of CD80-bearing tumor cells. These studies categorically demonstrate that, tumor regression is not only the consequence of CD80-mediated activation of CTLs and NK cells but signaling through CD80 can also trigger death of lymphomas.^{94,95}

Costimulatory Monoclonal Antibodies in Non-Hodgkin's Lymphoma and Chronic Lymphocytic Leukemia

The development of monoclonal antibodies, biological therapies that target tumor-associated antigens specifically, gives hope for improvement of survival in many cancers. Monoclonal antibodies make an important contribution to the therapeutic treatment of non-Hodgkin's lymphoma (NHL), CLL, breast

cancer, and colorectal cancer. These antibodies can bind to their receptors on tumor cells specifically and block the binding of endogenous ligands, which leads to inhibition of phosphorylation of receptor-protein tyrosine kinases and inhibition of downstream signaling events.⁹⁶ Monoclonal antibodies that act through this mechanism include trastuzumab, which targets v-erb-b2 erythroblastic leukemia viral oncogene homolog 2 (ERBB2); cetuximab, which targets EGFR; alemtuzumab a lymphocyte-depleting humanized monoclonal antibody with efficacy in the treatment of chronic lymphocytic leukemia targeting CD52;⁹⁷ and bevacizumab, which blocks angiogenic signaling by binding to and sequestering the ligand of VEGF receptor (VEGFR).⁹⁸ Rituximab is a highly specific mouse/human chimeric IgG1 antibody that targets CD20, a cell-surface protein expressed on 95% of B-cell lymphomas, has shown efficacy in patients with various lymphoid malignancies, including indolent and aggressive forms of B cell NHL and B-cell CLL. It inhibits cell proliferation and directly induces apoptosis.^{99,100} It has been shown that patients can be treated safely and effectively with multiple courses of rituximab without induction of human antichimeric antibodies, cumulative myelosuppression, adverse hematological effects, and serious or opportunistic infections that are related to standard chemotherapy or radiotherapy.^{99,101} Galiximab is a chimeric, anti-CD80, IgG1 antibody that targets CD80, which is constitutively expressed on a variety of lymphomas like diffuse large B-cell NHL, CLL, Hodgkin's lymphoma, and MM. Because of the favorable safety profile of galiximab and its lack of induction of myelosuppression, which is associated with typical standard chemotherapies, radiotherapy and other antibody therapies, it is an attractive agent for lymphoma therapy.^{59,60} Galiximab has also shown promising results when used in combination with rituximab. These monoclonal antibodies bind to their targets specifically on the surface of cells and do not circulate in the plasma as free protein that could competitively inhibit antibody binding to lymphoma cells. Other properties of these monoclonal antibodies like long half-life, high specificity, and high safety standards compared with other cancer therapeutics together with their ability to bind to and modulate key players in signaling pathways that drive malignant transformation and enhance antitumor immune functions make them highly desirable therapeutic agents.

Anti-CD80 Antibody in Treatment of Lymphoma Patients

Over the past decade, the use of mAbs has emerged as an important component in treatment algorithms for patients with cancer, especially patients with B-cell NHL. The advantage of mAbs is that they provide targeted and effective therapy against cancer with less toxicity than standard chemotherapeutic agents. With the discovery of hybridoma technology in the 1970s the use of mAbs to treat patients with cancer has gained increasing in popularity.¹⁰² Initial the results of preliminary clinical studies were disappointing due to

limited antitumor activity. However, advances in tumor immunology and molecular biotechnology have led to the development of primatized, chimeric, and humanized mAbs with better antitumor activity and decreased immunogenicity.¹⁰³ The observation published by our laboratory on the induction of apoptosis on stimulation through CD80 has constituted a rationale for development of an anti-CD80 therapy for lymphomas. Preclinical studies indicate that targeting CD80 with anti-CD80 antibody has antitumor effects.^{2,104,105} Anti-CD80 antibody therapy delays progression of lymphoma and clinical studies have shown promising results.^{59,105,106} Recently, Biogene has developed an anti-CD80 antibody known as galiximab. Galiximab is a chimeric, anti-CD80, IgG1 lambda monoclonal antibody with human constant regions and primate (*Cynomolgus macaque*) variable regions. It is structurally indistinguishable from human antibodies and, therefore, is unlikely to be significantly immunogenic in humans. Some characteristics of the antibody are summarized in [Table 5.5](#).

This indistinguishability from human antibodies makes it more suitable for potential repeated dosing in lymphoma patients. The specificity of galiximab binding to human CD80 has been validated in competitive binding studies with several lymphoma cell lines. Preclinical *in vitro* and *in vivo* studies evaluated galiximab as a targeted therapy for lymphoma, with promising results. A phase I/II, single-agent, dose escalation, and multiple-dose study demonstrated that galiximab administration can decrease tumor burden in nearly half of all patients suffering from relapsed and refractory follicular lymphoma.⁵⁹ Treatment of relapsed/refractory follicular lymphoma with galiximab in an open-label, multicenter, phase I/II monotherapy trial in

TABLE 5.5 Characteristics of Anti-CD80 mAb (Galiximab) in the Treatment of Lymphomas

Characteristic Type	Property of Antibody
Antibody name	Galiximab or IDEC-114 or Anti-CD80 antibody
Type of antibody	Human–primate chimeric IgG1 λ
Dose	125–500 mg/m ² and half-life of 2–4 weeks in serum
Target	CD80 (B7-1) 60-kDa trans-membrane glyco-protein
Diseases	Follicular lymphoma, CLL, MM, etc.
Mechanism	Upregulation of proapoptotic and downregulation of antiapoptotic molecules, ADCC, complement-mediated cytotoxicity (CDC), NF- κ B modulation, etc.
Side effects	Fatigue, asthenia, nausea, headache, dizziness, vomiting, etc. mostly grade 1 and 2

TABLE 5.6 Clinical Results of Patients with Relapsed or Follicular Lymphoma after Treatment with Anti-CD80 Antibody, Galiximab

Reduction in Tumor Burden		Percentage of Patients Showing Most Common Clinical AEs				
Percentage Reduction in Lesions	Number of Patients (%)	Adverse Event Type	Grade 1	Grade 2	Grade 3	Grade 4
75–100%	17 (49%)	Lymph node pain	3	3	0	0
≥50%	6 (17%)	Dyspepsia	5	0	0	0
≥50% but with disease progression	2 (6%)	Vomiting	3	3	0	0
New lesions	1 (3%)	Axillary pain	3	0	3	0
Nonmeasurable disease progression	1 (3%)	Dysguesia	5	0	0	0
Not available for measurement	4 (12%)	Loose stools	5	0	0	0
Headache			5	5	0	0

37 patients with median age of 56.5 years who received different doses of antibody has shown no dose-limiting toxicities and a favorable safety profile. The most common related adverse events (AEs) experienced were grade one or two (headache, fatigue, and nausea) with only two cases of grade three deep venous thrombosis and one case of grade 3 axillary pain. Only one (3%) of 37 patients showed cytopenias, which was also considered to be unrelated to galiximab but related to the studied disease.¹⁰⁷ Clinical results of this study have been summarized in [Table 5.6](#).

Furthermore, in a study covering 242 patients receiving galiximab as part of multiple-dose psoriasis studies it has been demonstrated that its safety profile was similar to placebo: no T-cell or B-cell depletion was observed and none of the patients developed antigaliximab antibodies. Clinical results of this study have been summarized in [Table 5.7](#).^{104,106} Further a combination of galiximab and rituximab did not appear to increase the overall incidence of AEs or alter the AE profile. Studies in animal models of lymphoma have shown that a combination therapy of galiximab (anti-CD80) and rituximab

TABLE 5.7 Clinical Results of Patients with Moderate to Severe Plaque Psoriasis Showing Reduction in Psoriasis Area and Severity Index (PASI*) and Improvement in Physicians Global Psoriasis Assessment (PGA) after Treatment with an Anti-CD80 Antibody, Galiximab

Study days after treatment of patients with galiximab	15	29	36	43	50	57	71	99	127
Number of patients assessed after treatment	35	20	15	20	15	20	35	35	35
Percentage of patients showing $\geq 50\%$ reduction in PASI*	3	10	7	15	20	20	23	26	29
Percentage of patients showing $\geq 75\%$ reduction in PASI*	3	5	NA	10	NA	10	9	9	11

Note: *PASI is calculated in ill patients after treatment by visually assessing the extent of psoriatic involvement in four main body areas (head, trunk, upper extremities, and lower extremities) weighted by % of total body area and the severity of signs in those areas.

Taking all time points and all dose groups, 20 out of 35 patients (i.e., 57%) achieved a PGA rating of Good and seven patients (i.e., 20%) achieved a PGA rating of Excellent. NA: Not available.

(anti-CD20) has more clinical benefits, and when galiximab was combined with chemotherapeutic agents like fludarabine or doxorubicin, it enhanced the efficacy of anti-CD80 therapy significantly.¹⁰⁵ Recently, Leonard and co-workers have used a combination therapy of galiximab and rituximab for treatment of relapsed or refractory follicular lymphoma.⁵⁹ They found that this combination therapy does not alter the pharmacokinetics or immunogenicity but instead has the potential to avoid or delay chemotherapy or its associated toxicity or to amalgamate with other lymphoma therapies. Hence galiximab can be combined safely with a standard course of rituximab for treatment of relapsed or refractory follicular lymphoma. Based on these observations a phase III, randomized study has been initiated to evaluate the clinical benefits of rituximab therapy versus combination therapy.¹⁰⁷ A phase I/II study of galiximab in combination with rituximab therapy in patients with relapsed or refractory disease has shown promising preliminary results for clinical activity without added toxicity compared with single-agent rituximab alone.⁶⁰ The future plans in this direction include evaluating maintenance therapy with the combination treatment of galiximab and rituximab in follicular NHL and galiximab, rituximab, plus chemotherapy for follicular and diffused aggressive NHL.

Experimental and clinical results, make it quite evident that anti-CD80 antibody treatment is effective against relapsed and refractory follicular lymphoma. Future studies in other CD80-positive hematological malignancies (e.g., diffuse large B-cell NHL, CLL, Hodgkin's lymphoma, and MM) should also be taken under consideration.⁵⁹ Thus, anti-CD80 antibody immunotherapy may have a potent role in the treatment of CD80-bearing cancer cells because their binding can modulate the key molecules in the signaling pathway and enhance antitumor response. Considering the importance of CD80 signaling in the regulation of immune responses against cancer, the manipulation of this signaling pathway to increase immunity against cancer represents a potential therapeutic approach.

The FDA in 1997 approved rituximab an anti-Cd80 mAb for the treatment of patients with B-cell lymphoma and heralded the onset of a new era in cancer therapeutics.¹⁰⁸ The early use of rituximab as a single agent or in combination with systemic chemotherapy regimens has resulted in improved response rates and survival in patients with follicular lymphoma.^{109,110} Prior to the incorporation of rituximab into standard antilymphoma regimens, estimated median overall survival time for follicular NHL patients was 8–10 years from initial diagnosis.¹¹¹ With the possible exception of allogeneic stem cell transplantation, there is no curative therapy for advanced follicular NHL.¹¹² Therapeutic options currently in use include single-agent rituximab, single-agent or combination chemotherapy \pm rituximab, radio-immunotherapy, and involved-field radiotherapy (generally used for localized or chemotherapy-refractory disease). Commonly used single-agent chemotherapies include oral-alkylating agents and purine analogues. More aggressive combination drug regimens offer higher response rates than single agents, albeit at the cost of greater toxicity and no discernible improvement in long-term survival.^{104,113} Historically, it has been shown that “watchful waiting” instead of immediate therapy did not adversely affect outcome in appropriately selected patients.¹¹⁴ The fact that successful lymphoma therapies can be based on targeting pathways engaged in lymphomagenesis with less nonspecific toxicities than standard chemotherapy is a significant advancement in the treatment of B-cell lymphoma patients. Ongoing challenges in the treatment of patients with NHL consist not only in identifying biomarkers of response to specific “targeted” therapies, but also in the timely discovery and development of novel therapeutic agents with improved activity and decreased toxicity. In addition to rituximab, other mAbs targeting cell-surface antigen present on B-cell neoplasms are being actively studied in a large number of ongoing clinical trials.

Understanding of mechanisms by which tumor cells escape immune surveillance will help us to establish new and effective approaches to vaccination and immunotherapy. The success of novel cancer therapies depends on the identification of functional targets that play an essential role in tumor growth and metastasis, survival, and evasion from immunosurveillance. One of the most important immune evasion mechanism employed by tumor cells

is the downregulation of the CD80 costimulatory molecule. Anti-CD80 therapies can be used to target tumors expressing CD80. The clinical success of anti-CD80 antibody for the treatment of refractory follicular lymphoma has stimulated great interest in the promise of antibody therapeutics for cancer. The qualities of galiximab, like long half-life and high specificity and safety compared with other cancer therapeutics, together with its ability to bind to and modulate key players in signaling pathways that drive malignant transformation and enhance antitumor immune functions, makes it a highly desirable therapeutic agent.

Most tumors do not regress but continue to grow in spite of the presence of spontaneous or antigen induced immune responses, due to downregulation of costimulatory molecules like CD80. It is now well established that T cells are rendered anergic due to the lack of costimulatory molecule(s) expression by tumor cells. Tumor cells are able to induce antigen-specific tolerance or anergy on the basis of MHC-I-restricted antigen presentation without the expression of costimulatory molecules. This unresponsiveness, however, can be reversed when tumor cells are genetically modified to express costimulatory molecules. A plethora of studies suggest that the insertion of genes encoding CD80 into tumors generally increases their immunogenicity and can be used as vaccine.

Recently, fusogene vectors were developed to encode multiple gene products like CD80 with cytokines or MHC molecules as fusion proteins from a single cistron to increase the immunogenicity of target tumor cells. The vectors generated could be used in immunotherapy for the treatment of MM, leukemias, and other cancers, as they have been shown to stimulate allogeneic mixed lymphocyte proliferation and augment increases in CTL and NK cell responses and IFN- γ release. Considering the importance of costimulation in the regulation of immune responses against relapsed cancer, the manipulation of this pathway to increase immunity represents a potential therapeutic approach. Furthermore, signaling by either anti-CD80 antibodies or through CD28-bearing T cells regresses the growth and augments the expression of proapoptotic molecules and induces apoptosis of CD80+ lymphomas. Therefore, immunotherapy utilizing anti-CD80 antibodies is a promising future treatment especially in the case of relapse and refractory lymphomas. Clinical studies have demonstrated that administration of anti-CD80 antibody can decrease tumor burden in patients suffering from lymphomas.⁵⁹ CD80 is expressed transiently on the surface of activated B cells and other APCs, including DCs, but is constitutively expressed on a variety of NHLs, including follicular lymphomas. Thus, CD80 is an attractive target for lymphoma therapy. Targeting CD80 on NHLs with anti-CD80 antibody can arrest their growth and therefore can serve as a therapeutic intervention for treating cancer.

A significant number of patients relapse or do not respond to rituximab due to intrinsic or acquired resistance. Hence, mAbs targeting other cell

surface antigens on B-cell lymphomas are being studied. CD80 serves as an attractive target in the continued development of mAbs against lymphoma. Preclinical studies with galiximab, an anti-CD80 primatized mAb, have been encouraging and have demonstrated antitumor activity against various B-cell lymphoma models, both as a single agent as well as in combination with rituximab. Data were reviewed from a PubMed literature search from 1975 to 2009 and also included a review of abstracts from published proceedings of annual meetings from the American Society of Hematology and International Conference of Malignant Lymphoma, Lugano.

Galiximab (Biogen Idec, Inc., San Diego, CA) is a chimeric, primatized IgG1 1 mAb consisting of human constant and primate (*cynomolgus macaque*) variable regions targeting CD80. It binds specifically to CD80, which is transiently expressed on the surface of activated B cells and APCs, including DCs, but is constitutively expressed on a variety of NHLs, including follicular lymphoma (FL).¹¹⁵ Thus, CD80 serves as an attractive target for the development of antilymphoma therapy. Galiximab is emerging as an interesting and effective therapy for the treatment of patients with NHL, especially in combination with rituximab. Its favorable toxicity profile makes it an attractive alternative to existing agents. This “primatized” antibody is structurally indistinguishable from human antibodies and, therefore, is not significantly immunogenic in humans. Although a number of *in vivo* as well as *in vitro* studies point towards possible “unique” mechanisms of action of galiximab, more studies are needed to fully understand this agent’s antitumor activity, as well as to establish strategies by which to optimize its efficacy. However to further assess the therapeutic role of galiximab in the treatment of patients with follicular and other lymphoma subtypes, as monotherapy or in combination with rituximab or systemic chemotherapy, additional studies designed are under active consideration.^{116,117}

FULLY HUMANIZED ANTI-CTLA-4 ANTIBODY IN TREATMENT OF TUMORS

Ipilimumab (Fully Humanized Anti-CTLA-4 Antibody) in the Treatment of Tumors

The encouraging results obtained in preclinical models led to the development of two fully humanized anti-CTLA-4 antibodies, MDX-010 (ipilimumab), and CP-675,206 (tremelimumab) (Table 5.3). Ipilimumab is an IgG1 with a plasma half-life of 12–14 days, and it was approved by the FDA in March 2011 for the treatment of advanced melanoma (Bristol-Myers Squibb, Princeton, NJ). Both agents are able to recognize human CTLA-4 and block the interaction of CTLA-4 with CD80 or CD86, but their exact mechanism

of action is not fully understood. Preliminary data about an increase in antigen-specific effector T cells following ipilimumab treatment in combination with vaccine in three melanoma patients are also published.¹¹⁸ Ipilimumab and tremelimumab were tested in ovarian, breast, prostate, colon carcinoma, and, mainly, in melanoma and renal cell cancer clinical trials.¹¹⁹ Several early phase II studies evaluated ipilimumab in metastatic melanoma, reported an objective response rate ranging from 6% to 21% and a disease-control rate of about 30%.^{34,35,120} Recently, a multicenter single-arm phase II study evaluated the efficacy and the safety of ipilimumab monotherapy in patients with pretreated advanced melanoma. Patients ($n = 155$) were treated with ipilimumab at 10 mg/kg and the authors showed that 1- and 2-year survival rates were 47.2% and 32.8%, respectively, with a median overall survival of 10.2 months.¹²¹ Moreover, a randomized, double blind, placebo-controlled, phase II trial of ipilimumab at 10 mg/kg, with or without budesonide, in 115 patients with unresectable stage III or IV melanoma, showed that 2-year survival rate was approximately 40% in each arm.¹²² In another randomized, double blind, multicenter, phase II, dose-ranging study, 217 melanoma patients were randomly assigned to receive ipilimumab at 10 mg/kg ($n = 73$), 3 mg/kg ($n = 72$), or 0.3 mg/kg ($n = 72$). The authors observed a dose-dependent antitumor effect of ipilimumab, with the best overall response of 11.1% in patients treated with 10 mg/kg.¹²⁰ Ipilimumab was also tested in combination with other therapeutic agents such as IL-2, vaccine or chemotherapy, such as dacarbazine.^{123–125} In particular, interesting results came from the phase III trial by Hodi et al.; in this study, melanoma patients were randomized to receive ipilimumab 3 mg/kg, with or without a gp100 peptide vaccine, or the vaccine alone. The primary endpoint of the study was the overall survival. The median overall survival was approximately 10.0 months among patients receiving ipilimumab (with or without gp100 vaccine), as compared with 6.4 months among patients receiving gp100 alone.¹²⁶ In another phase III study, the administration of ipilimumab (at a dose of 10 mg/kg) in combination with dacarbazine was associated with a significant improvement in overall survival among patients with previously untreated metastatic melanoma.¹²⁵ Ipilimumab was also tested in other types of malignancies. In a phase II study on 61 patients affected by metastatic renal cell carcinoma, ipilimumab was administered at a dose of 1 mg/kg or 3 mg/kg; five of 40 patients treated with 3 mg/kg had a partial response.¹²⁷ A recent phase II study compared the addition of ipilimumab to carboplatin and paclitaxel chemotherapy in nonsmall cell lung cancer patients. Ipilimumab was administered either concurrently or in a phased schedule after receiving the first two cycles of chemotherapy. Patients treated with ipilimumab plus chemotherapy, in concurrent and sequential regimens, showed an improved overall survival compared with patients receiving chemotherapy alone.¹²⁸

Tremelimumab (Fully Humanized Anti-CTLA-4 Antibody) in the Treatment of Tumors

Tremelimumab (CP-675,206) is fully humanized anti-CTLA-4 an IgG2 antibody with a plasma half-life of approximately 22 days (Pfizer, New York, NY). Tremelimumab was first evaluated in the treatment of metastatic melanoma. A phase I clinical trials evaluating the maximum-tolerated dose of tremelimumab showed antitumor activity of this drug in melanoma patients.³⁶ Other phase I/II studies reported stable disease (SD) occurring in about 30% of melanoma patients treated with tremelimumab and an objective antitumor response in 10% of patients.^{37,129} A phase II study evaluated the antitumor activity of 15 mg/kg tremelimumab in 246 patients affected by melanoma, and reported an objective response rate of 6.6%, with duration of response ranged from 8.9 to 29.8 months.¹³⁰ Tremelimumab was also tested in cancers others than melanoma, both as monotherapy and in combination therapy, such as metastatic renal cell carcinoma, metastatic colorectal cancer, and advanced gastric and esophageal adenocarcinoma, but not significant clinical improvements were reported.^{38,39,131} The treatment with both ipilimumab and tremelimumab is associated with inflammatory AE like rash, diarrhea, colitis, and hypophysitis; this side effect profile could be linked to the potentiation of Th17 cell differentiation following CTLA-4 blockade.¹³²

ICOS-L:ICOS Costimulatory Pathway

The inducible costimulator (ICOS, CD278) is a costimulatory receptor that is weakly expressed on naive T cells and quickly upregulated in activated CD4⁺ and CD8⁺ T cells. A constitutive expression of ICOS by CD25⁺ CD4⁺ Foxp3⁺ Tregs has also been reported.¹³³ The cognate ligand for ICOS is ICOS-L (B7h, B7RP-1, CD275), which is expressed by professional APCs and by peripheral epithelial and endothelial cells following TNF- α stimulation.¹³⁴ The ICOS:ICOS-L pathway provides a key costimulatory signal for T-cell proliferation and, mainly, for T-cell survival. Moreover, ICOS regulates development and response of T follicular helper (Tfh), Th1, Th2, and Th17 cells and plays roles in the maintenance of memory effector T cells and Tregs homeostasis.¹³⁵ Due to its role in sustaining T-cell activation and effector functions, targeting ICOS:ICOS-L could represent a plausible approach to enhance antitumor immunity. The ICOS-L costimulation, through its expression on tumor cells, was capable of inducing cancer regression in Sa1/N fibrosarcoma and J558 plasmacytoma models.^{136,137} Systemic treatment with murine ICOS-L-IgG fusion protein was effective in promoting INF- γ -dependent antitumor immunity in MethA fibrosarcoma and B16F1 melanoma tumor models.¹³⁸ Recent data showed an increase of ICOS^{hi} CD4⁺ effector T cells percentage after CTLA-4 blockade in several cancer models. In addition, upon CTLA-4 blockade, this cell population produced greater levels of INF- γ than ICOS^{lo} CD4⁺ T cells, suggesting that ICOS could be used as a marker for CD4⁺ effector T-cell response.^{16,139,140}

A downregulation of ICOS was shown in colon cancer patients and the expression of ICOS in tumors was associated with a greater survival of melanoma patients.^{141,142} A recent study investigated the role of ICOS in the Tregs in melanoma and demonstrated the selective expression of ICOS on a “hyperactivated” Treg population, which strongly inhibits T-cell response through IL-10-mediated APCs suppression.¹⁴³ Moreover, ICOS-L was expressed by both cultured and freshly isolated melanoma cells from stage IV melanoma patients and could provide costimulation through ICOS for the activation and expansion of Tregs in the tumor microenvironment, as another mechanism of escape from immune surveillance.¹⁴⁴ Thus, targeting costimulatory and coinhibitory molecules on Tregs might be a promising approach for modulating peripheral tolerance in cancer patients.

PD-L1/PD-L2:PD-1 Costimulatory Pathway

Programmed cell death 1 (PD-1) is a negative costimulatory receptor belonging to the B7/CD28 family. PD-1 expression is induced on activated T cells, B cells, monocytes, DCs, and, at low levels, on natural killer T cells (NKT). Negative costimulatory signals mediated via PD-1 and CTLA-4 are not redundant; in fact PD-1 mainly acts in regulating inflammatory responses in peripheral tissues, whereas CTLA-4 modulates T-cell priming in lymphoid organs. In addition, in contrast to CTLA-4, PD-1 is able to block TCR- and CD28-mediated activation through the recruitment of inhibitory phosphatases, such as SHP-2, which inhibits the induction of PI3K activity.¹⁴⁵ PD-1 has two known ligands belonging to the B7 family: PD-L1 (B7-H1) and PD-L2 (B7-DC). To date, one of the major differences between these ligands concerns their expression pattern. PD-L1 mRNA is broadly expressed in multiple peripheral tissues such as heart, placenta, muscle, fetal liver, spleen, lymph nodes, and thymus; PD-L1 protein has been found in activated T cells, B cells, monocytes, DCs, endothelial cells, and myocardium and can be upregulated following exposure to type I or type II interferon, providing a negative feedback mechanism to dampen immune response.¹⁴⁵ On the contrary, PD-L2 expression is largely restricted to activated macrophages and DCs, but only PD-L2 mRNA can also be observed in the human heart, placenta, lung, and liver.¹⁷

CD279 (PD-1), CD274 (PD-L1), and CD273 (PD-L2) Expression in Malignancy

Although initial *in vitro* work showed PD-L1/PD-L2.Ig transmitted stimulatory signals to activated lymphocytes binding to their ligand PD-1 conveyed potent suppressive signals.¹⁷ Evidence for bidirectional signaling emerged from murine B-cell- and T-cell-specific gene deletion and transplantation experiments. Knock-out of T-cell-specific PD-1 or B-cell-specific PD-L2 impaired long-lived plasma cell formation, which implied a stimulatory role

for reverse signaling. Receptor/ligand pairing is also more promiscuous than originally believed. A search for binding partners for CD80 using *Ctla-4*^{-d}*Cd28*^{-d2} activated CD4⁺ T cells suggested PD-1 as a candidate, with CD80 signaling through PD-1 independent of CTLA-4 or CD28, to inhibit T-cell proliferation.¹⁴⁶ The complexity of the B7 system is becoming clear: context and direction of signaling determines responder cell outcome.

CD279 is invariably expressed in angioimmunoblastic T-cell lymphoma, frequently in CLL, but absent in most other NHL. CD279 is expressed in the immune microenvironment of NHL, where it may have a prognostic significance.¹⁴⁷ Evidence for expression of CD274 in NHL is sparse and inconsistent. Immunohistochemistry and flow cytometric studies have shown expression by the malignant cell in the majority of primary mediastinal large B-cell lymphoma and anaplastic large cell lymphoma (ALCL), a minority of diffuse large B-cell lymphoma, and not at all in most other NHL. CD274 is overexpressed in the Hodgkin and Reed-Sternberg (HRS) cells of classic Hodgkin lymphoma (CHL) associated with copy number aberrations and activating translocations involving the 9p24 chromosomal region containing the CD273 and CD274 genes,¹⁴⁸ and upstream through the CD274 and CD273 expression-regulating JAK2 pathway.¹⁴⁹ Limited *in vitro* evidence suggests that there is CD274-dependent suppression of T cells in CHL, evidenced by suppressed proliferation and IFN- γ production.

There is variable expression of CD274 in myeloma with evidence for an important role in myeloma-induced immunosuppression. CTL generated *in vitro* by exposure to IFN- γ -treated myeloma cell lines show impaired cytotoxicity of subsequently presented target myeloma cells NK-cell trafficking, and cytotoxic potency *in vitro* is augmented by pretreatment with anti-PD-1 mAb. An autologous transplantation and costimulator/vaccine-based treatment strategy for myeloma in a cell line-based mouse model, which had showed evidence for the emergence of dysfunctional circulating CD279-expressing CD8⁺ T cells, is augmented by the administration of PD-L1 antibodies.^{150–152} While somewhat conflicting and lacking functional corroboration, these findings suggest that therapeutic modulation of this axis in hematologic malignancy may have a role.

Therapeutic Modulation of PD-1 (CD274), PDL-1 (CD279), and PDL-2 (CD273)

The PD-1/PD-L1/PD-L2 axis has been widely demonstrated to contribute to failed antitumor immunity. *Pd-1*^{-d} mice and *Pd-1*^{wt/wt} mice treated with anti-PD-L1 antibody demonstrate superior rejection of PD-L1-expressing tumors through PD-1-dependent and -independent pathways.^{14,18} Growth of pancreatic cancer cells in immunocompetent B6 mice is impaired by administration of anti-PD-1 or PD-L1 mAbs.¹⁵ Tumor-specific T cells fail to lyse melanoma cells in host *2C/Rag2*^{-ag} mice despite stimulatory IFN- γ

treatment because IFN- γ also induces upregulation of immunosuppressive PD-L1.¹⁵³ PD-L1 is expressed in many tumors and cell lines and appears to be associated with poorer outcome in solid malignancy, while PD-1 expression by tumor-infiltrating lymphocytes (TILs) may also confer poorer outcomes. After treatment using a tumor vaccination strategy of murine acute leukemia, long-lived residual leukemia cells were found to upregulate PD-L1, which suggests a mechanism of leukemia persistence and relapse.¹⁵⁴ These cells resisted CTL-mediated cell lysis, which could be overcome by anti-PD-L1 mAbs. However, blockade using anti-PD-1 mAbs did not reverse CTL resistance, while blocking CD80 had a similar effect to a PD-L1 block. The B7 system's complexity is further illustrated by these findings: CD274 appears to have further immunosuppressive binding partners, while CD80 may be capable of mediating suppressive as well as costimulatory signals.¹⁵⁵

There is sparse and conflicting evidence for CD274 (PD-L2) expression in acute leukemia,¹⁵⁶ although the molecule is apparently upregulated on malignant cell exposure to IFN- γ or Toll-like receptor (TLR) ligand, and at relapse. Evidence for reduced T-cell-mediated specific lysis *in vitro* correlates with CD274 expression and is reversible by blocking CD274.¹⁵⁷ Functional work in CLL has demonstrated a clear role for the PD-1/PD-L1 axis in mediating systemic, as well as antitumor, immune defects. Expression of CD279 by circulating T cells and CD274 by leukemia and FL cells is greater than in nonmalignant control cells, while CD274 expression is associated with poorer clinical outcome. T cells derived from patients with various malignancies have impaired *in vitro* ρ -GTPase signaling with reduced immune synapse formation. This defect is reversed by treatment with anti-PD-1 and anti-PD-L1 mAbs, along with blockade of further inhibitory molecules CD200, CD270, and CD276 (B7-H3). Lenalidomide exposure also reverses this defect, coincident with downregulation of CD274 and CD276 expression.¹⁵⁸ These findings, along with evidence that CD80 expression is upregulated on lenalidomide-exposed CLL cells,¹⁵⁹ provide some insight into the drug's mechanism of action. In patients with adult T-cell leukemia/lymphoma (ATLL), CD279 expression on CD8 + T cells is upregulated compared with ATLL-free human T-cell lymphotropic virus (HTLV1+) controls.¹⁶⁰

PD-L1 aberrant expression has been reported in many human cancers, such as glioblastoma, melanoma, and cancers of the head and neck, lung, ovary, colon, stomach, kidney, and breast.^{161,162} Moreover, in several follow-up studies, the expression of PD-L1 correlates with a poor prognosis of patients.^{15,163–165} Based on these experimental evidences, PD-L1 blockade has been proposed for cancer immunotherapy. Two independent studies have shown that forced expression of PD-L1 in the murine myeloma cell line P815 render them more resistant to *in vitro* cytolysis and less susceptible to rejection than control when inoculated in mice.¹⁸ In addition, the treatment with anti-PD-L1 mAbs was capable of inhibiting the growth of P815-PD-L1 *in vivo*.¹⁴ PD-1 blockade was able to restore the antitumor

immunity to accelerate tumor eradication in murine squamous cancer cell line SCCVII and in P815 cell line and to block both CT26 colon carcinoma metastasis to the lung and B16 melanoma metastasis to the liver. *In vivo* studies in these tumor models examined the combination of anti-PD-1 mAb with GM-CSF-secreting tumor cell vaccine and reported that the administration of anti-PD-1 mAb enhanced the efficacy of vaccine by increasing number and activity of tumor-specific CD8 + T cells.¹⁶⁶

Recent a study reported that the combination of PD-1 blockade and NKT cell activation results in increased antitumor responses in a melanoma model.¹⁶⁷ A reduced number of Tregs at the tumor site was observed after the treatment with anti-PD-1 mAb and the TLR agonist CpG, which suggests that it could be another mechanism by which PD-1 blockade exerts antitumor effects.¹⁶⁸ The expression of PD-L1 on DCs is also able to block antitumor T-cell response; myeloid DCs expressing PD-L1 isolated from ovarian cancer poorly stimulated T cells and PD-L1 blockade could revert this scenario.¹⁶⁹ In addition, the anti-PD-L1 therapy was observed to revitalize “exhausted” antiviral CD8 + T cells in animals with chronic viral infections.¹⁷⁰ The promising results in preclinical models have led to the development of two humanized antibodies against PD-1 receptor that block its interaction with PD-L1:CT-011 and MDX-1106 (Table 5.4). A preclinical study evaluated the combination of CT-011 with low dose of cyclophosphamide and with a tumor vaccine; the authors reported the complete regression of established tumors in most of the animals treated.¹⁷¹ Benson et al. reported that CT-011 enhances NK-cell migration toward malignant plasma cells in MM.¹⁵² To date, only phase I clinical trials have been conducted to evaluate both efficacy and safety of these two agents. In a phase I trial enrolling 17 patients with advanced hematological malignancies, the treatment with CT-011 at doses ranging from 0.2 to 6 mg/kg led to clinical improvement in 33% of patients.⁴⁰ Recently, the administration of CT-011 in combination with autologous DC/myeloma fusion-vaccine was demonstrated to stimulate T-cell responses after vaccine administration.²⁵ The efficacy of MX-1106 was evaluated in a phase I study enrolling 39 patients with advanced solid cancers (melanoma, colorectal cancer, castrate-resistant prostate cancer, nonsmall cell lung cancer, and renal cell carcinoma) obtaining very promising results; phase II and III trials are under evaluation.⁴¹ In particular, it will be interesting to evaluate the combination of anti-PD-1 with other agents, such as anti-CTLA-4 mAb, vaccines, and chemotherapy. Several phase II clinical trials are testing the safety and the efficacy of these two PD-1 antibodies in several types of cancer.¹⁷² MK-3475 and MDX-1105-01 are other two antibodies against PD-1 and PD-L1, respectively, which are currently being investigated in phase I clinical trials⁴² (also see Table 5.4).

Fully humanized IgG4 mAbs targeting the PD-1/PD-L1 axis have reached clinical trials. Anti-PD-L1 therapy (BMS936559) (see Table 5.4) was administered to 160 heavily pretreated patients with a diverse group of solid

tumors in a phase 1 trial¹⁷³ Objective and often durable responses were observed in up to 19% of patients, along with autoimmune and infusion reactions consistent with the murine PD-L1^{-/-} models. A similar clinical trial of an anti-PD-1 monoclonal antibody therapy (BMS936558) demonstrated comparable response rates in 296 patients.¹⁷⁴ There was an association between response and level of tumor PD-L1 expression. Further combination therapy approaches using anti-PD-1 mAbs are under way, which include combination with the anti-CTLA-4 mAb ipilimumab in melanoma (NCT01024231), with conventional combination chemotherapy in nonsmall cell lung cancer (NCT01454102) and with tyrosine kinase inhibitors sunitinib and pazopanib in renal cell carcinoma (NCT01592370). Another PD-1 antibody (MK-3475) has also shown activity in patients with advanced solid tumors.¹⁷⁵ A phase 1 safety and pharmacokinetic study of the anti-PD-1 mAb in advanced diverse hematologic malignancies showed a reasonable safety profile and a 33% overall response rate (ORR)⁴⁰ with further monotherapy and combination therapy trials proposed. The results are promising, but these drugs remain at an early stage of development and are likely to find their most active role as immune adjuvants to conventional therapy.

HVEM:BTLA/CD160 Costimulatory Pathway

The BTLA is another member of the B7/CD28 family acting as a negative costimulatory receptor. The constitutive expression of BTLA has been reported, at low levels, on naive B and T cells, Tfh, macrophages, DCs, NKT cells, and NK cells, but unlike CTLA-4 and PD-1, BTLA is not expressed on Tregs.¹⁷⁶ The HVEM, a member of TNFR superfamily, has been identified as a BTLA ligand. HVEM expression is high in naive T and B cells, but it decreases during T-cell activation. HVEM is also expressed on DCs, Tregs, monocytes, NK cells, and neutrophils, and in non-hematopoietic cells, such as parenchymal cells. In addition to BTLA, HVEM binds also CD160, another member of the B7/CD28 family. CD160 is highly expressed on CD56dimCD16 NK cells, NKT cells, $\gamma\delta$ T cells, CD8+ CD28- T cells, a small subset of CD4+ cells and intestinal intraepithelial T cells.¹⁷⁶ HVEM also interacts with LIGHT (described below in the text) and lymphotoxin alpha (LT α) of TNFR family, as and such is a unique example of a direct interaction between the two families. Therefore, HVEM is considered a molecular switch because of its ability to regulate the immune response, which depends on which cognate ligand binds. The role of BTLA as a negative costimulatory receptor has been shown by the phenotype of BTLA deficient mice, which were more susceptible to develop autoimmune disorders, and by the *in vitro* observation that anti-BTLA agonists drive negative signals to T cells.¹⁷⁷ The engagement of BTLA results in the inhibition of CD3/CD28 T-cell activation. BTLA signals through the recruitment of SHP-2 phosphatase, but the downstream target of SHP-2 is unclear. Anyway,

recent studies showed that triggering BTLA signaling in B cells resulted in blocking B-cell proliferation through the inhibition of phosphorylation of some transcription factors like NF- κ B.¹⁷⁸

The similarity between BTLA and PD-1 signaling could justify a possible use of BTLA blockade to enhance antitumor immunity. The expression of BTLA was found in CLL/small lymphocytic lymphoma.¹⁷⁹ Moreover, soluble BTLA seems to enhance antitumor efficacy of the HSP70 vaccine in murine TC-1 cervical cancer mice.¹⁸⁰ Recently, a study by Derré et al. demonstrated the potentiality of targeting BTLA for cancer immunotherapy, and reported that BTLA is expressed on tumor antigen-specific CD8 + T cells from melanoma patients and that this molecule inhibits their fully functionality; following the vaccination with CpG adjuvants, the authors observed a downregulation of BTLA, with a partial recovery of CD8 + T cells functionality.¹⁸¹ BTLA-HVEM blockade showed antitumor effects in murine TC-1 cervical cancer model *in vivo*, resulting in downregulation of IL-10 and TGF-beta and in activation of DCs in IL-12- and B7-1-dependent manner.

In addition, BTLA-HVEM blockade alone was not effective in eradicating the tumor, whereas the combination with HSP70 vaccine improved antitumor immunity by increasing IL-2 and INF- γ production and decreasing IL-10, TGF-beta, and Foxp3 transcription levels in the tumor microenvironment.¹⁸⁰ The evidences supporting a negative costimulatory function for CD160 come from *in vitro* studies because CD160 deficient mice have not been generated. CD160 agonists strongly inhibited CD4+ T-cell proliferation and cytokines production and reduced INF- γ secretion by NK cell line.¹⁸² Recently, Cai et al. reported a strong inhibition of CD3/CD28-induced T-cell activation after the use of CD160 agonists, but the downstream intracellular pathways involved are not known.¹⁸³ Indeed, Liu et al. reported that CD160 is expressed in B-cell CLL, in which its engagement mediates survival and growth signals. In fact, CD160 activation was associated with upregulation of antiapoptotic genes Bcl-2, Bcl-xL, and Mcl-1 and, consequently, with reduced mitochondrial membrane potential collapse and cytochrome c release. CD160 engagement also induced cell cycle progression and proliferation.¹⁸⁴ A recent study examined the expression of CD160 in 811 cases of B-cell lymphoproliferative disorders (B-LPD). The authors showed that CD160 was expressed in 98% of CLL cases, 100% of hairy cell leukemia cases, 15% of mantle cell lymphoma in the leukemic phase, and 16% of other B-LPD cases, whereas it was absent in the normal B-cell lineage.¹⁸⁵ Recently, Chabot et al. suggested a role for CD160 in tumor neoangiogenesis. CD160 was expressed on newly formed blood vessels in human colon carcinoma and mouse B16 melanoma, but not in the healthy vessels. Treatment with anti-CD160 monoclonal antibody CL1-R2 in combination with cyclophosphamide chemotherapy resulted in the regression of tumor vessels in B16 melanoma-bearing

mice.¹⁸⁶ Further studies are needed to clarify this pathway so as to design potential CD160/BTLA-based antitumor therapeutic strategies.

B7-H3 and B7-H4 Costimulatory Pathway

B7-H3 and B7-H4 (B7x, B7S1) are two of the newer members of the B7 family. B7-H3 expression has been found to be inducible on T cells, NK cells, and APCs. B7-H3 is also broadly expressed on osteoblasts, fibroblasts, and epithelial cells, as well as in liver, lung, bladder, testis, prostate, breast, placenta, and lymphoid organs. To date, only one potential receptor of B7-H3 on activated T cells named TLT-2 has been identified.¹⁸⁷ There are conflicting data about the functions mediated by B7-H3, as both stimulatory and inhibitory properties have been reported. Because of these controversial results, the possible existence of additional receptors for B7-H3 has been taken into consideration.

Recently, several works reported B7-H3 expression in different human cancers, as reviewed by Loos et al. The double stimulatory and inhibitory nature of B7-H3 signaling also appeared in murine cancer models and in human cancers. The authors showed that B7-H3 expression is associated both with favorable and adverse clinic-pathologic features.¹⁸⁷ Due to its immunomodulatory ability, B7-H3 blockade could be a potential anticancer immunotherapy, but the controversial findings about its role remain to be elucidated.

B7-H4 mRNA is broadly expressed in the peripheral tissues, whereas protein expression is restricted to activated B cells, T cells, and monocytes. Several studies reported the expression of B7-H4 in many human cancers, such as nonsmall cell lung cancer, ovarian cancer, prostate cancer, breast cancer, and renal cancer. To date, the cognate receptor of B7-H4 on activated T cells remains unclear, although BTLA has been reported as a possible receptor. Unlike B7-H3, which shows opposite functions, B7-H4 mediates a negative costimulatory signal. B7-H4 strongly inhibited T-cell proliferation and IL-2 secretion, and its blockade with antagonistic mAbs resulted in *in vitro* enhanced T-cell response and *in vivo* exacerbation of EAE.^{188,189}

Recent studies indicate that B7-H4 could be a potential diagnostic/prognostic marker and/or therapeutic target for several cancers. In fact, B7-H4 expression was found to correlate with stage, pathological types, and biological behavior of many tumors, as shown by retrospective analyses on 13 types of human cancers. Moreover, the expression of B7-H4 reverse correlated with the survival of patients in most cancer analyzed.¹⁹⁰ A recent study by Quandt et al. demonstrated that the overexpression of B7-H4 in melanoma cells impaired the antitumor immune response by decreasing IFN- γ , TNF- α , and IL-2 production.¹⁹¹

In ovarian cancer, the inhibitory role of B7-H4 might be due to B7-H4 expression on tumor-associated macrophages (TAMs). Kryczek et al. showed that B7-H4-expressing TAMs could block activation of T cells in the setting

of ovarian cancer. B7-H4 expression in TAMs was upregulated by IL-6 and IL-10 present in the tumor microenvironment, whereas its expression was inhibited by GM-CSF and IL-4. It has been shown that Treg cells induced APCs to produce IL-10 and IL-6, providing a new mechanism by which Tregs exert their suppressive action. B7-H4 blockade by antisense oligonucleotides restored T-cell antitumor immune response and led to tumor regression *in vivo*.¹⁹² Recently, Qian et al. reported the development of a monoclonal antibody to B7-H4 and preliminary data showed that it could effectively inhibit the activity of B7-H4, promoting the growth of T cells and the secretion of IL-2, IL-4, IL-10, and IFN- γ .¹⁹³ Based on these preliminary data, the blockade of B7-H4 could be an attractive opportunity to enhance antitumor immunity in a subset of human cancers.

TNF:TNFR SUPERFAMILY OF COSTIMULATORY MOLECULES IN ANTITUMOR IMMUNITY

CD40L:CD40 Costimulatory Pathway

CD40 receptor and its ligand CD40L (CD154) were the first members belonging to the TNF:TNFR family to be identified. CD40 expression was originally found on B cells, but it is also expressed on DCs, monocytes, platelets, macrophages as well as myofibroblasts, fibroblasts, epithelial, and endothelial cells.¹⁹⁴ CD40L is expressed on activated T and B cells, by platelets, monocytes, NK cells, mast cells, and basophils, where CD40L is induced following proinflammatory stimuli. CD40L:CD40 signaling in B cells is also important for the generation of long-lived plasma cells and memory B cells, as well as for their survival. CD40 intracellular signaling is mediated by the recruitment of TNFR-associated factors (TRAFs), which in turn activate different pathways, such as the canonical and noncanonical NF- κ B pathway, MAPKs, PI3K, and the phospholipase C γ pathway.¹⁹⁵ The engagement of CD40 by CD40L on APCs has been shown to promote cytokines production and upregulation of costimulatory molecules, crucial events for T-cell activation, and differentiation.¹⁹⁶

CD40/CD40L pathway is crucial for the development of antitumor immunity. CD40L blockade resulted in lacking of protective immune response following administration of a GM-CSF-expressing B16 melanoma cells vaccine. In addition, low expression of CD40L was sufficient to induce a long-lasting antitumor immune response via CTLs in a small number of cancers.¹⁹⁵ Another aspect to take into consideration is the direct effect of CD40:CD40L pathway on tumor cells. Elgueta et al. reviewed that CD40 is broadly expressed in several tumors, such as melanoma, prostate cancer, lung cancer, as well as carcinoma of nasopharynx, bladder, cervix and ovary, Hodgkin's and NHL, MM, and acute myeloid leukemia. The engagement of CD40 on tumor cells can provide growth arrest and apoptosis of malignant

cells, as dictated by the type of malignancies and the microenvironment.¹⁹⁵ The combination of CD40L expression with other immunomodulators (IL-2, GM-CSF, and INF- γ) has been found to promote antitumor immunity in several cancer models. Gene therapy approach with the use of recombinant adenovirus encoding CD40L was also effective in colorectal, lung, and melanoma cancer models. An early study reported that the use of CD40 agonistic antibodies triggered CTL-4 responses in a lymphoma system, with the consequent tumor eradication.¹⁹⁷ Recently, the use of activators of both adaptive and innate immunity, such as CD40 agonists and TLR agonists, induced antitumor-specific immunity in many tumor models.¹⁹⁸ Currently, the combination of CD40 tumor therapy with other approaches, such as cancer vaccines, chemotherapy, radiation, CTLA blockade, TLR agonists, and cytokines, is becoming overriding.^{199,200}

To date, three humanized CD40 agonistic antibodies have been developed: CP-870,893, SGN-40, and HCD122 (Table 5.8). CP-870,893 is a fully human, IgG2 antibody that selectively binds to CD40. It enhances the expression of MHC class II, CD54, CD86, and CD23 on human B cells *in vitro*. CP-870,893 also enhances DC activity as demonstrated by secretion of IL-12, IL-23, and IL-8, by the upregulation of CD86 and CD83, and by the capacity to prime T cells to secrete IFN- γ .^{211,212} Results from a phase I

TABLE 5.8 Tumor Models and Clinical Trials of TNF-TNFR Costimulatory Molecules

Tumor Model	Costimulatory Molecule	Therapeutic Strategy	Reference
Melanoma, advanced cancers	CD40	CP-870,893 agonistic Ab anti-CD40	[201,202]
Myeloma, B-cell lymphoma, diffuse large cell lymphoma	CD40	SGN-40 agonistic Ab anti-CD40	[203–205]
MM, B-cell CLL	CD40	HCD122 agonistic Ab anti-CD40	[195]
Solid tumors	4-1BB	BMS-663513 agonistic Ab anti-4-1BB	[206]
Metastatic prostate cancer	OX40	Agonistic Ab anti-OX40	[42]
Melanoma	GITR	TRX518	[42]
CD30-positive lymphoma	CD30	Brentuximab vedotin Ab-drug conjugates anti-CD30	[207–210]

study showed that administration of CP-870,893 was associated with early signs of clinical efficacy, especially in patients with melanoma.²⁰¹ The same authors reported that weekly infusions of this agonist CD40 antibody were associated with little clinical activity in advanced cancer patients.²⁰² SGN-40 (dacetuzumab) is a humanized anti-CD40 monoclonal antibody with multiple mechanisms of action. In non-Hodgkin lymphoma, Dacetuzumab activates two distinct proapoptotic signaling pathways; on the one hand, it constitutively activates the NF- κ B and MAPK signaling pathways producing the sustained downregulation of the oncoprotein B-cell lymphoma 6, which loss results in c-Myc downregulation and activation of early B-cell maturation, concomitant with reduced proliferation and cell death. On the other hand, dacetuzumab induces the expression of the proapoptotic p53 family member TAp63 α and downstream proteins associated with the intrinsic and extrinsic apoptotic machinery.²¹³ *In vitro*, dacetuzumab exhibited antitumor activity against several B-cell lymphoma and MM cell lines, and induced direct apoptosis as well as the engagement of effective antibody-dependent cell-mediated cytotoxicity (ADCC).²¹⁴

Early clinical trials have evaluated the pharmacokinetics, safety, and efficacy of dacetuzumab in patients with relapsed/refractory B-cell lymphomas, MM, and CLL.^{203–205} Dacetuzumab resulted in modest antitumor activity in B-cell lymphomas and, to a lesser extent, in MM. In CLL dacetuzumab showed modest activity as monotherapy, while better results were obtained by using combination therapy with lenalidomide.²¹⁵ Dacetuzumab is currently in multiple phase II trials for the treatment of myeloma and diffuse large cell lymphoma. HCD122 is a human IgG1 monoclonal antibody and currently in a phase I trial for the treatment of MM and B-cell CLL. In B-cell CLL, HCD122 exerts antitumor activity by killing leukemia cells through ADCC and inhibiting CD40L induced survival and proliferation of tumor cells.²¹⁶

1BBL:4-1BB Costimulatory Pathway

4-1BB is an inducible costimulatory receptor expressed on activated CD4 + and CD8 + T cell, NKT, NK cells, DCs, macrophages, eosinophils, neutrophils, and mast cells, as well as Tregs. In the most cases, 4-1BB is induced on the cellular surface following activation, except for APCs and Tregs, where its expression is constitutive. The ligand of 4-1BB is 4-1BBL, which is expressed on activated professional APCs.^{217,218} 4-1BB:4-1BBL pathway seems to amplify the existing costimulatory signals, even if the engagement of 4-1BB in the presence of a strong TCR signaling can induce IL-2 production in a CD28-independent manner.²¹⁹ Following stimulation with its ligand, 4-1BB provides costimulatory signals to both CD4 + and CD8 + T cells, with a greater effect on the expansion of CD8 + due to the upregulation of antiapoptotic genes, such as *bcl-XL* and *bfl-1*. 4-1BB signals are mediated by the activation of NF- κ B, c-Jun, and p38

downstream pathways.¹⁵⁵ 4-1BB has also a role in activation of DCs, which induces IL-6 and IL-12 production and upregulating B7 costimulatory molecules.²²⁰ 4-1BB plays roles in activating non-T-cells other than DCs, such as monocytes, B cells, mast cells, NK cells, and neutrophils and its engagement is associated with cellular proliferation, cytokine induction, bactericidal activity, and sustenance of T-cell effector functions.

Targeting of 4-1BB:4-1BBL pathway in cancer reveals itself as a promising approach. The adoptive transfer of *ex vivo* 4-1BB- and CD28-costimulated T cells induced antitumor immune response in some preclinical studies.^{221,222} 4-1BB agonistic antibodies as antitumor therapy were broadly tested in several animal models with encouraging results. Melero et al. reported that the intraperitoneal injection of an antimurine 4-1BB mAb resulted in the eradication of established P815 mastocytoma and Ag104A sarcoma in mice.²²³ Driessens et al. reviewed of subsequent studies that demonstrated the efficacy of anti-4-1BB- or 4-1BBL-expressing tumor cells vaccines in inducing specific antitumor T-cell response, suppression of tumor growth and regression of pre-established tumors in different animal models. Therapeutic effects of agonistic anti-4-1BB mAb are due to enhanced NK and CD8+ T-cell activation and IFN- γ production.²¹⁸ The current direction toward which 4-1BB-directed anticancer immunotherapy is moving is the use of anti-4-1BB mAbs in combination with other therapeutic approaches, such as antitumor necrosis factor-related apoptosis inducing ligand (TRAIL), CD40 mAbs, intratumoral delivery of IL-12 gene, DC vaccines, CTLA-4 blockade, anti-CD4 therapy, chemotherapy, and radiotherapy.²²⁴ To date, one agonistic anti-4-1BB-humanized mAb BMS-663513 has been developed and the functional effects were demonstrated on human and monkey T cells and peripheral blood mononuclear cells, where IFN-production was enhanced compared to controls (Table 5.8). BMS-663513 is under evaluation in several phase I and II trials in patients with solid tumors and showed clinical activity. A phase II randomized study in melanoma patients with stage IV disease was stopped due to the occurrence of hepatitis.^{206,225}

OX40:OX40L Costimulatory Pathway

OX40 is an inducible costimulatory receptor expressed on activated CD4+ and CD8+ T cells, but also on activated Tregs, NKT cells, NK cells, and neutrophils. OX40L expression is induced on professional APCs, as well as on T cells, with the aim of amplifying T-cell responsiveness during T-cell/T-cell interactions. In addition to APCs, other cell types can induce OX40L expression, such as Langerhans cells, mast cells, NK cells, endothelial cells, and smooth muscle cells. Based on experimental evidences from OX40 deficient mice, it has been reported that this receptor promotes effector T-cell proliferation and survival, cytokines production, as well as the generation, and the maintenance of memory T cells. Moreover, OX40 inhibits Treg functions and counteracts the

generation of inducible Tregs.²²⁶ OX40 seems to act as a late positive costimulatory receptor, which goes on after CD28 signal, in a sequential manner.²²⁷ The prosurvival activity of OX40 is in part due to its ability to upregulate antiapoptotic genes of the Bcl-2 family. In fact, OX40 engagement by OX40L activates both PI3K/Akt and NF- κ B downstream pathways.²²⁸

OX40 represents a promising candidate for cancer immunotherapy. As reviewed by Croft et al., different approaches have been evaluated, such as agonistic OX40 mAbs or OX40L-Ig fusion protein, tumor cells, and DCs transfection with OX40L, and agonist RNA aptamer-binding OX40. Treatment with agonistic OX40 mAbs or OX40L-Ig fusion protein resulted in enhanced antitumor immunity in several cancer models, such as sarcoma, melanoma, colon carcinoma, and glioma.²²⁶ Several studies also reported encouraging results following the use of agonistic OX40 mAbs in combination with IL-12, anti-4-1BB, GM-CSF, DC vaccine, IL-12, and CD80 costimulation and chemotherapy.^{229–231} Combination therapy with agonistic OX40 mAbs and cyclophosphamide induces a profound Tregs depletion in concomitance with an increased infiltration of effector CD8⁺ T cells in B16 melanoma model.²³²

A phase I/II trial is ongoing to evaluate the safety and the efficacy of a murine anti-human OX40 in combination with cyclophosphamide and radiation in patients with progressive metastatic prostate cancer (see Table 5.8). To avoid immune response to the murine mAb, a humanized OX40 agonist has been developed by Agonox, a spinoff biotech company, and it will be tested in future clinical trials.⁴²

Light: HVEM Costimulatory Pathway

LIGHT, along with LT α , was identified as a ligand of the aforementioned HVEM receptor and it is a member of TNF family. LIGHT expression has been reported on activated T cells, immature DCs, monocytes, and NK cells. LIGHT is not expressed on B cells, but it can be induced following activation.²³³ Experimental evidences suggested that the interaction HVEM/LIGHT results in a positive costimulatory signaling, which induces T-cell proliferation and cytokines production.¹⁷⁷ In fact, the constitutive expression of LIGHT in T cells of transgenic mice leads to accumulation and activation of DCs and expansion of activated effector and memory T cells.²³⁴ Moreover, the manifestation of lymphoproliferative disorders and autoimmune disease was also observed.²³⁵ LIGHT deficient mice have been generated and showed defects in CD8⁺ T-cell activation and in thymic selection. LIGHT is also a critical ligand for activating NK cells to produce IFN- γ .²³³ The intracellular signaling of HVEM following LIGHT binding is mediated by TRAF proteins, which in turn activate NF- κ B and c-Jun/AP-1 pathways, leading to the transcription of prosurvival and proliferative genes, as well as genes regulating cytokines secretion. LIGHT can also engage the

lymphotoxin- β -receptor (LT β R) on DCs and provide a crucial signaling that results in DCs expansion, activation and IL-2 production.¹⁷⁷

Thus, LIGHT can modulate the immune response both directly by signaling via HVEM on T cells and indirectly by activating DCs through LT β R. Due to its role as immunomodulator, LIGHT could be a suitable target for cancer immunotherapy. The overexpression of LIGHT in P815 myeloma cell line induces regression of established tumors in a CD28-independent manner;²³⁶ similar results were obtained upon LIGHT overexpression in Ag104 sarcoma cell line; in fact, this forced expression caused rejection of tumor through NK-cell activation, which, in turn, triggered tumor-specific CD8+ T-cell proliferation at the tumor site.²³⁷ In addition, the injection of LIGHT expressing adenoviral vector into primary 4T1 mammary carcinoma has been found to promote T-cell recruitment, immune surveillance of the tumor, and elimination of metastasis.²³⁸ A recent study showed that the LIGHT/HVEM costimulation through both LIGHT-transfected cells and HVEM agonistic mAb-induced apoptosis in fresh B-CLL cells along with an increased production of IL-8.²³⁹ Another therapeutic approach targeting LIGHT/HVEM signaling was reported by Park et al., which developed P815 tumor cell expressing a single-chain variable fragment (scFv) of an anti-HVEM agonistic monoclonal antibody on their surface. These authors showed that tumor cells expressing anti-HVEM scFv spontaneously regress in a CD4+ and CD8+ T-cell-dependent manner when inoculated in mice and stimulated tumor-specific long-term T-cell memory. Moreover, the combination of anti-HVEM scFv expressing tumor vaccines and 4-1BB costimulation caused the regression of established tumors *in vivo*.²⁴⁰ Further studies are needed to clarify the true potential of targeting this pathway in cancer immunotherapy.

CD70:CD27 Costimulatory Pathway

CD27 is another costimulatory receptor belonging to the TNF family, and it is expressed on naive T and B cells and on NK cells. CD70 has been identified as CD27 ligand and its expression is restricted to APCs. The engagement of CD27 by CD70 promotes a positive costimulatory signaling, which results in T-cell proliferation and survival, maybe in concert with CD28.²²⁷ Like other members of the TNFR family, CD27 signaling is mediated by the recruitment of TRAF proteins.²⁴¹ Targeting CD27 could represent an attractive strategy in the field of cancer immunotherapy. Earlier studies have reported that early studies reporting that the overexpression of CD70 promoted cancer elimination through the activation of T cells and NK cells. In addition, the potential of costimulatory ligand CD70 to boost DC-based vaccine capacity to evoke effective CD8+ T-cell immunity has been explored.²⁴² Glouchkova et al. suggested that the modulation of the CD70/CD27 pathway might represent a novel therapeutic approach for enhancing the antileukemic response in B-cell precursor acute lymphoblastic leukemia.²⁴³

Recently, agonistic anti-CD27 antibodies has shown to be effective as monotherapy in reducing the outgrowth of experimental lung metastases and established subcutaneous melanoma tumors *in vivo*.²⁴⁴ In addition to CD70 agonists, the soluble form of CD70 has been evaluated as powerful adjuvant in a glioblastoma model.²⁴⁵ The aberrant expression of CD70 in a broad range of hematological malignancies and in some solid tumors has led to the development of CD70-specific T cells, having a chimeric antigen receptor (CAR) consisting of CD27 fused to the CD3-chain. Recently, adoptively transferred CD70-specific T cells have been found to induce regression of established murine xenografts through the recognition of CD70-expressing tumor cells.²⁴⁶

GITRL:GITR Costimulatory Pathway

Glucocorticoid-induced TNFR-related protein (GITR) is a costimulatory receptor expressed on activated T cells and, constitutively, on Tregs. Its ligand GITRL is expressed at low levels on APCs, but it gets induced following TLR stimulation.²⁴⁷ Several studies have reported that GITR signaling promotes the proliferation of naive T cells and cytokines production through the recruitment of TRAF proteins and the activation of downstream pathways. Moreover, one of the first described GITR functions was the ability to protect T cells from activation-induced cell death. Controversial data have been reported about the regulation of Treg functions by GITR. In fact, experimental evidences suggest both an inhibitor and a stimulatory role for GITR. The use of GITR agonistic antibody (clone DTA-1) is also effective in stimulating antitumor immunity *in vivo*.²⁴⁸ The modulation of GITR pathway is an intriguing therapeutic possibility.

The treatment with GITR-expressing adenovirus vector has been shown to be able to induce T-cell response and to reduce tumor size in mice inoculated with B16 tumor cells.²⁴⁹ Nishikawa et al. reported that triggering GITR through GITRL-expressing plasmid resulted in the inhibition of tumor growth in a CMS5 sarcoma model. The protection of CD8 + T cell against Treg-mediated suppression was also observed by the authors.²⁵⁰ Recently, Zhou et al. suggested that the antitumor effect of anti-GITR antibody was dependent on its ability to positive costimulates T cells rather than to suppress Treg functions, but the question is still debated.²⁵¹ Last year, a clinical study of an agonist anti-GITR antibody (TRX518) in melanoma was started but the trial was put on hold because of a major business setback of the company that makes the antibody (see [Table 5.8](#)).

CD30L:CD30 Costimulatory Pathway

CD30 receptor is an inducible costimulatory receptor expressed on activated and memory T cells following TCR/CD28 or IL-4 stimulation. The ligand of

CD30 is CD30L, which is expressed on activated T cells, as well as on macrophages, DCs, and B cells. CD30L/CD30 signaling seems to be involved in Th1 and Th2 cell responses and plays a critical role in Th17 differentiation.²⁵² Due to the expression of CD30 on all malignant HRS cells, this receptor represents an important target for the immunotherapy of hematological malignancies. SGN-35 (brentuximab vedotin) is an anti-CD30 antibody that has been modified by the addition of a dipeptide linker to permit attachment of microtubule polymerization monomethylauristatin E (MMAE)²¹⁰ (also [Table 5.8](#)). SGN-35 has been evaluated in phase I dose escalation study in 45 patients with relapsed or refractory CD30-positive hematologic malignancies and the maximum-tolerated dose was determined to be 1.8 mg/kg.²⁰⁷ In a pivotal phase II study of SGN-35, 102 patients with relapsed or refractory Hodgkin lymphoma were treated with 1.8 mg/kg dose of SGN-35 every three weeks. A reduction in tumor volume was observed in 95% of patients and the ORR was 75%.²⁰⁸ The efficacy of SGN-35 has been also evaluated in a phase II single-arm study in 58 patients with ALCL. The authors reported that the ORR was 86%.²⁰⁹ A multicenter randomized phase III trial of SGN-35 (AETHERA) in posttransplant classical Hodgkin lymphoma patients at high risk for recurrence was started in April 2010.

In the light of these impressive result, the FDA approved SGN-35 for the treatment of Hodgkin lymphoma in August 2011. The high efficacy of this antibody-drug conjugate could be due to the fact that cytotoxic effect of MMAE targets not only CD30-expressing HRS cells, but also the immune-suppressive Tregs present in the tumor microenvironment because of a bystander effect; moreover, SGN-35 delivers itself an additional apoptotic signal, mainly in ALCL cells.²¹⁰

COSTIMULATION IMMUNOTHERAPY AND TUMOR TOLERANCE

Developing tumors are typically recognized and eliminated by a protective mechanism of immunosurveillance, failure of which results in the prevalence of tumor tolerance over immunity.^{253,254} The goal of tumor immunotherapy is to break such tolerance and initiate a robust, prolonged tumor-specific immune response without the harsh toxicities that traditional cancer therapies impart. High-profile accomplishments in the field have recently been made, including FDA-approval of the first cellular vaccine for cancer,²⁵⁵ and demonstration that a small minority of patients with metastatic melanoma can be, for all intents and purposes, cured upon antibody mediated blockade of the T-cell inhibitory molecule CTLA-4.¹²⁶ Median overall survival in both trials, however, was only increased by 4 months. While these milestones should be celebrated as proof-of-concept that T-cell-based immunotherapy of cancer is feasible and relatively effective (compared to standard of care treatment), there is clearly room for improvement.

Blocking negative costimulatory molecules to boost antitumor T-cell responses in cancer patients is an attractive new strategy in cancer immunotherapy. Antibodies that block CTLA-4 on T cells have proven effective in treating melanoma patients.³⁵ The efficacy of CTLA-4 antibodies at least suggests that tumor-specific T cells can be activated in patients bearing tumors. When tumors escape immune surveillance and progress to become a solid cell mass, immune tolerance mechanisms are established in the tumor microenvironment that strongly inhibit infiltrating T-lymphocyte function.²⁵³ For example, prostaglandin E2 secreted by monocytes and TAMs inhibits the proliferation of T cells and promotes T helper (Th)2 responses, which down-regulate the antitumor Th1 response.²⁵⁶

Moreover, prostaglandin E2 promotes the secretion of IL-10 by macrophages, DCs, and tumors, which maintains the immunosuppressive environment in the tumor. Macrophages from tumor stroma also produce the immunosuppressive cytokine TGF- β , which inhibits the antitumor activities of infiltrating CTLs, NK cells, neutrophils, and macrophages. In addition, transforming growth factor- β induces the production of IL-10 by the tumor cells and downregulates the expression of the activating receptor NKG2D—an important regulator of antitumor immune surveillance—on CD8+ T cells and NK cells.^{192,257} B7-H4 is the newest addition to the B7 family that has been shown to negatively regulate T-cell activation. The expression of B7-H4 in multiple nonlymphoid tissues suggests that it might mediate tolerance at the tissue level. In addition, human breast and ovarian cancers have been found to express B7-H4.^{258,259} The cytokines IL-6 and IL-10 present in the ascites were shown to induce expression of B7-H4 on normal blood monocytes. However, tumor ascites had no direct effect on intracellular or cell surface expression of B7-H4 in the tumor cells. The authors also found that the cytokines IL-4 and GM-CSF suppressed IL-10-induced B7-H4 expression. Thus, the downregulation of B7-H4 may contribute to the potent adjuvant effects of GM-CSF in cancer immunotherapy. Expression of B7-H4 by TAMs inhibited the *in vitro* proliferation and effector function of CD8+ T cells specific for the tumor antigen Her-2/neu. Using morpholino antisense oligonucleotides, the authors showed that the immunosuppression was mediated specifically by B7-H4 and not other immunosuppressive effectors known to be expressed by macrophages, such as PDL-1, arginase, or inducible nitric oxide synthase.²⁵⁶ The B7-H4-dependent immunosuppression by macrophages at the tumor site was then confirmed in a mouse ovarian tumor model. Macrophages treated with oligonucleotides to block B7-H4 expression reduced the growth of tumors implanted in lymphocyte-deficient mice that received T cells specific for tumor antigens. This further indicates that B7-H4 is the critical immunosuppressive molecule expressed by TAMs. A possible scenario that might explain these findings is that ovarian cancer cells secrete IL-6 and IL-10 to

induce the expression of B7-H4 on infiltrating macrophages, which would then inhibit the proliferation of and cytokine production by infiltrating T cells via B7-H4. This suggests that blockade of B7-H4 enhances the activity of tumor-infiltrating T cells, shifting the balance of immunity and immune evasion in favor of tumor destruction.¹⁹²

The above experimental evidence reveals the crucial roles of negative costimulators in preventing tissue and tumor attack by infiltrating T cells.^{256,260} Enhancing the function of these molecules may be of benefit in the treatment of autoimmune diseases, whereas inhibition of their function might be an effective therapeutic strategy to boost antitumor immunity. Tumor immunity and autoimmunity are alike to some extent. Tumor reactivity and autoreactivity exist in our TCR repertoire but are restricted by many inhibitory mechanisms such as peripheral tolerance and the action of regulatory T cells. The studies discussed in this chapter, along with others, suggest that further checkpoints exist in tissues and tumors that are mediated by negative costimulatory molecules such as PD-L1 and B7-H4. These studies also suggest that cytokines (pro- or anti-inflammatory) are crucial regulators of the expression of these inhibitory molecules. A lot still needs to be learned about negative costimulation in tissues and tumors and its role in regulating T-cell activity. For instance, is positive costimulation in tissues or tumors required for the function of self- or tumor reactive T cells? And what are the differences in the function and regulation of the various negative costimulators? From our experience in studying costimulation in the context of the regulation of naive T-cell activation, we may expect to observe combinations of various positive and negative costimulatory factors that regulate T-cell activities in tissues and tumors. Thus, it may eventually be possible to design effective immunotherapies that specifically perturb autoimmune responses without impinging on immunity to infection or that break tumor tolerance without provoking autoimmune diseases.

MODULATION OF B7 PATHWAY TO AUGMENT CANCER IMMUNOTHERAPY

The B7 system is one of the most important secondary signaling mechanisms and is essential in maintaining the delicate balance between immune potency and suppression of autoimmunity. Potential therapeutic applications include immune-boosting adjuvants to conventional anticancer therapy, HSCT, anti-tumor vaccines, bioengineered T cells, as well as attenuation of GVHD. The B7 family is only one aspect of a complex signaling network that comprises other IGSF members, the TNFRSF, chemokines, cytokines, and adhesion molecules. However, based on a substantial evidence base and a growing therapeutic armory, the B7 family requires particular attention, as summarized in [Table 5.3](#). The standardized approach to nomenclature is the cluster

of differentiation (CD), which has not yet been applied to all members of the family. While many researchers are more familiar with their original names, which are still widely used in the literature, for the purposes of this review, the CD designation is used where ever possible.

Allogenic HSCT

Failure of allogenic HSCT may in part be mediated by suppressive B7 signaling. Anti-CTLA-4 mAb therapy has been applied successfully in phase 2 clinical trials to treat relapsed CHL, myeloma, and leukemias after allogenic HSCT without the anticipated exacerbation of GVHD seen in murine transplantation experiments.^{19,261} Use of PD-L1/PD-1 axis blockade to enhance allogenic T-cell responses has not yet reached clinical trial, although there is promising preclinical data in a murine AML model using TCR-engineered AML antigen-specific allogenic T cells.²⁰ Late transfer of these cells led to poorer antitumor responses accompanied by PD-1 overexpression. Responses were restored by treatment with anti-PD-L1 mAb. PD-1 blockade, however, accelerated GVHD lethality.²⁶²

Tumor Vaccination Strategies

B7 molecules may play an essential role in cancer vaccination strategies. Mouse model immunization strategies using highly immunogenic cancer cell lines showed early success, particularly with chemokine costimulatory approaches.²⁶³ Tumor lysate-pulsed DCs have been incorporated into some vaccination strategies,²⁶⁴ showing modest success in patients with advanced lymphoma.²⁶⁵ Idiotypic-specific tumor vaccines have been produced for myeloma²⁶⁶ and NHL²⁶⁷ while immunogenic tumor-specific antigens have been identified in CLL and acute leukemia.^{268,269} However, vaccination has yet to make any real impact on cancer therapy.

Combining vaccination strategies with immunosuppressive B7 pathway blockade may improve these disappointing outcomes. CTLA-4 mAb-enhanced melanoma rejection has been demonstrated in a clinical trial,^{36,37,129} but studies in NHL have been disappointing, with minimal responses achieved in vaccinated patients treated with ipilimumab. Encouraging *in vitro* evidence suggests that a similar strategy could be applied to PD-1/PD-L1 axis blockade, although this has yet to be incorporated into any clinical trial. One group showed PD-L1/PD-L2 siRNA silencing improved DC-induced leukemia-specific T-cell responses,²⁶ while another demonstrated enhanced T-cell responses after PD-1 antibody treatment in a myeloma/DC fusion vaccination strategy.²⁵

BIOENGINEERING OF T CELLS FOR CANCER IMMUNOTHERAPY

In contrast to the active immunization strategies outlined in the previous paragraph, an alternative strategy is to expand and reinfuse autologous bioengineered T cells. Autologous TILs extracted from melanoma biopsies, expanded and reinfused into immunosuppressed recipients, have shown some therapeutic success but failed to translate to other malignancies. Attempts to augment autologous T-cell specificity for tumor antigens led to TCR bioengineering approaches, first against Epstein–Barr virus (EBV)-driven lymphoproliferative tumors and subsequently other tumor-specific antigens, while clinical responses are reported, reactive clones tend to rapidly lose efficacy or die off altogether.^{270–274} Subsequent developments used tumor antigen-inoculated transgenic mice whose T cells express human MHC, which expanded tumor-specific T-cell clones and hence derived TCR genes for transfection into patient-derived T cells.²⁷⁵ Newer techniques incorporate novel receptor constructs into autologous T cells using CAR. Here, a rearranged, antigen-specific immunoglobulin variable region gene is combined with intracellular domains of the TCR gene, thus bypassing MHC-restricted antigen recognition. CARs have the advantage of being amenable to further augmentation with costimulatory motifs, of which CD28 and CD80 have demonstrated efficacy.^{23,24} CARs incorporating costimulatory modifications have reached phase I clinical trials in NHL using an anti-CD19-CD28-TCR ζ chain construct, and CLL, using an anti-CD19-CD137 construct²⁷⁶ (a TNFRSF member). Promising results have been observed, even in refractory patients.²⁷⁷ Further optimization is needed and B7 family members will play a central role in this.

ICOS and B7-H2 Costimulatory Pathway in Cancer

Angioimmunoblastic T-cell lymphoma cells express ICOS, PD-1, and CXCR5, consistent with being the malignant counterpart of T follicular helper cells (TFH) lymphocytes.²⁷⁸ ICOS expression has also been described as a potential survival mechanism in nucleophosmin-anaplastic lymphoma kinase (NPM-ALK) fusion ALCL cells.²⁷⁹ Although induced ICOS-L expression enhances cytotoxic T-cell-mediated tumor rejection,^{136–138} there is sparse and conflicting evidence for its expression in solid tumors, while expression in myeloma and acute leukemia is widely heterogeneous. Despite these sometimes contradictory findings, the ICOS/ICOS-L axis may yet have an indirect role in enhancement of tumor immunity. ICOS is inducible on NK cells, which demonstrate cytotoxicity against ICOS-L-transfected murine leukemia cell lines.²⁸⁰ Cytokine-induced killer cells demonstrate MHC-independent cytotoxicity and antitumor activity, enhanced after ICOS

transfection.²⁸¹ Optimal immunostimulatory therapy using CTLA-4 mAbs may be dependent on an intact ICOS/ICOS-L pathway, with CTLA-4 blockade increasing tumor-specific IFN- γ -secreting T cells in cancer patients.^{16,139}

Therapeutic Modulation of ICOS/ICOS-L

ICOS is overexpressed during acute allogeneic solid organ rejection and tolerance is improved with pathway blockade²¹ which suggests that this pathway may have importance in GVHD. Murine acute GVHD is reduced if donor *Icos*^{-co} mice or ICOS-blocking antibodies are used, with less gut and liver GVHD and improved survival^{22,282} despite intact *Icos*^{-co} donor cell cytotoxic function, cytokine-secretion, and tissue trafficking, while *Icos*^{-/-} recipient mice better tolerate allogeneic HSCT with more rapid and complete donor chimerism.²² However, a further study demonstrated more severe GVHD arising from allogeneic CD8+ *Icos*^{-/-} cell transplantation compared with wild type, a contradiction perhaps explained by the differing mouse genetic backgrounds and variable use of Treg-depleting anti-CD25 mAbs.²⁸³ ICOS appears central to GVHD modulation, although because of these controversies, a therapeutic application has yet to emerge.

COMBINATION OF COSTIMULATORY ANTIBODY–LIGAND FUSION PROTEINS FOR TARGETED CANCER IMMUNOTHERAPY

Combinatorial strategies are becoming of increasing interest in cancer immunotherapy. Costimulation by individual members of the immunoglobulin-like (Ig) and TNF superfamily have already shown promising antitumor potential, thus prompting the exploration of their synergistic abilities in combinatorial approaches. Interfering with the costimulatory/inhibitory ligand-receptor network that regulates the immune response has become a promising approach in cancer immunotherapy. Thus, antagonistic antibodies directed against inhibitory receptors of the Ig superfamily (e.g., CTLA-4, PD-1) and agonistic antibodies directed against costimulatory receptors of the TNFR superfamily (e.g., 4-1BB, OX40, GITR, CD127, CD40) are being evaluated in clinical trials.^{284,285} In addition, it is becoming apparent that strongest antitumor effects might be achieved by combinatorial treatments.^{286,287} In preclinical models, synergistic actions have been described for the combined activation of different costimulatory receptors (e.g., 4-1BB with OX40 or CD40)^{229,288,289} as well as for combined activation and blockade of costimulatory and coinhibitory receptors, respectively (e.g., anti-4-1BB mAb with either anti-CTLA-4 or anti-PD-1).^{290–292} The evaluation of indicated combinations is complicated since the expression pattern of these receptors and corresponding ligands orchestrating the immune response is cell-type specific, time-dependent, and cell context-related.¹⁰ Thus, seeking maximal

antitumor potential of costimulatory/inhibitory reagents in combinatorial settings constitutes a major challenge in the field.

For immunotherapeutic approaches, costimulation has been provided either by agonistic antibodies for particular receptors or respective ligands.^{293,294} Studies with ligands involved mainly multimeric recombinant proteins composed of the extracellular domain of the ligand fused to antibody Fc or multimerization domains (e.g., surfactant protein D (SP-D), modified core streptavidin (SA), and tenascin (TNC)).^{295–298} Although effective costimulatory activity was achieved, undirected costimulation bears the risk of unexpected AE and autoimmunity (e.g., liver toxicity observed after anti-4-1BB mAb treatment in preclinical and clinical studies).¹⁷² Thus, targeted approaches are being developed, creating tumor-specific antibody-fusion proteins with costimulatory ligands in order to direct, confine and improve the immune response at the tumor site.²⁹⁵ Enhanced antitumor effects could be shown by target-directed costimulation with B7.1 or 4-1BBL in diverse tumor mouse models. Furthermore, combination therapies with regulatory T-cell (Treg) depletion and a bispecific antibody resulted in improved therapeutic effects.^{299–301}

Previously a combinatorial approach of two costimulatory antibody–ligand fusion proteins, that target B7.2 and 4-1BBL to the tumor cell via two independent antigens, has been reported. In this model system, T cells were retargeted by a bispecific antibody, and stimulation was enhanced by target mediated costimulation of B7.2 and 4-1BBL.³⁰² Here we report further combinations of antibody–ligand fusion proteins with a focus on the combination of B7.1 and 4-1BBL with the costimulatory members OX40L, LIGHT and GITRL of the TNF superfamily, and we will analyze cytokine release, proliferation, and the cytotoxic potential of T cells. Because encouraging results were obtained by the combination of B7.1 and 4-1BBL, we adapted the model system for a time-shift setting, where the benefit of costimulation-assisted restimulation was shown. Finally, the antitumor potential of this combinatorial setting was confirmed *in vivo* in a lung metastasis mouse model. A targeted strategy with antibody-fusion proteins composed of a tumor-directed antibody and the extracellular domain of the costimulatory ligand B7.1, 4-1BBL, OX40L, GITRL, or LIGHT, respectively, was carried out.³⁰³ Costimulatory activity was assessed in an experimental setting where initial T-cell activation was induced by a bispecific antibody (tumor-related antigen 9 CD3). Advantage of combined targeted costimulation was shown for either B7.1 or 4-1BBL with OX40L, GITRL, LIGHT, and 4-1BBL in terms of T-cell proliferation and IFN- γ release. Since encouraging results were obtained by the combination of B7.1 and 4-1BBL, a model system for a time-shift setting was adapted. There was enhanced proliferation and granzyme B expression as well as reduced PD-1 expression on the T-cell population, which demonstrated the benefit of costimulation-assisted restimulation. Finally, the antitumor potential of this combinatorial setting was

confirmed *in vivo* in a lung metastasis mouse model.³⁰³ Thus, combinatorial approaches with costimulatory antibody–ligand fusion proteins seem a promising strategy to be further investigated for cancer immunotherapy.

Thus, for comprehensive costimulatory ligand combination assessment, spatiotemporal considerations need to be taken into account. The data from the time-shift setting system indicate that costimulation by B7.1-Db and scFv-4-1BBL is most effective when applied during scDb-mediated restimulation. In this system, costimulation showed not only the ability to further raise the proliferation and cytotoxic potential of T cells in support of repeated CD3-induced T-cell stimulation, it also showed to counteract the scDb driven enhancement of T cells expressing the inhibitory receptors PD-1, commonly associated with immune restraint. PD-1 is expressed on antigen-experienced T cells in the periphery to limit their activity during inflammatory response and autoimmunity.³⁰⁴ Upregulation of its ligand PD-L1 in tumors has been identified as a mechanism of immune evasion. The importance of this checkpoint receptor in the context of cancer immunotherapy is reflected by the fact that several antagonistic antibodies are in clinical trials.³⁸

It was shown in a B16 lung metastasis tumor mouse model that the application of the costimulatory antibody-fusion proteins could enhance the anti-tumor effect of the bispecific antibody *in vivo*, which provides evidence for the benefit of such a combinatorial approach. This suggests that even powerful tumor-directed polyclonal T-cell activation can benefit from additional costimulation.³⁰³ On the other hand, metastases reduction, although to a much lesser extent was observed for the treatment with the costimulatory fusion proteins only, which indicates also an enhancement of the endogenous tumor-specific immune response in this rather poorly immunogenic model. Thus, the potential to enforce a broad spectrum of cancer immunotherapeutic strategies could be expected. However, extensive *in vivo* studies will be required to understand in detail the underlying immunological mechanisms. By now, the antitumor potential of individual costimulation has been widely acknowledged. Nevertheless, it has also become apparent that it might take combinatorial approaches to fully exploit this strategy. As we could corroborate in this study, the evaluation of configurations and spatiotemporal considerations are crucial. Thus, addressing these issues by establishing experimental settings with costimulatory antibody–ligand fusion proteins constitutes a valuable contribution to the field.

SUMMARY AND CONCLUSION

Improving the knowledge of T costimulatory and coinhibitory pathways over the past decade had shed light on the central roles that these costimulatory molecules play in the generation of an effective immune response. Many tumors escape from immune surveillance through the downregulation of positive costimulatory molecules and the upregulation of coinhibitory

signals. Blockade of coinhibitory pathway on the one hand and the stimulation of the positive signals on the other hand have been found to enhance antitumoral immunity, both alone and in combination with traditional therapy in preclinical and clinical trials. So engaging costimulatory pathways, as well as blocking coinhibitory pathways, has therapeutic potential for treatment of cancer. But for tumor immunotherapy, there are multiple barriers to the antitumor response, and coinhibitory pathways participate in shielding tumors from immune eradication. The differential expression of coinhibitory molecules (such as B7-H4) on certain cancers provides an opportunity for selective immunotherapeutic intervention. Interference with multiple coinhibitory pathways may be needed for optimal therapeutic benefit. Despite the expression of antigens by tumor cells, spontaneous immune-mediated rejection of cancer seems to be a rare event. TCR engagement by peptide/MHCs constitutes the main signal for the activation of naive T cells but is not sufficient to initiate a productive generation and maintenance of effector cells. Full activation of T cells requires additional signals driven by costimulatory molecules present on activated APCs but rarely on tumors. Following the discovery of B7-1 (CD80), several other costimulatory molecules have been shown to contribute to T-cell activation and have relevance for improving antitumor immunity. Moreover, increasing understanding of coinhibitory receptors has highlighted key additional pathways that can dominantly inhibit antitumor T-cell function. Improving positive costimulation, and interfering with negative regulation, continues to represent an attractive immunotherapeutic approach for the treatment of cancer. The pathways with the highest potential for clinical application and translation to the clinic, are clinical trials aimed at boosting positive costimulation and antitumor immunity with agonistic CD40, 4-1BB, or OX40 antibodies, and clinical trials that block the coinhibitory receptors CTLA-4 and PD-1. However, challenges are posed by the development of inflammation or autoimmunity following immune intervention.

As established by clinical and experimental results, it is quite evident that anti-CD80 antibody treatment is effective against relapsed and refractory follicular lymphoma. Future studies in other CD80-positive hematological malignancies (e.g., diffuse large B-cell NHL, CLL, Hodgkin's lymphoma, and MM) should also be taken under consideration. Thus, anti-CD80 antibody immunotherapy may have a potent role in the treatment of CD80-bearing cancer cells because their binding can modulate the key molecules in the signaling pathway and enhance antitumor response. Considering the importance of CD80 signaling in the regulation of immune responses against cancer, the manipulation of this signaling pathway to increase immunity against cancer represents a potential therapeutic approach. Understanding the mechanisms by which tumor cells escape immune surveillance will help us to establish new and effective approaches to vaccination and immunotherapy. One of the most important immune evasion mechanism employed by tumor

cells is the downregulation of the CD80 costimulatory molecule. The success of novel cancer therapies depends on the identification of functional targets that play an essential role in tumor growth and metastasis, survival, and evasion from immunosurveillance. Anti-CD80 therapy can be used to target tumors expressing CD80. The clinical success of anti-CD80 antibody for the treatment of refractory follicular lymphoma has stimulated great interest in the promise of antibody therapeutics for cancer. The qualities of galiximab, like long half-life and high specificity and safety compared with other cancer therapeutics together with its ability to bind to and modulate key players in signaling pathways that drive malignant transformation and enhance antitumor immune functions, makes it a highly desirable therapeutic agent.

Most tumors do not regress but continue to grow in spite of the presence of spontaneous or antigen induced immune responses, due to downregulation of costimulatory molecules like CD80. The existence of systemic immune responses may not by itself be sufficient to deal with the complex nature of tumor–host interactions because factors such as insufficient costimulation to induce T-cell response may further contribute to the lack of effective immunity. It is now well established that T cells are rendered anergic due to the lack of costimulatory molecule(s) expression by tumor cells. Tumor cells are able to induce antigen-specific tolerance or anergy on the basis of MHC-I-restricted antigen presentation without the expression of costimulatory molecules. This unresponsiveness, however, can be reversed when tumor cells are genetically modified to express costimulatory molecules. A plethora of studies suggest that the insertion of genes encoding CD80 into tumors generally increases their immunogenicity and can be used as vaccine. Recently, fusogene vectors were developed to encode multiple gene products like CD80 with cytokines or MHC molecules as fusion proteins from a single cistron to increase the immunogenicity of target tumor cells. The vectors generated could be used in immunotherapy for the treatment of MM, leukemias, and other cancers, as they have been shown to stimulate allogeneic mixed lymphocyte proliferation and augment increases in CTL and NK cell responses and IFN- γ release. Considering the importance of costimulation in the regulation of immune responses against relapsed cancer, the manipulation of this pathway to increase immunity represents a potential therapeutic approach. Furthermore, signaling by either anti-CD80 antibodies or through CD28-bearing T cells regresses the growth, augments the expression of proapoptotic molecules, and induces apoptosis of CD80 + lymphomas. Therefore, immunotherapy utilizing anti-CD80 antibodies is a promising future treatment especially in the case of relapse and refractory lymphomas.

Our understanding of the B7 family has expanded enormously since the costimulatory pathways were first described. Contemporary models of immune regulatory networks reflect this complexity. The promise of immune-mediated enhancement of autologous and allogeneic responses through pharmacologic manipulation of this pathway shows substantial

promise. Ongoing studies investigating the role of the most recently described members of the family: B7-H3, B7-H4, B7-H6, butyrophilin, and Skint will likely reveal further potentially modifiable pathways as therapeutic targets for intervention. Whereas manipulation of the B7 family members and their ligands in suppressing GVHD and augmenting GVL after allogeneic HSCT has shown benefit in murine studies, and promising results in limited phase 2 clinical studies, this remains an area for clinical trial development. The contribution of ICOS/ICOS-L to GVHD appears to be fundamental; however, based on our current understanding of this pathway, the complexities of the interactions make predictions of clinical responses difficult to predict. After decades of study, strategies incorporating B7 family members, such as CTLA-4 and PD-1 pathway-inhibiting molecules and CAR-T cells, have finally demonstrated clear clinical benefit, fulfilling the early promise of the importance of this family in malignancies. Further studies are necessary to evaluate the safety and the efficacy of these approaches before using them in the clinical practice for human use.

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T-Cell Costimulation and Its Applications in Diseases

INTRODUCTION TO T-CELL COSTIMULATION

T cells are dependent on antigen presenting cells (APCs) for at least two distinct stimuli for their optimum activation and effector function. The first signal is provided by the engagement of clonotypic T-cell receptor (TCR) with major histocompatibility complex (MHC)–peptide complex on APCs. The second signal is non-MHC-restricted and delivered by costimulatory molecules. The best-defined costimulatory molecules are the CD80 and Cd86 on APCs and their ligands CD28/CTLA-4 on T cells. The significance of costimulatory molecules in T-cell activation gained considerable impetus following the observation that occupancy of TCR alone is generally inadequate for exerting complete T-cell activation. Thus an encounter with an antigen can lead to two quite distinct outcomes in T cells: proliferation and differentiation into effector cells; or inactivation or death. Which outcome occurs is determined by the appropriate delivery of costimulatory signals.^{1–4}

Activation of T lymphocytes is a critical component of the immunological response to foreign protein molecules. TCR recognition of peptide-loaded MHC molecules provides antigen specificity and initiates the required steps for T-cell activation, although additional signals are needed for complete T-cell activation. A two-signal model for lymphocyte activation was first proposed as a mechanism for self:nonsel self discrimination in B cells by Bretscher and Cohn in 1970⁵ and was subsequently expanded to incorporate CD8 T-cell activation by Lafferty et al.⁶ in 1975. They showed that engagement of the TCR by peptide-loaded MHC in the absence of other signals is insufficient to activate the T cell, and in fact may render it unresponsive to further antigenic stimulation, a condition termed anergy.^{7,8} They further showed that provision of additional, non-MHC-restricted signals provided by professional APC, resulted in the T cell becoming fully activated leading to clonal expansion and development of effector function. Thus, the two-signal

model, where signal one is defined as the binding of peptide/MHC complexes and signal two refers to the additional, costimulatory signal(s), provided a potential mechanism for peripheral tolerance.^{9,10} The most studied and well characterized costimulatory receptor/ligand complex is the CD28/B7 interaction.^{11,12}

COSTIMULATION MEDIATED BY CD28 COSTIMULATORY PATHWAY IN T CELLS

CD28 was first identified by antibodies recognizing a 44 kDa protein on the surface of human T cells as type 1 transmembrane protein. In humans, CD28 is expressed on approximately 80% of CD4+ T cells and 50% of CD8+ T cells and has been detected on plasma cells, neutrophils, and eosinophils.^{13,14} In contrast to humans, in mouse 100% of both CD4 and CD8 T cells express CD28.¹⁵ CD28 is the founding member of the immunoglobulin (Ig) family of costimulatory receptors that now includes the receptors: cytotoxic T lymphocyte-associated antigen 4 (CTLA-4, CD152), inducible costimulator (ICOS), programmed death receptor 1 (PD-1), and B & T lymphocyte attenuator (BTLA).

CD28 signaling causes the initial activation of naive CD4 T cells by increasing the sensitivity of the T cell to antigen receptor engagement, and as a result proliferation is induced at otherwise submitogenic concentrations of antigen.^{16,17} Cytokine production is greatly increased, most significantly IL-2. CD28 costimulation results in a 50-fold increase in IL-2 secretion by both transcriptional and posttranscriptional regulation of expression.¹⁸ In addition, cell survival is enhanced by CD28 costimulation, in part by inducing expression of anti-apoptotic proteins including Bcl-X_L.^{19,20} Thus, by a variety of mechanisms, CD28-mediated costimulation greatly enhances the T cells effector response to antigen.

Whether CD28 can activate signaling independent of TCR engagement or function only in a TCR dependent manner has been controversial. The ability of certain “superagonistic” anti-CD28 antibodies to independently activate T cells suggests that in some circumstances CD28 can function in the absence of a TCR derived signal.²¹ Importantly, the existence of both TCR dependent and independent pathways is not mutually exclusive, and it is likely that both are operative and important.

CD28 COSTIMULATION ENHANCES T-CELL SURVIVAL

A major consequence of CD28 signaling is protection from cell death. The two dominant pathways of apoptosis in T cells are (1) receptor mediated (i.e., tumor necrosis factor (TNF) family) or (2) mitochondrial-associated proteins (i.e., the Bcl family) both of which are modulated via CD28 signaling.¹¹ CD28 costimulation prevents activation induced cell death (AICD) by inhibiting the expression of CD95L expression (*cis*-mediated death) on the

T cell and by decreasing the formation of the death-inducing signaling complex (DISC). CD28 stimulation increases the protein expression of c-FLIPs, which compete with procaspase 8 for binding to the death domains of CD95 and FADD, and thereby preventing the conversion of procaspase 8 to active caspase 8 and the induction of cell death.²² CD28-deficient cells are more sensitive to Fas-mediated cell death. CD28 stimulation also induces the upregulation of anti-apoptotic factors including Bcl-X_L. The overexpression of Bcl-X_L in CD28-deficient mice restores cellular survival but not proliferation; thereby, separating these two CD28-dependent functions.^{19,23}

CD28 T-CELL COSTIMULATION IN DISEASES

Perhaps surprisingly, mutations in CD28 have not been identified as the genetic basis for any human disease. However, mouse models of disease have been shown to depend on CD28 function. The use of CD28- and B7-deficient cells has implicated this pathway in the generation of a TH2 response required for the destruction of islet cells (autoimmune diabetes in the NOD mouse model)¹ and in the activation of newly recruited T cells responsible for epitope spreading (relapsing EAE in the EAE mouse model).²⁴ CD28 has been shown to be required in a model of allergic airway inflammation.^{25,26} In the absence of CD28 or in the CD28-AYAA knockin mice, antibody production and germinal center formation is markedly reduced.²⁷ Thus, in a number of model systems, CD28 plays a critical role.

So, an understanding of costimulation and the mechanism by which it regulates immune responses provides new avenues for the development of therapeutics. As CD28 decreases the threshold required to activate naive T cells, as well as increasing the survival of activated T cells, one can envision development of drugs that either potentiate or downregulate an immune response. Although global inhibition of CD28 function would be anticipated to profoundly inhibit T-cell function, more specific targeting of specific regions of the extracellular or intracellular regions of CD28 via antibodies, small molecule mimetics or siRNAs could potentially alter some aspects of CD28 effects whereas leaving others intact.

CD28 COSTIMULATION IN PRIMARY T-CELL RESPONSES

The importance of CD8 + T cells in the resolution of viral infection is widely accepted. Activation of naive CD8 + T cells during virus infection occurs in local draining lymph nodes, where dendritic cells (DCs) present viral antigens to CD8 + T cells.^{28,29} During naive CD8 + T cell–DC interactions, costimulatory signals delivered by molecules such as CD28 (signal two) determine whether CD8 + T cells will become activated and expand, or they will be suboptimally activated. Studies examining primary infection of mice with viruses such as VSV, MHV-68, and influenza type A virus,

indicated that CD28 was required for primary expansion of antiviral CD8+ T cells.^{30–33} In one of the earliest CTLA-4-Ig blocking studies, using influenza virus, Lumsden et al. identified that the loss of CD28 signaling negatively impacted both CD4+ and CD8+ T cells.³⁴ In this study, there was a significant decrease in the production of antiviral antibodies, decreased expansion of virus-specific CTLs, and a loss of IFN- γ and cytotoxic function by those cells which did expand. Ultimately these CTLA-4-Ig blocked mice resolved the infection, yet it was delayed in comparison to controls. In a complimentary study by Bertram et al., influenza virus-infected CD28 $-/-$ mice exhibited substantially decreased expansion of virus-specific CD8+ T cells at the peak of the primary response whether virus was delivered intraperitoneally or intranasally.³² Halstead et al. also showed that both dominant and subdominant primary CD8+ T-cell responses against influenza virus are greatly reduced in CD28 knockout mice.³³ In contrast, studies using lymphochoriomeningitis virus (LCMV) infection of mice, initially showed that an efficient primary CD8+ T-cell response could be generated in the absence of CD28 costimulation. CD28 knockout mice (CD28 $-/-$) were infected with LCMV, and despite the absence of CD28 signaling, virus CD8+ T cells expanded and viral burden was eliminated at levels comparable to wild-type controls. Similar CD8+ T-cell expansion was observed against all measured epitopes of LCMV, including subdominant epitopes.^{35,36} The reason for this discrepancy in the lack of requirement for a CD28-mediated signal in LCMV infection became apparent from studies that showed that if sufficiently high levels of TCR stimulation were obtained, the need for costimulation could be overcome.³⁷ Viola and Lanzavecchia elegantly showed in *in vitro* studies, that independent of the nature of the TCR stimuli, TCR stimulation must exceed a minimum threshold in order to achieve complete activation of a T-cell clone. However, in the presence of CD28 costimulation, that threshold is significantly lowered.³⁸ Kundig et al. utilized LCMV infection and showed that the disparity in requirement for CD28 in primary LCMV infection versus VSV infection was due to differences in TCR signal duration.³⁷ Indeed, of all the viruses examined, LCMV is the only virus whose natural host is the mouse and therefore it replicates much more rapidly and extensively than any of the other viruses examined. As a result, antigen presentation persists for a longer period of time and at higher levels, providing a strong and sustained TCR signal which overcomes the need for CD28 costimulation.³⁷

When viral infection is cleared the effector T-cell population undergoes a steady contraction until a small stable memory pool is formed. Memory cells respond faster and more effectively in the event of secondary insult to the host.³⁹ One of the major contributing factors to the rapidity of memory T-cell responses is their potentially higher affinity for antigen that leads to a lower threshold of activation.^{40,41} Given that the strength of TCR signaling and predetermined threshold of activation can affect the need for

costimulation, it is reasonable to question whether or not memory T cells have a requirement for CD28 costimulation during re-activation.

CD28-B7 Costimulation and Its Effect on CD8 + Memory T-Cell Responses

Costimulation signals have been recognized as critical for optimal T-cell responses and result from important interaction between receptors on the surface of T cells and their ligands on APCs. CD28 receptor costimulatory family has been found to be a major player in providing costimulation to CD8+ T cells. Recent studies using viral infection models have highlighted the importance of CD28 costimulation signals during memory responses against viruses. PD-1 another member of the CD28 family may contribute to functional defects of helpless memory CD8+ T cells. The delivery of costimulatory molecules such as CD28, 4-1BB, and OX40 can help boost the generation and function of virus-specific memory CD8+ T cells. Taken together this suggests that the use of costimulatory molecules as adjuvants along with viral antigens in vaccines may facilitate the generation of effective antigen-specific memory CD8+ T-cell responses. Understanding the costimulatory requirements of memory CD8+ T cells therefore may lead to improved vaccines that target antiviral CD8+ T-cell memory.

After pathogen clearance effector T cells undergo contraction and a memory T-cell pool is formed. These memory cells respond faster and more effectively in the event of secondary insult to the host because of their higher precursor frequency and higher affinity for antigen that leads to a lower threshold of activation. As the requirement for costimulation affects the strength of TCR signaling and the threshold of activation it is important to determine whether or not memory T cells have a requirement for CD28 costimulation during re-activation. Early studies relied on *in vitro* experiments to address this. More recently however, the availability of new reagents and of genetically modified mice have allowed the direct assessment *in vivo* of the requirement of CD28 signaling by memory CD8+ T cells. During antiviral CD8+ T-cell responses costimulatory signals from the CD28 family affect different phases of the immune response. It has now become apparent that different members of this family play important roles in the initiation phase, the generation and maintenance of memory, quality of memory, and the secondary response.

Memory CD8+ T-cell immune responses have been considered for a number of years not to require costimulation based on several studies that used *in vitro* systems of restimulation or CD28-deficient mice.^{42,43} It has been demonstrated recently that CD28 plays a critical role in the secondary CD8+ T-cell response,^{44,45} thus challenging the notion that costimulation is not required by memory CD8+ T cells. To circumvent the problem of generating virus-specific memory cells in CD28-deficient mice

that have an impaired primary response, memory CD8⁺ T cells, were generated in wild-type mice by *in vivo* viral infections. Following the development of intact primary T-cell responses, and the generation of memory CD8⁺ T cells, the requirement of the memory population for CD28 costimulation during a secondary response was examined by blocking CD28 binding to its ligands with either CTLA-4-Ig, anti-B7 or anti-CD28 monoclonal antibodies or transfer of memory cells to CD80/CD86 double deficient mice (B7.1 and B7.2 knockouts).^{44,45}

Interestingly, virus-specific memory CD8⁺ T cells that were generated in the absence of CD28 costimulation expanded approximately 9 times, whereas virus-specific memory CD8⁺ T cells that were generated in C57Bl/6 mice expanded more than 40 times. These findings were further substantiated when virus-specific memory CD8⁺ T cells were generated in C57Bl/6 mice and transferred into CD80/CD86-deficient mice or wild-type control mice, challenged, and analyzed five days later for expansion of the virus-specific memory population. The importance of CD28 signaling during secondary responses was further supported with vaccinia virus infections where the lack of CD28 signaling impaired the responses of memory virus-specific CD8⁺ T cells.⁴⁵

Thus the CD28 requirement of memory CD8⁺ T cells for expansion has been shown in multiple viral infections such as influenza type A virus, HSV, vaccinia virus, and murine gamma herpes virus and is required for rapid pathogen clearance. The reduced expansion of virus-specific memory CD8⁺ T cells in the absence of costimulation challenges the paradigm that memory immune responses occur independently of costimulatory signals. CD28 signaling during primary response may be affecting the quality of memory CD8⁺ T cells generated while the expansion of memory T cells clearly requires CD28 costimulation signals for optimal secondary responses and normal pathogen clearance.^{44,45} This CD28 requirement by memory cells is not restricted to CD8⁺ T cells, CD4⁺ T cells have also been shown to require *in vivo* CD28 costimulation.⁴⁶ This information has important implications in designing efficient vaccination strategies against pathogens and tumors that can downregulate costimulatory signals.

The Future Prospectus

The need for potent CD8⁺ T-cell eliciting vaccines remains largely unfulfilled. Discovery of adjuvants and strategies that enhance the generation, maintenance, and quality of memory CD8⁺ T cells is important for the development of effective vaccines against viruses, intracellular pathogens, and tumors. Costimulatory molecules from the CD28 and the TNFR family augment the immune response during viral infections and contribute to different phases of CD8⁺ T-cell responses as seen in [Figure 6.1A–C](#). It appears that the CD28 and CD27 are required to initiate a primary CD8⁺ T-cell response.⁴⁷

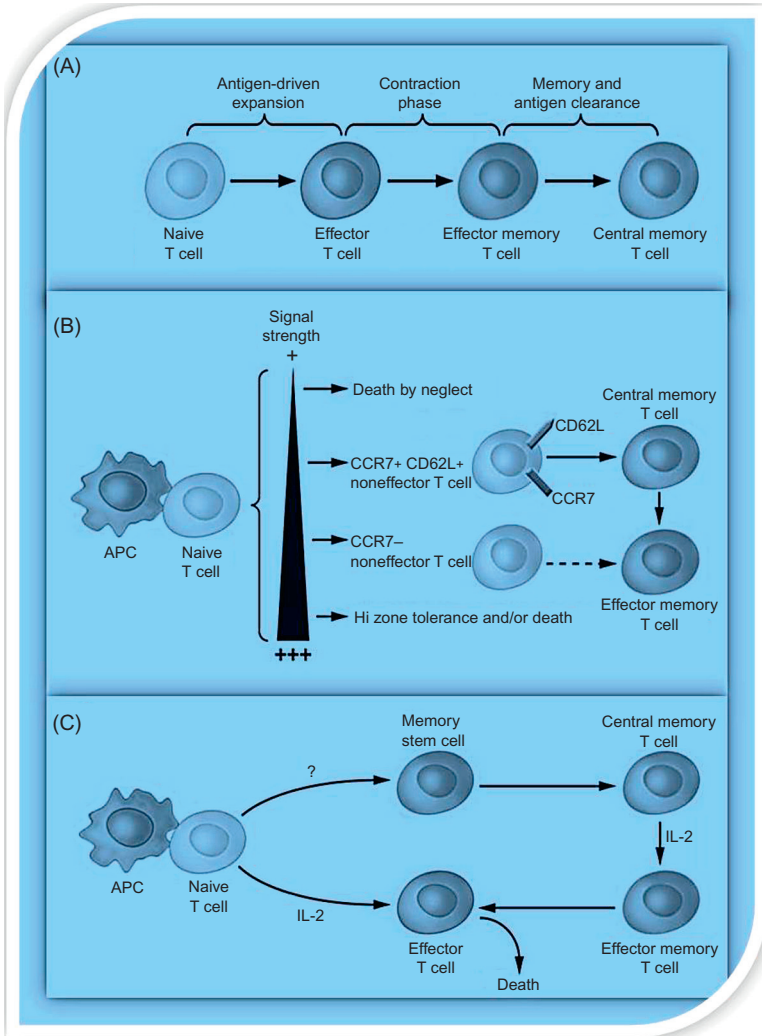


FIGURE 6.1 (A–C) Models of CD8+ T-cell differentiation to distinct memory cell subsets. (A) In the linear differentiation model, an autonomous antigen-triggered differentiation process consisting of conversion from naive to effector to TEM cell occurs, followed by the appearance of TCM cells after antigen clearance through a process of dedifferentiation.¹¹⁷ (B) The signal strength model proposes that naive T cells progress through hierarchical thresholds for proliferation and differentiation as the strength and duration of the interaction with APCs is increased. T cells receiving the weakest signals do not survive, whereas high-intensity signaling causes the development of terminally differentiated effector T cells that cannot survive into the memory phase. The TCM cells, being the least differentiated of the antigen-stimulated T cells, retain the developmental options of naive T cells, including their capacity for marked clonal expansion.^{118,119} (C) The memory stem cell model proposes that the cells within the TCM cell compartment are self-renewing and serve as a source of effector T cells.^{197,198}

For secondary responses of memory CD8+ T cells, costimulatory molecules such as CD28, 4-1BB, OX40, and CD27 all seem to play a role in shaping the memory responses either by providing pro-survival signals or by enhancing the quality of the memory CD8+ T cells. Secondary expansion of memory CD8+ T cells requires professional APC66,⁴⁸ and CD28 costimulation.^{44,45} TNFR family members as costimulatory molecules could also be used as adjuvants. Indeed DNA and adenovirus-based vaccines have shown that expression of 4-1BBL, OX40L, and CD70 leads to increased T-cell expansion, enhanced CTL activity, and antibody response.^{49,50} Agonistic antibodies to TNF family members such as 4-1BB can also provide an adjuvant effect and enhance memory CD8+ T-cell generation.⁵¹ Direct delivery of ligands such as using 4-1BBL to “decorate” tumor cells may provide costimulatory signals that enhance antitumor CTL.^{52–54} An alternative strategy may be the use of oligonucleotide-based ligands known as aptamers, an example being multivalent 4-1BBL aptamers that act as agonists that can directly trigger CD8+ T cells and inhibit tumors in mice.⁵⁵ Chronic viral infections are characterized by accumulation of functionally impaired antigen-specific CD8+ T cells, and studies have shown that 4-1BBL in combination with CD80 can induce the expansion of the antigen-specific CD8+ T cells from this impaired pool.⁵⁶ Therefore such strategies may prove valuable for the design of effective vaccines not only against acute viral infections but also against chronic viral infections.

CONCLUDING REMARKS

Animal models of viral infections have shown that costimulatory molecules of both the CD28 and the TNFR family, as indicated in [Table 6.1](#), help in the generation and maintenance of virus-specific memory CD8+ T cells, but are also important for the re-activation of memory CD8+ T cells and secondary responses. These costimulatory molecules may act in a redundant fashion, but they may also provide an opportunity to augment virus-specific memory CD8+ T-cell responses and may prove useful in designing effective vaccines against chronic viral infections such as HBV and HIV.⁵⁷ A number of questions still remain about the function of costimulatory molecules in memory antiviral CD8+ T-cell responses (see “Unanswered Questions” below). The function of CD28 and TNFR family members has been studied for RNA and DNA viruses, and studies have been done to delineate their functions at the acute and chronic stages of disease. However the redundancy in the function of these molecules questions whether there is a hierarchy in the expression and function of the costimulatory molecules. Although agonistic antibodies may enhance responses when CD28 signaling is absent.⁵⁸ This does not necessarily suggest a switch in costimulation requirements. It may however suggest that the lack of critical costimulatory molecules may be overcome by targeting alternative receptors with monoclonal antibodies or aptamers.

TABLE 6.1 Chimeric Antigen Receptor (CAR) T Cells Using CD28 Costimulatory Molecules in Clinical Trials

Cancer	CAR Target	CAR Endodomains	Number of Patients	Clinical Outcomes
Lymphoma	CD19	CD28 and CD3- ζ	1	PR
Lymphoma	CD19	CD28 and CD3- ζ	4	3 PR, 1 died of influenza
CLL	CD19	CD28 and CD3- ζ	4	1 CR, 2 PR, 1 SD
CLL	CD19	CD28 and CD3- ζ	4	1 CR, 1 PR, 2 PD
Lymphoma	CD19	CD28 and CD3- ζ	6	1 PR, 5 SD
CLL	CD19	CD28 and CD3- ζ	8	1 PR, 2 SD, 3 NR, 1 PD, 1 died of sepsis-like disease (symptoms preceded T-cell transfer)
ALL	CD19	CD28 and CD3- ζ	1	CR
ALL	CD19	CD28 and CD3- ζ	5	5 CR
ALL	CD19	CD28 and CD3- ζ	16	10 CR, 4 CRi, 2 NR
CLL	CD19	4-1BB and CD3- ζ	1	CR
CLL	CD19	4-1BB and CD3- ζ	3	2 CR, 1 PR
ALL	CD19	4-1BB and CD3- ζ	2	2 CR

CLL, chronic lymphocytic leukemia; ALL, acute lymphocytic leukemia; PR, partial response; CR, complete response; SD, stable disease; PD, progressive disease; NR, no objective response; CRi, complete response with incomplete count recovery; NED, no evidence of disease.

The recent discovery of the important role of costimulatory molecules such as CD28 during re-activation of memory CD8 + T cells would raise the danger that therapeutic blocking of costimulation in transplantation and autoimmunity may impair the host's ability to respond to viral infection. On the other hand, some viral infections manipulate costimulation to evade the immune response. A number of viruses such as measles and HIV down-regulate B7 family members so as to hamper CTL responses.^{59–61} This would suggest that successfully priming CTL vaccines may not be very effective against such viruses. A strategy against such pathogens and tumors may be to “add back” costimulatory ligands to reverse the impairment of CTL responses, or to generate memory, if possible, which is less dependent on CD28 costimulation. One of the hallmarks of chronic viral infections is the occurrence of systemic inflammation; the effect of different pro-inflammatory cytokines on the expression of costimulatory receptors and their ligands during different phases of viral infections is still poorly understood.

The expression of TNFR or CD28 costimulatory molecules or their corresponding ligands on the APCs is necessary for an effective antiviral CD8 + T-cell response, especially for vaccinations against viral infections. The important interplay between different costimulatory molecules for the generation, maintenance, and functionality of memory CD8 + T cells, indicates that the design of preventive vaccines will require we further understand the important contributions of costimulation to effective antiviral memory CD8 + T-cell responses.

UNANSWERED QUESTIONS

- Can one cell provide multiple costimulatory signals?
- Do different APC provide different costimulatory signals?
- What is the importance of redundant signaling?
- Do central and effector memory CD8 + T cells differ in their requirement for costimulation?
- Does persistent viral load alter expression of costimulatory molecules on APC?
- What is the effect of costimulation on tertiary memory?
- Does chronic infection change costimulation requirements of CD8 + T cells?
- Can costimulatory ligand combination enhance adjuvant activity?

PROMOTING T-CELL FUNCTION BY MODULATING COSTIMULATION OR CO-INHIBITION

The development of an antitumor T-cell mediated immune response is a multistep process in which tumor-associated antigens (TAAs) expressed by the tumor cells are processed and presented by professional APCs to circulating T cells, which become activated. The quality and potency of this T-cell

response will depend on the nature of the antigen presented, the functionality of the APC itself, and the cytokines and costimulatory interactions that occur in the T-cell/APC microenvironment.⁶² Once activated, TAA-specific effector T cells can migrate to the tumor site and target malignant cells expressing the cognate antigen epitope displayed in the context of a MHC molecule.

Optimally effective cancer immunotherapy may require combinatorial approaches. Synergistic combinations could include means of activating T cells, such as vaccination or the transfer of *ex-vivo* activated and expanded tumor-reactive T cells, coupled with means of sustaining T-cell activation, such as agonistic antibodies to costimulatory receptors or blocking antibodies to co-inhibitory receptors. As many clinical-grade reagents are now available, potential combinations are abundant. Currently, there are over 20 registered clinical trials to study the combination of ipilimumab with vaccines, radiation, chemotherapy, hormonal therapy, cytokines, anti-PD-1, and targeted therapy agents in patients with a broad range of cancers.

Immune activation is tightly regulated by co-receptors expressed on T cells (Figure 6.1A–C). Costimulatory receptors include CD28 and ICOS (inducible T-cell co-stimulator) of the Ig superfamily, as well as 4-1BB, OX40, CD27, CD30, CD40, GITR (glucocorticoid inducible TNF receptor-related protein), and HVEM (herpes-virus entry mediator) of the TNFR superfamily.^{63,64} These costimulatory signals are counterbalanced by co-inhibitory members of the Ig superfamily including CTLA-4, PD-1, BTLA (B and T lymphocyte attenuator), lymphocyte activation gene-3 (LAG-3), TIM3 (T-cell immunoglobulin and mucin domain-containing protein 3), and VISTA (V-domain immunoglobulin suppressor of T-cell activation) on T cells.^{65–69} The idea of blocking the immune co-inhibitors as a therapeutic anticancer strategy was suggested by James Allison over a decade ago.⁷⁰ Anti-CTLA-4 was used as a prototype but antibodies that either stimulate costimulatory TCRs or block other inhibitory immune-checkpoint molecules have been examined more recently.

TURNING ON THE STIMULATORS: ANTIBODIES TO 4-1BB (CD137), OX40 (134), GITR, AND CD40

Animal models and clinical trials have focused on targeting the 4-1BB, OX40, GITR, and CD40 members of the TNFR superfamily. 4-1BB is expressed on activated T cells, activated NK cells, DCs, and endothelial cells.⁶² 4-1BB enhances T-cell function by promoting T-cell survival and memory generation,⁷¹ activating APCs and endothelial cells, reducing T regulatory cell (Treg) suppressive capacity, and enhancing effector T-cell resistance to Treg suppression.^{72,73} Agonistic anti-4-1BB monoclonal antibodies alone or in combination with vaccination can enable rejection of syngeneic tumors in preclinical models.^{74–77} Phase I and II clinical trials with a fully humanized monoclonal anti-4-1BB antibody in patients with melanoma and

renal cell carcinoma (RCC) showed only very mild antitumor activity,⁷⁸ suggesting that anti-4-1BB antibody may require combinations with other agents in order to demonstrate relevant antitumor activity.

OX40, expressed on activated CD4+ and CD8+ T cells, functions as a late costimulatory receptor.^{62,79} Ligation of OX40 enhances immune responses by prolonging CD4+ and CD8+ T-cell survival and memory generation, which prevents T-cell tolerance that impairs the suppressor function of Treg and suppresses the generation of inducible Treg. An agonistic anti-OX40 antibody increases antitumor activity in a number of animal models.^{80–83} Other reports indicate that anti-OX40 antibodies can work in combination with anti-4-1BB or vaccination to augment antitumor immune responses.^{84,85}

GITR is also expressed after T-cell activation, is constitutively expressed at high levels on Tregs, and can be further induced after Treg activation.^{48,86} GITR enhances proliferation and function of CD4+ and CD8+ T cells, inhibits Treg function, and elicits effector T-cell resistance to Treg-mediated suppression.^{87,88} GITR may increase intratumor effector T cells to Tregs ratio and anti-GITR antibodies elicit antitumor responses in a number of animal models.^{89,90} In addition, anti-GITR antibodies can be combined with vaccination, chemotherapy, or anti-CTLA-4 antibody to augment antitumor activity.⁹¹

CD40 is mainly expressed on APCs and endothelial cells, and its binding to its ligand (CD40L/CD154) on activated T cells results in persistent T-cell activation as well as expansion and activation of APCs. An agonistic anti-CD40 antibody has been shown to cause regression of pancreatic cancer in both human and mice via activation of macrophages, suggesting a T-cell independent antitumor mechanism.⁹² CD40 is also expressed on many types of tumor cells and its ligation may promote tumor cell apoptosis and growth arrest. An initial phase I clinical trial using an agonistic anti-CD40 antibody in patients with advanced solid tumors demonstrated modest antitumor activity.⁹³ Overall, agonistic antibodies against immune co-stimulators have demonstrated modest antitumor activities in early phases of clinical trials. The combination of agonistic antibodies with other treatment modalities may be needed to achieve optimal antitumor activity.

TURNING OFF THE BRAKES: ANTIBODIES AGAINST CTLA-4 (CD152), PD-1 (CD279), AND PD-L1

CTLA-4 is expressed by activated CD4+ and CD8+ T cells. Upon T-cell activation, CTLA-4 is rapidly mobilized from intracellular vesicles to the immune synapse where it outcompetes co-stimulator CD28 for binding to its ligands B7-1 and B7-2 and, as a result, ablates T-cell activation (Figure 6.2B). In addition, CTLA-4 mediates cell cycle arrest and is essential for Treg suppression.⁶³ In human studies, anti-CTLA-4 enhances frequency of CD4+ and CD8+ T cells as well as the antibody response to tumor antigens.^{94,95} In animal models, blocking monoclonal antibody against CTLA-4

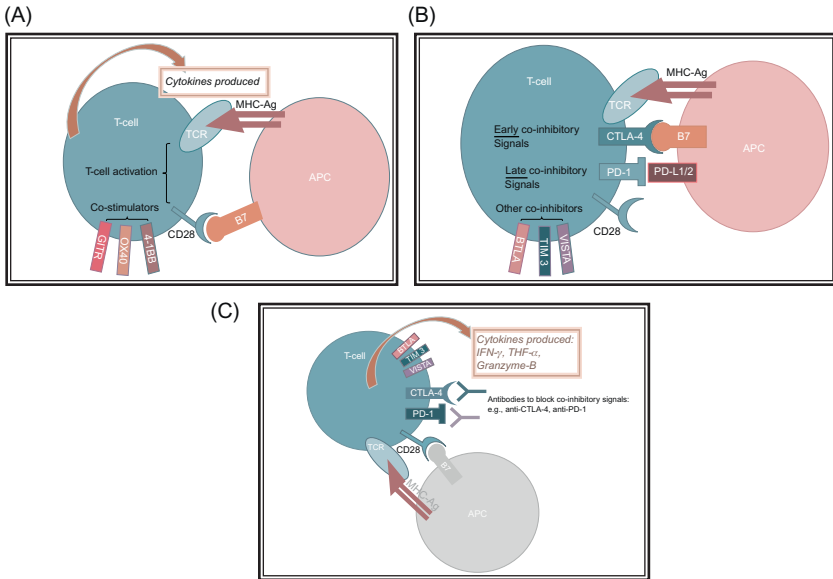


FIGURE 6.2 Modulation of T-cell activation and current strategies promoting effector T-cell functions. (A) Augmenting T-cell activation by positive costimulation. Antigenic presentation triggers T-cell activation and occurs when a peptide bound to MHC molecule on an APC interacts with T-cell receptor (TCR) on the surface of a T cell. In order to achieve optimal activation, additional costimulatory signals are required and primarily involve interaction between CD28 on T cells and B7 on APCs. Other T-cell positive co-stimulators include 4-1BB, OX40, and GITR. (B) Limiting T cell activation by negative costimulation. After T cell activation, cytotoxic T lymphocyte-associated protein 4 (CTLA-4) is mobilized to the cell surface and binds to B7 with greater affinity than CD28 and therefore prevents signaling through CD28. Later inhibitory signals can be provided by co-inhibitors such as programmed cell death-1 (PD-1), which binds to PD-1 ligand 1 (PD-L1). Other co-inhibitors of T cell activation include VISTA, TIM3, and BTLA. (C) Sustaining T-cell activation through blockade of negative costimulatory molecules. Blocking antibodies against CTLA-4 or PD-1 are currently employed to neutralize co-inhibitory receptors and prevent dampening of the T-cell response. Blockade of these inhibitory immune checkpoints results in enhanced and sustained activation of tumor-specific T cells that produce cytokines including TNF- α , interferon- γ (IFN- γ), and granzyme-B.

promotes antitumor activity.^{63,96,97} Phase I and II clinical trials indicate that an anti-CTLA-4 antibody, ipilimumab, has significant antitumor activity in patients with advanced melanoma.^{98,99} A phase III randomized controlled trial in patients with metastatic melanoma showed that ipilimumab improved median overall survival by 3.7 months,¹⁰⁰ leading to FDA approval of ipilimumab for the treatment of patients with metastatic melanoma. A second randomized phase III clinical trial recently showed that addition of ipilimumab to standard dacarbazine chemotherapy improved overall survival by 2.1 months in patients with metastatic melanoma.¹⁰¹

PD-1 is expressed by activated CD4⁺ and CD8⁺ T cells, NK T cells, and APCs. It has two ligands, PD-L1 and PD-L2, with distinct expression profiles.¹⁰² PD-L1, the main target for PD-1, is expressed broadly on T cells, B cells, APCs, and nonhematopoietic cells. PD-L2 is largely restricted to activated macrophages, myeloid DCs, and mast cells. PD-1^{-/-} mice develop autoimmunity with high titers of autoantibodies, consistent with PD-1's role as an inhibitor of T and B cells.^{103,104} Anti-PD-1 antibodies reduce tumor metastasis and growth in animal models,^{105,106} whereas forced expression of PD-L1 in murine tumor cell lines enhances *in vivo* tumor growth.¹⁰⁷ In a phase I trial in 39 patients with refractory metastatic melanoma, colorectal cancer, prostate cancer, nonsmall-cell lung cancer (NSCLC), or RCC, anti-PD-1 antibody (MDX-1106) resulted in complete or partial response in three patients and less significant tumor regression in two other patients.¹⁰⁸ A recent phase I clinical trial with an anti-PD-1 antibody (BMS-936558) showed promising antitumor activity in patients with advanced NSCLC, RCC, and melanoma.¹⁰⁹ Another phase I trial with anti-PD-L1 antibody showed mild to modest antitumor activity in patients with advanced NSCLC, melanoma, and RCC.¹¹⁰ Blocking antibodies against other T-cell co-inhibitors including LAG-3 and TIM3 are still at early stages of development as potential anticancer agents.^{65,69}

ADOPTIVE T-CELL TRANSFER

The adoptive transfer of T cells is a promising approach to treat cancers. Primary human T cells can be modified using viral and non-viral vectors to promote the specific targeting of cancer cells via the introduction of exogenous TCRs or chimeric antigen receptors (CARs). This gene transfer displays the potential to increase the specificity and potency of the anticancer response while decreasing the systemic adverse effects that arise from conventional treatments that target both cancerous and healthy cells.

The transfusion of T cells, also called adoptive T-cell therapy, is an effective treatment for viral infections and has induced regression of cancer in early stage clinical trials. Adoptive T-cell therapy for cancer is a form of transfusion therapy consisting of the infusion of various mature T-cell subsets with the goal of eliminating a tumor and preventing its recurrence. However, recent advances in cellular immunology and tumor biology are guiding new approaches to adoptive T-cell therapy. For example, use of engineered T cells is being tested as a strategy to improve the functions of effector and memory T cells, and manipulation of the host to overcome immunotoxic effects in the tumor microenvironment has led to promising results in early stage clinical trials.^{111–113}

Allogeneic and autologous sources of T cells derived from several anatomic sites have been tested. Indeed, in the 1970s, Chester Southam and colleagues demonstrated that subcutaneous growth of human tumor autografts to patients bearing advanced cancers was inhibited by the co-transfer of

autologous leukocytes in about half of the patients.¹¹⁴ This finding suggested that lymphocytes with a specific inhibitory effect on the implantation and growth of cancer cells were present in many patients and could be mined as potential candidates for adoptive immunotherapy. The primary advantage of using CD8+ T cells for adoptive therapy, as opposed to other cytolytic cells, such as NK cells, is their ability to specifically target tumor cells through the recognition of differentially expressed tumor proteins presented on the cell surface. Using T cells for adoptive therapy is also attractive due to the long clonal life span of T cells,^{115,116} which allows both therapeutic and immunoprophylactic scenarios to be envisioned. In addition, T cells are well suited for genetic manipulation, which has enabled the evaluation of genetically enhanced or retargeted T cells in pilot clinical trials for cancer as well as other diseases.

Adoptive T-cell therapy depends on the ability to optimally select or genetically engineer cells with targeted antigen specificity and then induce the cells to proliferate while preserving their effector function and engraftment and homing abilities. T-cell populations are heterogeneous and comprise memory cells, effector cells, and Tregs exist in several distinct stages of differentiation. Naive CD4+ and CD8+ T cells undergo unique developmental programs after antigen activation, which results in the generation of effector memory and long-lived central memory T cells, TEM cells, and TCM cells, respectively. Three models by which memory CD8+ T cells can be generated have been proposed (Figure 6.1A–C). Models of CD8+ T-cell differentiation to distinct memory cell subsets. **(A) Linear differentiation model**, an autonomous antigen-triggered differentiation process consisting of conversion from naive to effector to TEM cell occurs, followed by the appearance of TCM cells after antigen clearance through a process of dedifferentiation.¹¹⁷ **(B) The signal strength model** proposes that naive T cells progress through hierarchical thresholds for proliferation and differentiation as the strength and duration of the interaction with APCs is increased.^{118,119} T cells receiving the weakest signals do not survive, whereas high-intensity signaling causes the development of terminally differentiated effector T cells that cannot survive into the memory phase. The TCM cells, being the least differentiated of the antigen-stimulated T cells, retain the developmental options of naive T cells, including their capacity for marked clonal expansion. **(C) The memory stem cell model** proposes that the cells within the TCM cell compartment are self-renewing and serve as a source of effector T cells.

IMPROVING SIGNALING CAPACITIES—SECOND- AND THIRD-GENERATION CARs

T cells play a major role in antitumor immunity. CD8+ CTLs efficiently destroy tumor cells, whereas CD4+ T cells improve the antigen-presenting

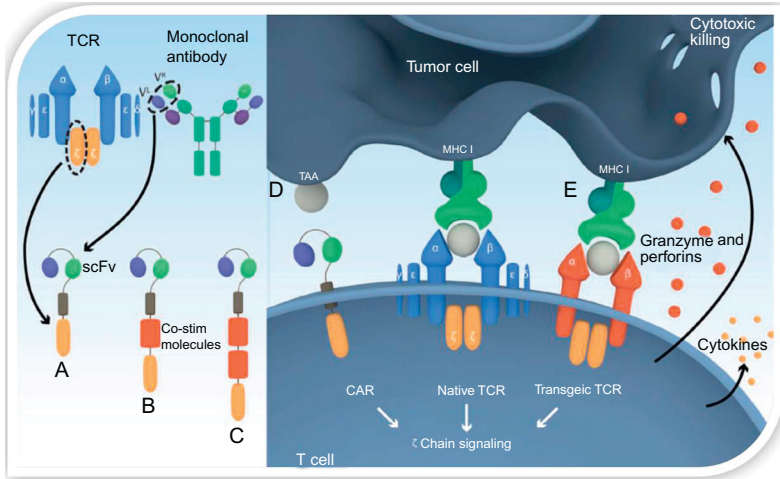


FIGURE 6.3 Genetic engineering and adoptive transfer of patient T cells. Lymphocytes are isolated from the peripheral blood of a cancer patient and transduced with a vector encoding either a new antigen-specific TCR or CAR. The engineered T cells are then expanded *ex vivo* before being adoptively transferred back to the patient. Important factors to consider during optimization of a clinical protocol are indicated.

capacity of DCs and support the stimulation of tumor-reactive CTLs. Following these findings, T-cell-based immunotherapeutic strategies for the treatment of tumor patients were developed. CD4⁺ and CD8⁺ T lymphocytes are powerful components of adaptive immunity, which essentially contribute to the elimination of tumors. Due to their cytotoxic capacity, T cells emerged as attractive candidates for specific immunotherapy of cancer. A promising approach is the genetic modification (see Figure 6.3) of T cells with CARs. The development of an antitumor T-cell mediated immune response is a multistep process in which TAAs expressed by the tumor cells are processed and presented by professional APCs to circulating T cells, which become activated. The quality and potency of this T-cell response will depend on the nature of the antigen presented, the functionality of the APC itself, and the cytokines and costimulatory interactions that occur in the T-cell/APC microenvironment. Once activated, TAA-specific effector T cells can migrate to the tumor site and target malignant cells expressing the cognate antigen epitope displayed in the context of a MHC molecule.¹²⁰

T-cell proliferation and survival require antigenic stimulation by APCs expressing costimulatory molecules. Many tumor cells lack expression of costimulatory molecules, such as CD80 and CD86, and substantially reduce the opportunity of T cells to successfully encounter the antigen in a suitably costimulatory environment.^{121,122} One way to circumvent this problem is to use CARs or transgenic $\alpha\beta$ TCRs to retarget virus-specific CTLs, rather than

“random” T cells. Such virus-specific CTLs should receive optimal activation and costimulation following engagement of their native (virus-specific) $\alpha\beta$ TCRs with viral latent antigens expressed by professional APCs, thus enabling superior antitumor activity mediated by their tumor-reactive transgenic receptor. This hypothesis has been tested in a clinical trial performed at our institution, in which we compared the longevity of polyclonal activated T cells and polyclonal EBV-specific CTLs each expressing a CAR targeting the GD2 antigen, which is expressed on neuroblastoma.^{123,124} The EBV-specific CTLs remained detectable substantially longer (>18 months) than their activated T-cell counterparts (<1 month), suggesting the beneficial effects of costimulation of EBV-specific CTLs provided by APCs expressing EBV-latent antigens.¹²⁴ An alternative means for providing T-cell costimulation is to force the expression of costimulatory ligands, such as CD80 and 4-1BBL, for the costimulatory receptors they already possess.¹²⁵ Post-infusion, these molecules will ensure there is “autocostimulation” and a bystander “transcostimulation,” which substitutes for the lack of appropriate costimulation from the tumor cells. Preclinical studies have shown that incorporation of these molecules in CAR-modified T cells enhances their ability to produce effector cytokines (IFN γ , IL-2, TNF α) and, in an *in vivo* mouse model, promotes tumor rejection in mice with large tumor burden.¹²⁵

A CAR, sometimes referred to as a T-body, chimeric immune receptor, or chimeric artificial receptor, is a transmembrane molecule, which is composed of an extracellular binding domain derived from a single chain antibody fragment (scFv), for recognition of a TAA and intracellular signaling domains for T-cell activation. Hence, upon CAR binding to a TAA on the cell surface of a target cell, the CAR T-cell, will induce apoptosis in the target cell using the same mechanisms as ordinary T cells [Figure 6.2B]. In contrast to a TCR, which recognizes a peptide fragment of an antigen presented by an HLA molecule on the surface of target cells, a CAR molecule recognizes an intact cell surface antigen, thus tumor cell recognition is HLA independent so there is no restriction in terms of patient selection.¹²⁶

CARs are antibody-based extracellular receptor structures anchored into the cell membrane of T cells with a cytoplasmic domain which mediates signal transduction. Eshhar and colleagues introduced the concept of CARs as early as 1989. Several groups have since confirmed the ability to redirect T cells using receptors encompassing different scFvs fused to the CD3 zeta or Fc receptor gamma (FcR γ) signaling domains. CARs are molecules combining antibody-based specificity for tumor-associated surface antigens with TCR-activating intracellular domains with specific antitumor cellular immune activity.^{47,127} These CARs allow a T cell to achieve MHC-independent primary activation through single chain variable fragment (scFv) antigen-specific extracellular regions fused to intracellular domains that provide T-cell activation and costimulatory signals. Most investigators

have achieved efficient CAR gene transfer of human tumor and HIV antigens into human T cells via retrovirus or HIV-derived lentivirus, and some of these cell therapy products have advanced to phase I/II trials.^{124,128–130} Adoptive transfer of CTLs has shown great promise in both viral infections and cancers. Results from recent clinical trials indicate improved clinical results with CARs introduced with retroviral vectors.^{129,130} Second- and third-generation CARs also provide appropriate costimulatory signals via CD28 and/or CD137 (4-1BB) intracellular activation motifs, which augment cytokine secretion and antitumor activity in a variety of solid-tumor and leukemia models.^{131,132} After many years of disappointing results with CAR T-cell therapy, improved culture systems and cell engineering technologies are leading to CAR T cells with more potent antitumor effects.¹³³

First-generation CARs consist of a binding moiety specifically recognizing a tumor cell surface antigen and a lymphocyte activating signaling chain. The CAR-mediated recognition induces cytokine production and tumor-directed cytotoxicity of T cells. Second- and third-generation CARs include signal sequences from various costimulatory molecules resulting in enhanced T-cell persistence and sustained antitumor reaction. Clinical trials revealed that the adoptive transfer of T cells engineered with first-generation CARs represents a feasible concept for the induction of clinical responses in some tumor patients. However, further improvement is required, which may be achieved by second- or third-generation CAR-engrafted T cells.

FIRST-GENERATION CARs

CARs are generated by fusing two distinct functional fragments. The most widely used combination is the antigen binding specific fragment composed of the single chain (scFv) obtained from the variable domains of the heavy and light chains of a monoclonal antibody, and the signaling domain derived from the CD3 chain¹²⁷ (Figure 6.4). These simple molecules are currently defined as first-generation CARs, and when expressed by CD8 and CD4 T-cell subsets, they provide specific binding to antigens expressed on the cell surface of tumor cells and simultaneous activation of the cytotoxic machinery (granzyme-B and perforin) of T cells through the CD3 zeta chain signaling immunoreceptor tyrosine-based activation motif (ITAM). Due to the antibody-mediated antigen recognition, the cytotoxic function of CAR-redirected T cells does not depend on the canonical antigen processing and MHC-restricted presentation typical of native TCRs. The first-generation CAR molecules, with only an scFv against a cell surface antigen expressed on tumor cells and the cytoplasmic CD3 zeta chain signaling domain, were found to have limited clinical activity for the treatment of ovarian cancer,¹²⁹ lymphoma,¹³⁰ renal cancer,⁵⁵ and neuroblastoma.¹³⁴ In a study first-generation CAR T cells demonstrated transient cell division and suboptimal cytokine production and failed to produce prolonged T-cell expansion and

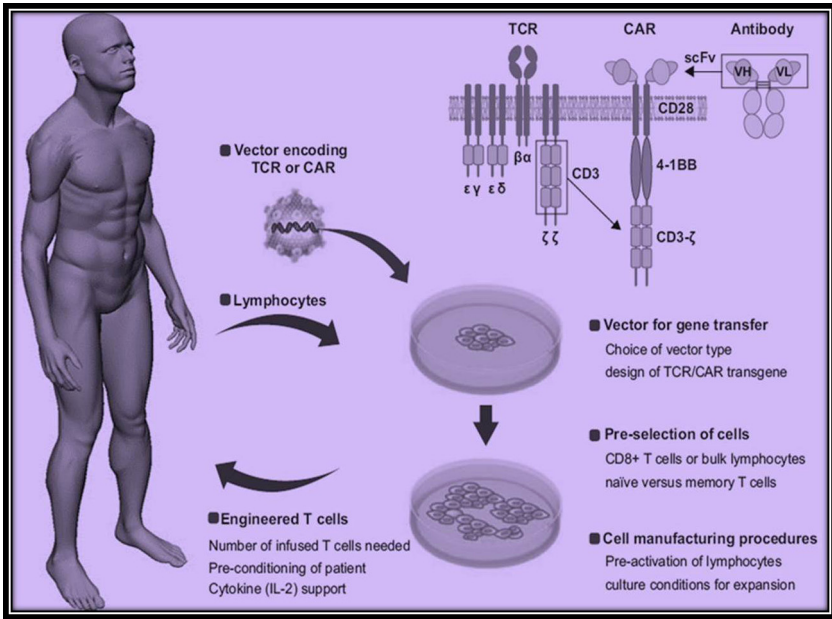


FIGURE 6.4 Redirecting T-cell specificity. T-cell specificity can be redirected by the expression of chimeric antigen receptors (CARs) or transgenic α - and β -TCR chains. (A) CARs are composed of two distinct fragments. The first is obtained by linking the variable heavy and light regions of a monoclonal antibody of known specificity into a single chain (scFv) and the second is the signaling region of the native TCR ζ -chain. (B) Second-generation CARs incorporate costimulatory molecules such as CD28 and 4-1BB. (C) Third-generation CARs include two or more costimulatory molecules. (D) CARs are activated after the scFv portion binds to an extracellular antigen thus the antigen recognition is MHC-independent. (E) Transgenic TCRs recognize specific peptides presented in the context of MHC class I molecules.

sustained antitumor effects. This may not be surprising given that the signal through the TCR-CD3 zeta chain alone is insufficient for priming resting T cells.⁴⁷ Although first-generation CARs could initiate a cytotoxic antitumor response in grafted T cells, it soon became obvious that signals from ITAM-bearing receptors alone can only induce transient cell division and suboptimal cytokine secretion but cannot provide prolonged polyclonal expansion and sustained antitumor reaction *in vivo*. According to the prevailing two-signaling model for lymphocyte activation, full activation and proliferation of T cells require a costimulatory signal through CD28-B7 interaction in addition to signaling through the TCR/CD3 complex.

SECOND-GENERATION CARs

However, several preclinical models showed that first-generation CARs do not fully activate modified T cells upon engagement with the antigen, as

these molecules lack costimulatory properties. To overcome this obstacle, second-generation CARs have been generated in which intracytoplasmic domains, derived from costimulatory molecules such as CD2837, or CD134/OX40 or CD137/4-1BB, are incorporated within CARs to fully activate T cells.^{135–137} Second-generation CARs were constructed to provide signaling both through the CD3 zeta chain and, primarily, the CD28 costimulatory molecule by placing the signaling domains in series as a single gene multidomain product. Constructs with the CD28 signaling domain proximal and the zeta chain distal to the membrane were found to be better expressed than constructs with the opposite orientation and were capable of mediating up to 20 times more interleukin (IL)-2 production upon stimulation with solid-phase antigen compared with first-generation CARs.^{138,139} Subsequently, CAR constructs with costimulatory signaling domains from CD28, ICOS, OX40 (CD134), or 4-1BB (CD137) in series with the CD3 zeta signaling region were evaluated using resting human primary T cells.¹⁴⁰ It was found that second-generation CARs, providing any of these B7 or tumor necrosis factor receptor (TNFR) family costimulatory signals in series with CD3 zeta, confer self-sufficient antigen-driven clonal expansion and enhanced effector function in resting human T cells. Furthermore, addition of the CD28 signaling domain to CARs has been shown to enhance CAR T-cell resistance to regulatory T cells.¹⁴¹

It has been reported that CAR T cells with a 4-1BB signaling domain have improved *in vivo* persistence, tumor localization, and antitumor activity compared with CAR T cells with the CD28 signaling domain.^{142,143} Furthermore, a CAR with the CD27 signaling domain together with the CD3 zeta domain was recently evaluated. The greatest impact of CD27 was noted *in vivo*, where transferred CAR T cells with CD27 demonstrated heightened persistence after infusion, which facilitated improved regression of human cancer in a xenogeneic allograft model.¹⁴⁴ However, side-by-side comparisons of otherwise identical CAR T cells with either CD28, ICOS, OX40, 4-1BB, or CD27 signaling domains, in clinical trials under equivalent conditions, need to be performed before a general conclusion can be drawn as to which costimulatory domain is the most appropriate for CAR constructs.

THIRD-GENERATION CARs

Third-generation CARs have also been constructed containing CD3 zeta, CD28, and the OX40 or the 4-1BB signaling domain.^{135,145} Third-generation CARs, in which two costimulatory molecules (CD28 and 4-1BB or CD28 and OX40) are encoded in tandem, have also been generated, and preclinical experiments suggest that these molecules may be even more potent than second-generation CARs^{136,137,145} (Figure 6.4). These receptors may provide a full complement of activation, proliferation, and survival signals for enhanced antitumor activity. Despite encouraging preclinical results and

some early clinical data, the use of third-generation CARs might have some disadvantages. One concern is that low avidity “off-target” binding may trigger third-generation CARs with potent activation signals that can lead to a lethal “cytokine storm.” One patient treated with a third-generation CAR targeting Her2 died from adverse events due to Her2 expression in the lungs that led to excessive cytokine release and respiratory distress.¹⁴⁶ In addition, third-generation CARs may reduce the signal threshold to a level at which the activation of grafted T cells can occur without triggering antigens. Signal leakage may be a problem for clinical applications of these CARs. Moreover, the exact amino acid sequence and order of the intracellular signaling domains are based on empirical findings, and the optimal CAR format for T-cell activation remains unclear.

Virus-mediated gene transfer seems to be the most efficient gene delivery method, as it allows robust expression of CARs by T cells and minimizes the time of their *ex vivo*. In addition the incorporation of costimulatory endodomains (CD28 or 4-1BB) within CARs is essential in supporting their *in vivo* persistence.^{147,148} Trials that include lympho-depleting chemotherapy regimens prior to the T-cell infusion may have better clinical outcomes as they reduce the tumor burden and provide an environment for the infused T cells that favors homeostatic expansion.¹⁴⁹ For example, impressive results were reported in three patients with aggressive, heavily pre-treated chronic lymphocytic leukemia (CLL) who received chemotherapy 1–4 days prior to the infusion of T cells transduced with a lentiviral vector encoding a CD19-specific CAR, incorporating the 4-1BB costimulatory endodomain.¹⁴⁷ In this trial, the infused cells showed logarithmic expansion *in vivo* and migration to the site of disease with remarkable cytotoxic effects against tumor cells (two complete and one partial remission), and persistence for more than 180 days. In contrast, CAR T-cell expansion observed *in vivo* in clinical trials using CD19-specific CARs that incorporate the CD28 signaling endodomain have produced less striking results,⁴⁵ which suggests that the “late activation” signaling pathways recruited by 4-1BB may be critical for functional, long-term expansion and engraftment of CAR-redirectioned T cells. However, given the diversity of the trials developed so far, systematic assessment of whether 4-1BB is responsible for the better outcome observed in one clinical trial may require a formal comparison between CAR T cells encoding CD28 and CAR T cells encoding 4-1BB infused in the same patient, in a similar way to the direct comparison of first- and second-generation (CD28 costimulation) CD19-specific CAR reported previously.¹⁴⁸

Is the current design of CAR molecules ideal? Potent third-generation CARs have been generated to combine “early” (CD28) and “late” costimulatory signals (4-1BB or OX40) for complete T-cell activation.^{135,137} However, combinations of these molecules raise safety concerns since they may induce T cells to produce excessive amounts of cytokines and promote severe cytokine storms as observed in a patient treated at the National Cancer

Institute.¹⁵⁰ One trial has been published using a CD19-specific third-generation CAR (CD28 and 4-1BB). The infusion of these cells did not produce significant side effects, but the CAR for this study was delivered in T cells by DNA electroporation, and required extensive *ex vivo* culture before infusion, which may exhaust these T cells and thus underestimate their potential toxicities.¹⁵¹ Many CARs may also be immunogenic, as the scFv are usually derived from mouse hybridomas. Although CAR T cells can still be detected for months after infusion, humanized scFv may be preferable in the future to reduce the immunogenicity. Several significant issues still remain to be resolved before designing large phase II studies using CAR technology.

Although first-generation CARs could initiate a cytotoxic antitumor response in grafted T cells, it soon became obvious that signals from ITAM-bearing receptors alone can only induce transient cell division and suboptimal cytokine secretion but cannot provide prolonged polyclonal expansion and sustained antitumor reaction *in vivo*. According to the prevailing two-signaling model for lymphocyte activation, full activation and proliferation of T cells require a costimulatory signal through CD28-B7 interaction in addition to signaling through the TCR/CD3 complex.¹⁵² Providing the costimulatory signal in trans through B7-expressing cells sustained proliferation of single ITAM-bearing CAR-expressing lymphocytes and led to enhanced antitumor response in mouse models.¹⁵³ A more straightforward strategy comprised the CD28 signaling in the CAR concept, either in two separate molecules or combined in a single CD28-ITAM receptor fusion protein.^{154,155} Placement of the CD28 domain proximal to the CD3 ζ chain and immediately distal to the transmembrane domain resulted in good surface expression, whereas in the opposite way the surface expression seems to be inhibited.¹⁵⁶ Therefore, all published second-generation CARs follow the general outline TM domain-CD28-ITAM-bearing signaling chain (Figure 6.4). Subsequently, it has been shown that the synergistic action of the two-signaling domains in one single CAR resulted in sustained proliferation of grafted lymphocytes and increased levels of IL-2, interferon (IFN)- γ , and granulocyte macrophage colony-stimulating factor (GM-CSF) secretion [73] independent of exogenous B7/CD28 costimulation. Due to CD28 costimulation the expression of anti-apoptotic proteins like B-cell lymphoma- (Bcl-) 2 is upregulated and induction of AICD is delayed.^{140,157–159}

Finally, studies in mouse models supported the *in vitro* findings that T cells grafted with recombinant CD28-ITAM receptors have also higher proliferation capacities resulting in an enhanced antitumor activity.^{136,159–161} A transgenic mouse model proved that even naive resting T cells can become fully activated, proliferate in antigen-induced manner, and secrete IL-2 through a chimeric receptor incorporating the costimulatory CD28 sequences in addition to an activating ITAM receptor chain.¹⁶² One major drawback of cellular immunotherapy might be that adoptively transferred cells are rapidly impaired in their effector function by the immunosuppressive milieu, which

tumor cells create in their surroundings.¹⁶³ However, engrafting T cells with second-generation CARs increases substantially the threshold of the modified cells against inhibitory effects mediated by transforming growth factor- β and against the suppressive action of Tregs.^{141,164} Resistance to these cells is an important issue for immunotherapy, as tumor-infiltrating Tregs counteract against adoptively transferred tumor-specific T cells.¹⁶⁵ Another way of improving signaling capacities is the incorporation of Src family kinases or their adaptor molecules, which regulate the very beginning of the signaling cascade resulting from TCR triggering. The 56 kDa Lck kinase is one of the Src-homology kinases, which most likely phosphorylates CD3 ITAM motifs after TCR engagement and is noncovalently linked to either the CD4 or CD8 molecule. Incorporation of the p56Lck or its adaptor CD4 in a CAR with CD3 ζ or CD28-CD3 ζ signaling chains decreased signaling threshold of receptor grafted T cells.¹⁵⁶

In recent years, further costimulatory molecules from the B7 family and the TNFR superfamily came into the focus of research.^{166,167} Not surprisingly, a number of these molecules were also tested in the context of CARs, which led to the development of the third-generation of CARs with three different signaling moieties in the intracellular chain (Figure 6.4). Brentjens et al. generated a series of chimeric receptors bearing signaling sequences from CD28, DAP10, CD134 (OX40), and CD137 (4-1BB) in addition to the CD3 ζ chain.¹⁶¹ However, only the CD28-CD3 ζ chain construct was able to induce significant proliferation of grafted T cells after antigen contact independently of exogenous B7 costimulation. In contrast, when cells expressing B7 molecules were used as targets, all T cells grafted with CARs having CD3 ζ and costimulatory sequences proliferate better than those cells grafted with CD3 ζ CAR.¹⁶¹ Also in terms of IL-2 and IFN- γ secretion, T cells grafted with CD28-CD3 ζ CAR always outcompeted T cells modified with receptors having other signaling combinations.¹⁶¹ Similar results were obtained by another study, which also included the ICOS in the analyzed series of signaling moieties.¹⁴⁰ Secretion of inflammatory cytokines like TNF- α , IFN- γ , and GM-CSF was similar for T cells engrafted with CD28-CD3 ζ and ICOS-CD3 ζ CARs; however, IL-2 secretion and induction of proliferation was better with the CD28-CD3 ζ construct.¹⁴⁰

Thus, both studies have proven that for antigen-induced proliferation and high-level cytokine secretion the signal from CD28 is required and sufficient, but inclusion of further costimulatory molecules enhances proliferation capacities and apoptosis resistance in the grafted T cells.¹⁶⁸ It is now established that physiologically optimal activation requires CD28 engagement followed by costimulation through other T-cell signaling molecules.¹⁶⁹ Consequently, CARs which can simultaneously transmit multiple signals from CD28, CD3 ζ , and an additional costimulatory molecule in the engrafted cell have been constructed. One of the most important “secondary” costimulatory molecules is OX40 (CD134), for which studies have shown

that its signaling can further augment CD28-activated T-cell responses, enhancing proliferation, cytokine secretion, and survival.¹⁷⁰ Integration of OX40 in a third-generation CAR in combination with CD28 and CD3 ζ chain leads to sustained *in vitro* proliferation and increased IL-2 secretion by grafted human primary T cells.¹³⁵ A number of studies pointed out the important role of the CD137/CD137L interaction for T-cell survival and AICD resistance.^{125,171,172} Integration of CD137 signaling in CARs increased expression of anti-apoptotic protein Bcl-xL, induced sustained proliferation and survival of the grafted T cells, and was associated with higher effector cytokine production and antigen-specific tumor cell lysis.^{132,173} In a mouse model, an established xenograft was controlled by retargeted T cells containing a CD28-CD137-CD3 ζ CAR.¹⁴⁵ If targeted cancer cells are B7.1/B7.2 positive, chimeric CD137-CD3 ζ CARs can be even more sufficient than a CD28-CD3 ζ CAR.¹³² Despite these encouraging results, the use of CARs with a tripartite signaling domain has to be carefully investigated, and it remains to be seen if CARs with tripartite signaling moieties reduce the signal threshold to a level where activation of grafted T cells can occur without antigen triggering.

In contrast to TCRs, CARs recognize TAAs independent of HLA. The typical CAR consists of four separate domains that are assembled from individual units: (1) the single chain variable fragment (scFv) domain, (2) the extracellular scaffold/linker, (3) the transmembrane domain, and (4) the signaling endodomain.¹⁷⁴ The specificity of the CAR is imparted by the scFv domain, which is directed against a TAA. Other molecules can be substituted for the scFv domain, including cytokines, pattern-recognition receptors, and cell surface molecules, to dock the CAR to the respective ligands of these molecules.^{175–177} The scFv domain is attached to the transmembrane region via an extracellular scaffold. Optimal recognition of the TAA is affected by the extracellular length of the CAR. TAAs presented proximal to the plasma membrane may be more efficiently recognized by a CAR containing a longer scaffold, and conversely, TAAs presented further from the cell membrane may be more efficiently recognized by a CAR containing a shorter scaffold.^{178,179} Moreover, the source of the scaffold may greatly affect the effectiveness and *in vivo* function of the CAR. CARs targeting the B-cell antigen CD19 that have been used for clinical trials contain scaffolds derived from CD28, CD8, IgG1, or IgG4. The IgG motifs found in CARs may bind to endogenous Fc receptors, leading to clearance of CAR⁺ T cells expressing the IgG motifs and the nonspecific activation of NK cells. *In vivo* binding between CAR⁺ T cells containing the IgG scaffolds and endogenous cells expressing Fc receptors can apparently be reduced by mutagenesis of the IgG scaffolds.¹⁸⁰ The transmembrane domain links the extracellular scFv domain and scaffold to the signaling endodomain. Oligomerization of the transmembrane domain may favor CAR signaling, but few studies have systematically analyzed different transmembrane domains.¹⁸¹

Signaling via the CAR molecule occurs via the addition of various endodomains to the cytoplasmic portion of the chimeric molecule. First-generation CARs signaled via immunoreceptor ITAMs from the CD3- ζ chain or FcR- γ .¹²⁷ A second costimulatory signal is included in second-generation CARs: the signaling endodomains of proteins such as CD28, 4-1BB, OX40, and ICOS.¹⁸² CARs containing two or more costimulatory molecules in addition to CD3- ζ are commonly referred to third-generation CARs. A competitive repopulation experiment demonstrated a survival advantage for second-generation CAR⁺ T cells infused into lymphoma patients compared to first-generation designs.¹⁴⁸ The ability to infuse more than one population of CAR⁺ T cells into a given recipient is appealing to help understand the immunobiology which favors the sustained persistence and, thus, the enhanced therapeutic activity of a given CAR species. However, the expense associated with generating two infusion products from two recombinant viral vectors diminishes the enthusiasm for this otherwise appealing experimental approach.

To help reduce the barriers to combining gene therapy with T-cell therapy, our laboratory at MD Anderson Cancer Center in Houston, Texas, has employed electroporation using a commercially available Nucleofector device (Lonza Inc., Allendale, New Jersey, USA) to stably express CARs in primary human T cells for clinical trials.¹⁸³ Circulating T cells are electrotransferred with both the CAR-containing transposon and a hyperactive SB transposase, which catalyzes the cut-and-paste insertion of the transposon into the host genome. The apparent transient expression of the SB transposase in manufactured T cells helps to minimize the genotoxicity that arises via the continuous reintegration of the transposon. Transient expression of this fish-derived transposase also reduces the risk of the recipient developing an immune response to the transposase after the cells are infused. This improvement is significant, as an anti-transgene immune response could prompt the clearance and rejection of the genetically modified cells.¹⁸⁴ After gene transfer, the T cells are selectively propagated on activating and propagating cells (AaPCs) in the presence of stimulatory soluble cytokines for 4 weeks. The clinical-grade AaPCs are genetically modified K-562 cells that are available as a master cell bank. The AaPCs support the selective numeric expansion of CAR⁺ T cells by expressing a TAA and the costimulatory molecules CD86, 4-1BB, and membrane-bound IL-15.¹⁸⁵

Currently, to maintain patient safety, the manufacture of clinical-grade SB-modified CAR⁺ T cells occurs over a period of 3–5 weeks to enable the loss of the DNA plasmid containing the SB transposase.^{185–187} Indeed, polymerase chain reaction (PCR) is used to confirm the absence of the SB transposase prior to infusion. Redirected T-cell specificity mediated by the CAR is evaluated via a chromium release assay. Sterility is verified via visual inspection, a negative result for mycoplasma based on a PCR assay, and a negative result for endotoxin based on the limulus amoebocyte lysate test. The absence of genotoxicity is determined by the lack of growth of the

genetically modified T-cell population upon removal of the AaPCs and cytokines, as well as by demonstrating a normal karyotype. The electroporated and propagated T cells are evaluated via flow cytometry for (1) viability, (2) the expression of the CAR, (3) the memory phenotype, (4) the absence of AaPCs, (5) exhaustion markers, and (6) telomere length. The presence of a polyclonal TCR repertoire is validated by flow cytometry or using bar-coded probes to confirm the lack of skewing of the numerically expanded T cells.¹⁸⁸

CAR T CELLS IN CLINICAL TRIALS

CAR⁺ T cells typically engineered by introducing DNA via a virus or a SB system have been successfully infused into patients with either solid or hematologic malignancy (see Table 6.1). Transfer of autologous T cells transduced with retrovirus to enforce the expression of a CD19-specific CAR containing the CD28 and CD3- ζ endodomains after a preparative chemotherapy regimen led to the prolonged eradication of CD19⁺ cells in a follicular lymphoma patient.¹⁸⁹ Later studies conducted using CD19-specific CAR⁺ T cells in eight lymphoma and CLL patients found favorable responses in six of these patients, including 1 CR lasting more than 15 months and 5 partial responses (PRs) lasting at least 7 months. The expected adverse events associated with targeting CD19 include long-term B-cell aplasia in about half of the patients. In addition to these long-term toxicities, acute adverse events manifested as symptoms of inflammation, such as fever, fatigue, and hypotension. The loss of humoral immunity can be life-threatening, as 1 patient has died of culture-verified influenza 18 days after receiving an infusion of CAR⁺ T cells.¹⁴⁹

Current T-cell immunotherapy relies on harvesting T cells either from the patient, who may produce few or defective T cells due to the malignancy itself and prior treatment of the malignancy, or from an HLA-matched allogeneic donor, which includes the risk of GVHD. Due to their inherent antitumor activity and lack of HLA restriction, $\gamma\delta$ T cells are an intriguing candidate therapy that can be pre-prepared from unrelated third-party donors and infused on demand. Artificial nucleases are another means to improve T cells for adoptive immunotherapy. Zinc-finger nucleases, transcription activator-like effector nucleases, and CRISPR/Cas9 introduce double-stranded DNA breaks at specific sites, which leads to repair by nonhomologous end joining, which may result in gene inactivation.^{190–194} These technologies display potential for modulating the immune response to generate a favorable outcome in cancer immunotherapy. Artificial nucleases, such as CTLA-4 and programmed cell death-1 (PD-1), can be used to remove negative regulators of the anticancer response.^{109,110,195} Furthermore, artificial nucleases could be used to remove the TCR from the $\alpha\beta$ T cells used for immunotherapy to eliminate the HLA alloreactivity of the infused T cells to reduce the risk of GVHD and to generate an off-the-shelf T-cell product.¹⁹⁶

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