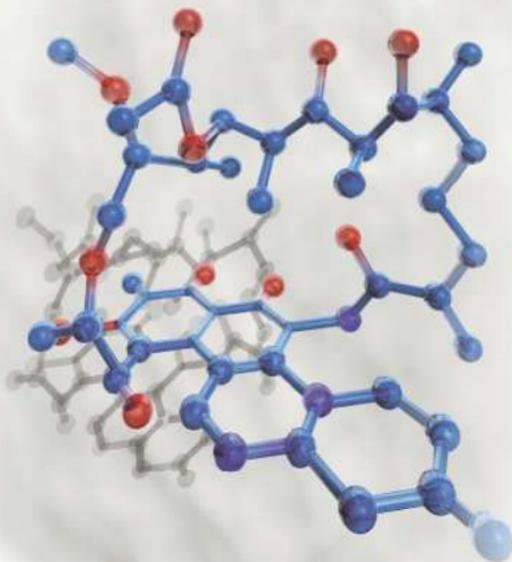


Rifaximin: A Poorly Absorbed Antibiotic

Pharmacology and Clinical Use

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The Pathogenesis of Gastrointestinal Bacterial Overgrowth

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Key Words

Bacterial overgrowth · Pathogenesis · Gastrointestinal motility · Gastric acid · Malabsorption syndromes

Abstract

The normal indigenous intestinal microflora consists of about 10^{15} bacteria that under physiological conditions reside mainly in the lower gastrointestinal tract. Bacterial overgrowth implies abnormal bacterial colonization of the upper gut, resulting from failure of specific defense mechanisms restricting colonization under physiological conditions. At present two types of bacterial overgrowth with defined pathogenesis can be distinguished: (1) gastric overgrowth with upper respiratory tract microflora resulting from selective failure of the gastric acid barrier, and (2) gastrointestinal overgrowth with Gram-negative bacilli (enteric bacteria) resulting from failure of intestinal clearance. *Helicobacter pylori*-induced gastritis of the oxyntic mucosa is the main cause of acquired failure of the gastric acid barrier, which is common among the healthy elderly. Intestinal clearance may fail as the result of impaired intestinal peristalsis or anatomical abnormalities that alter luminal flow. Impaired peristalsis is associated with conditions interfering with intestinal neuromuscular function including myopathic, neuropathic, autoimmune, infectious, inflammatory, metabolic, endocrine, and neoplastic diseases. Anatomical ab-

normalities are mainly the result of gastrointestinal surgery, intestinal diverticula or fistula. Combined failure of intestinal clearance and the gastric acid barrier results in more severe colonization with Gram-negative bacilli. Gram-negative bacilli are uncommon in the upper gut of otherwise healthy individuals with gastric hypochlorhydria, being acquired (*H. pylori*) or drug-induced. Significant bacterial overgrowth with Gram-negative bacilli is a rationale in the search for an explanation to optimize clinical management. The clinical significance of colonization with upper respiratory tract microflora remains unclear. Translocation of live bacteria, their metabolic products, or antigens from a small bowel colonized by Gram-negative bacilli play a role in the pathogenesis of spontaneous bacterial peritonitis in hepatic disease and in certain types of sepsis, indicating that further studies can point to new patient populations with potential benefit from medical treatment.

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Introduction

The oral cavity and the lower gastrointestinal tract are densely colonized by bacteria with counts exceeding 10^9 colony-forming units (CFU)/ml, whereas the density in the stomach and proximal small bowel is normally below 10^5 CFU/ml (fig. 1). Bacterial density increases through

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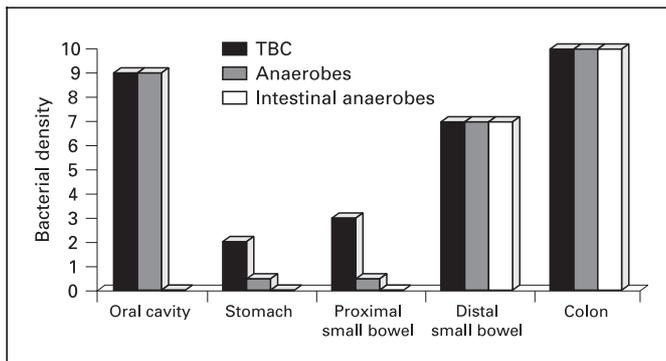


Fig. 1. The density of bacteria along the gastrointestinal tract of man is shown schematically based on data from references 1–5 in the text. Density is given by log₁₀ CFU/ml of luminal contents in the fasting state. TBC = Total bacterial count.

the ileum to approximately 2 log below cecal counts in the distal ileum. Bacterial overgrowth implies abnormal bacterial colonization of the upper gut.

There is also a segmental distribution of the types of bacteria. Strict anaerobic species are normally confined to the oral cavity and the colon, habitats they densely colonize and predominate [1–5] (fig. 1). Bacteria indigenous to the upper respiratory tract (URT flora) and anaerobic bacteria of oral origin are swallowed with saliva and recovered from the upper gut at densities below 10⁵ CFU/ml. Under physiological conditions, they are considered transitory rather than indigenous to the upper gut. Facultative anaerobic bacteria are usually confined to the distal small bowel and colon, but transient species entering the gut with nutrients are occasionally recovered from the healthy upper gut at low counts.

When the mechanisms restricting bacterial colonization in the upper gut fail, due to disease or dysfunction, bacterial overgrowth develops. The segmental distribution may be gastric, intestinal or both depending on the type of failure. The consequences for the host vary from none to life-threatening complications, caused by severe water and electrolyte deficiencies and septic manifestations.

Definition of Bacterial Overgrowth

The predominant quotation in the literature is purely quantitative with 10⁵ CFU/ml of small intestinal aspirate as a limit [2, 6–8]. In symptomatic bacterial overgrowth, Gram-negative bacilli are present in the small intestine, making the flora ‘colonic-like’ [2, 7]. The term ‘bacterial

overgrowth syndrome’ has been used to define bacterial overgrowth leading to clinical symptoms [7], without reference to the pathogenesis of the disorder.

In the present review, an increase in bacterial density above 10⁵ CFU/ml of small intestinal aspirate is considered the general definition of bacterial overgrowth, in accordance with the current standard [2, 6–8]. Based on this definition, recent data make it possible to distinguish between two types of bacterial overgrowth with distinct pathogenesis, microflora and clinical presentation: bacterial overgrowth with URT flora and with Gram-negative bacilli, respectively (table 1). With cultures from both the stomach and small intestine, the segmental distribution can also be defined. Unless the segment is specified, bacterial overgrowth is synonymous with small intestinal bacterial overgrowth.

Testing for Bacterial Overgrowth

Culture of intestinal contents is the gold standard for detecting bacterial overgrowth [2, 7, 9]. This technique allows both segmental localization and the identification required to distinguish between URT and Gram-negative bacilli, respectively. The labor intensity and cost, however, make its clinical use difficult.

Of the indirect tests the ¹³C or ¹⁴C-*D*-xylose or lactulose breath test and the glucose, lactose or lactulose hydrogen breath tests are available alternatives. These tests are in general developed to recognize Gram-negative bacilli rather than URT overgrowth. There are, however, pitfalls involved.

Rapid intestinal transit may result in a false-positive breath test, in particular when hyperosmolar nonabsorbable substrates are used. A false-negative outcome in patients with culture-proven Gram-negative bacilli in the upper gut further query the sensitivity and usefulness of breath tests for clinical practice [10–13]. Positive microbial culture from small intestine is thus advantageous when major alterations of clinical management are considered.

The Main Defense Mechanisms

The pathogenesis of bacterial overgrowth is reviewed by considering separately the consequences of failure of the two main defense mechanisms in the upper gut responsible for the two types of bacterial overgrowth (table 1): the gastric acid barrier and intestinal clearance.

Table 1. Developing the concept of bacterial overgrowth^a

Type	URT flora	GNB
Pathogenesis	Failure of the gastric acid barrier	Failure of intestinal clearance
Etiology	<i>H. pylori</i> -induced atrophy of gastric mucosa, drug-induced etc.	Failure of small bowel motility or intestinal anatomical abnormality
Bacteria	Mainly Gram-positive bacteria	Enterobacteriaceae In severe forms strict anaerobic species of colonic type
Tracer species	α -Hemolytic streptococci	<i>E. coli</i> \pm <i>Bacteroides fragilis</i> group
Location and extent	Gastric stomach Similar flora present in duodenum and proximal jejunum	Small intestine, segmental or global Backwards colonization of the stomach in severe forms

Features of the two main types of bacterial overgrowth, defined by the underlying pathogenesis (see text for details of the failure required to alter the microflora of the upper gut, and the diseases and clinical conditions that can lead to failure of the gastric acid barrier and intestinal clearance, respectively). GNB = Gram-negative bacilli.

^a $>10^5$ CFU/ml of fasting luminal contents.

The significance of oral bacterial carriage, degree of illness, malnutrition and immunological disorders will also be addressed.

The Gastric Acid Barrier

Defining the Gastric Acid Barrier

Gastric acid can be quantified by the capacity of secretion (peak or maximal acid output) or by the concentration of H_3O^+ ions generating the acidity of gastric juice (pH). It is the acidity that regulates microbial growth [1, 14–16], which is further emphasized by the observation that bacterial counts in the stomach correlate with basal but not with peak acid output [17]. Failure of the basal acid secretion that determines fasting gastric pH is therefore of particular importance. Accordingly, patients with a preserved ability to secrete acid in response to maximal stimulation may still have fasting hypochlorhydria [18].

At pH 4 most bacteria are killed within 30 min, and at physiological luminal pH, 99% of bacteria are killed within 5 min [14]. Certain bacteria, like lactobacilli, are more acid-resistant, and some microbes survive the hostile gastric environment by colonizing luminal niches at the mucosal surface, protected by gastric bicarbonate secretion. This is the case for *Helicobacter pylori*, related spiral-

shaped bacteria, and particular fungi [5]. Although the gastric acid barrier is acidic enough to kill all bacteria ingested, dynamic changes of gastric pH and emptying related to the intake of nutrients explain survival through the gastrointestinal tract. Passage of live bacteria is physiological, and a prerequisite for maintaining a normal indigenous gut microflora [19].

Reduced gastric acidity with pH 3–5 during and just after meal intake [1] and the rapid initial phase of gastric emptying [20] both contribute to the gastric passage of live bacteria. The meal-induced increase of bacteria in the stomach and upper small bowel disappears about 1 h after meal intake, when gastric emptying is slower and gastric pH has returned towards fasting levels [1]. This occurs hours before the recurrence of the migrating motor complex in the upper gut [21], a motility pattern associated with luminal clearance of the small bowel [12, 22, 23] (see below).

The short-lasting temporal variations in gastric pH in concert with the migrating motor complex during fasting [24] are less likely to result in significant changes in gastric microflora, although the secretory component [24, 25] of the migrating motor complex contributes to intestinal clearance.

There are also segmental variations of intragastric acidity. Because the antrum is usually empty in the fasting state, local pH is substantially influenced by duodenogas-

tric reflux and also by other factors [26] making this location less suitable for reliable measurements of the gastric acid barrier. The fundic reservoir, however, is capable of acidifying considerable amounts of refluxate. If, for example, 10 ml of duodenal chyme at pH 7 refluxes into a fundic reservoir of 50 ml gastric juice at pH 2.00, the increase to pH 2.08 is negligible in terms of microbial growth. The pH of fundic aspirate is thus a robust indicator of fasting gastric acidity with respect to the control of luminal microbial growth.

There is also a gradient from the low luminal pH through the mucus layer, under which gastric bicarbonate secretion maintains neutral conditions. Mechanically, this is explained by the acid secretion occurring like small finger-like ejections penetrating the thick gel-like mucus layer into the gastric lumen [27].

Bacterial colonization of the mucosal surface by, for example, *H. pylori*, other spiral-formed bacteria, and fungi reflect the microbial ability to pass the mucus layer and to adhere, rather than a failure of host defense. Accordingly, in developing countries with poor hygienic conditions, the great majority of people are colonized by *H. pylori* from early childhood [28, 29], whereas the prevalence in industrial countries is steadily falling with improved standards of living [29]. This type of colonization thus differs from bacterial overgrowth of the lumen that reflects microbial adaptation to the failure of host defense.

In the present review the gastric lumen is confined to the habitat above the mucus layer, for which the pH of fasting gastric juice is the major defense mechanism against bacterial colonization. This defense mechanism is henceforth denoted the gastric acid barrier.

Testing the Gastric Acid Barrier

The gastric acid barrier is tested by measuring the acidity of gastric aspirate or by an intragastric pH probe [30]. Serial aspirations during fasting over 24 h [31] gave results comparable to those obtained by intragastric pH probes during 24 h with four meals [30]. The average 24-hour pH is thus mainly determined by the fasting pH, confirming the importance of basal acid secretion in this regard. The ease and superior data acquisition when using an intragastric pH probe connected to a portable data logger make this test attractive [30], but it is expensive, time-consuming, uncomfortable for the patient, and requires expertise.

Measurement of pH in gastric juice aspirated during endoscopy can be used as a rough albeit robust indicator of the gastric acid barrier. In 29 consecutive outpatients

undergoing routine endoscopy, aspirates were collected from the fundic reservoir by entry of the stomach and again before withdrawal of the endoscope [32]. The increase from the first to the second aspiration was only 0.22 pH units (range -0.99 to 1.39). For the 24 patients with fasting gastric pH <4, the mean was pH 1.87, which fits in well with the average intragastric pH 1.98 observed during 24-hour recordings in healthy individuals eating four meals [33]. The mean + 2 SD was pH 2.95 [32], corresponding to the recommended upper limit of pH 3 for normal pH of fasting gastric aspirates [1, 14, 34, 35].

A single aspirate from the fundus during fasting is also a valid indicator. Fasting gastric aspirates were obtained from 51 patients participating in an acid secretion study (unpubl. data kindly provided by L. Blomquist at the Karolinska Hospital, Stockholm, Sweden). In 26 of 51 patients pH > 3 was found in the first aspirate after intubation. The average of the four succeeding basal aspirates taken at 15-min intervals showed 100% agreement: the same 26 patients had at least one of four succeeding samples with elevated pH, using pH 3 as a cutoff. The clinical relevance of this limit is confirmed by the correlation between bacterial counts and time of pH > 3 from 24-hour pH recordings [36].

Measuring pH in fundic juice aspirated when entering the stomach during endoscopy is thus a simple, robust, and valid means of testing the gastric acid barrier, and pH >3 indicates failure.

Failure of the Gastric Acid Barrier

Causes of Failure of the Gastric Acid Barrier

Drug-Induced Inhibition of Acid Secretion **H₂-Receptor Blockers**

Although H₂-receptor blockade markedly inhibits maximal acid output, the reduction of gastric acidity is modest because basal output remains and tolerance develops during chronic use [37]. With a standard dosage of cimetidine of 800 mg [38] or nizatidine of 300 mg [36] gastric pH will increase modestly to about pH 2 in gastric aspirate [36, 38], which is too acidic to allow for clinically significant bacterial colonization of the stomach. Increased bacterial density in gastric juice has been reported during H₂-receptor blockade in some studies [17, 35, 39, 40], although others have found no significant change [34, 36]. The limited effect of H₂-receptor blockers that explains this discrepancy was clearly shown in a recent comparison with proton pump inhibitors [38].

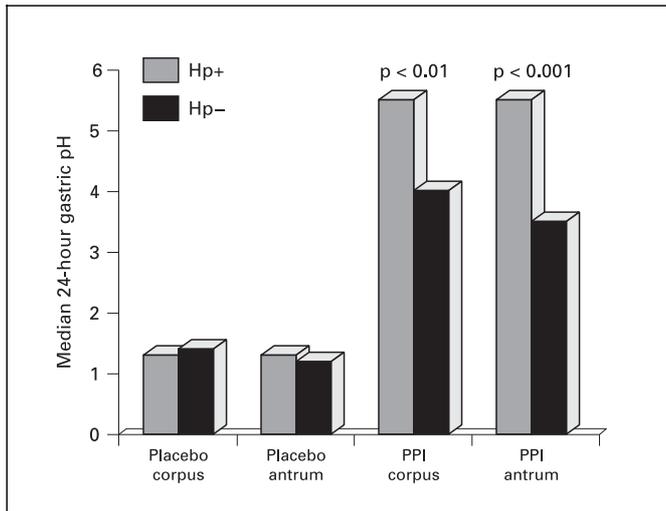


Fig. 2. Median gastric pH is elevated about 2 log by 20 mg of omeprazole in *H. pylori*-negative healthy subjects, and by 4 log in *H. pylori*-positive ones [based on data from 42]. Hp = *H. pylori*; PPI = proton pump inhibitor.

Proton Pump Inhibitors

Proton pump inhibitors are potent inhibitors of gastric acid secretion, resulting in an increase of gastric pH that interferes significantly with the gastric acid barrier. It is now well established that *H. pylori* is of major importance for the magnitude of this response, an effect that relates to the extension of the gastritis into the gastric corpus [41]. In *H. pylori*-negative individuals, 20 mg of omeprazole daily increases gastric pH about 2 pH units to pH 3–4 [42] (fig. 2). This results in a 50–100-fold increase of bacterial density in the stomach [43]. In *H. pylori*-positive individuals, however, the same dose will raise gastric pH by about 4 pH units to pH 5–6, which will almost completely abolish the gastric acid barrier. Accordingly, the bacterial density increases more than 1,000-fold [42, 43]. Comparable results were obtained by Sharma et al. [44] when 30 mg of omeprazole was given to healthy volunteers without knowing their *Helicobacter* status. Based on the available literature, figure 3 shows how the density of bacteria in the stomach increases with gastric pH, to reach a plateau of about 10^8 CFU/ml beyond pH 6.

H. pylori Colonization

When the gastritis induced by *H. pylori* is confined to the antrum, the increase of gastrin and the reduction of somatostatin released by the G and D cells in the antrum, respectively, will increase the drive for acid secretion from the preserved oxyntic mucosa [45]. This increased

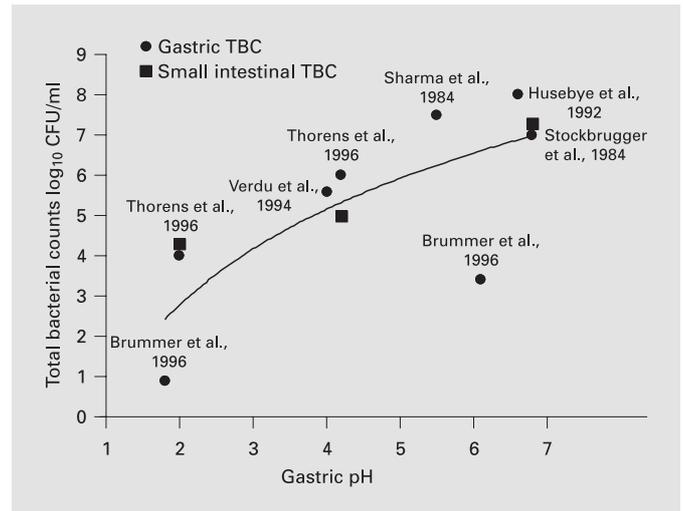


Fig. 3. The relationship between gastric pH and total bacterial counts in the stomach is shown by studies of patient populations and healthy volunteers with different gastric pH levels. Verdu et al. [43], 1994: *H. pylori*-negative healthy subjects on omeprazole 20 mg. Sharma et al. [44], 1984: Healthy individuals on omeprazole 30 mg. Thorens et al. [38], 1996: Patients on omeprazole 20 mg or cimetidine 800 mg (lower pH). Brummer et al. [36], 1996: Patients on omeprazole 20 mg or nizatidine 300 mg (lower pH). Stockbrugger et al. [17], 1984: Patients with pernicious anemia. Husebye et al. [32], 1992: Healthy old individuals with hypochlorhydria related to gastritis. Thorens et al. and Stockbrugger et al. also give data for duodenal cultures; corresponding values are found at the same pH as for gastric TBC. Logarithmic trend line for gastric bacterial counts is given.

acid secretion contributes to the development of duodenal ulcer and maintains the gastric acid barrier.

In another subpopulation, the *Helicobacter* gastritis extends into the corpus resulting in atrophy of the oxyntic mucosa and reduced acid secretion. It is not yet clear to which extent these manifestations reflect different stages or different courses of *Helicobacter*-induced gastritis [41, 45]. *H. pylori* thus emerges as the main cause of acquired gastric hypochlorhydria [46–48].

The Role of Aging

Achlorhydria, implying reduced peak acid output, was found in only 17.5% of 348 patients above 70 years of age [49]. Evidence for elevated gastric pH, however, was found in 82% of 657 patients above 65 years using the azuresin test: achlorhydria in 68% and hypochlorhydria in 14% [50]. Differences in techniques and definitions explain this divergence. Elevated fasting gastric pH is thus prevalent in the elderly. Accordingly, in healthy old people >75 years of age, 80% had hypochlorhydria defined as

fasting gastric pH > 3 with average gastric pH of 6.6 (Husebye et al. in fig. 3) [32].

The observation that gastric acid secretion declines with age [49, 50] is biased because of the influence of *H. pylori*. Accordingly, the reduction of acid secretion in the elderly is a cohort effect caused by *H. pylori*-associated atrophic gastritis of the oxyntic mucosa [46–48]. In *H. pylori*-negative individuals, gastric acid secretion persists during aging [46, 48, 51, 52] in the absence of autonomic diseases and other conditions interfering with acid secretion [53].

Autoimmune Disease

Pernicious anemia is the classical autoimmune disease associated with immunologically mediated injury of the oxyntic mucosa resulting in achlorhydria [52]. Parietal cell antibodies are also present in other autoimmune diseases [52, 53] and immunopathies [54] that can be associated with hypo- or achlorhydria.

Malnutrition and Degree of Illness

Malnutrition per se is associated with both gastric hypochlorhydria and bacterial overgrowth with both URT flora and Gram-negative bacilli [55]. The degree of illness, which determines oral colonization with Gram-negative bacilli [56], contributes to this change of microflora in severe malnutrition. Accordingly, when severely malnourished children were nourished, the gastric Gram-negative colonization disappeared after initial treatment, before gastric acidity was restored [55]. Malnutrition, therefore, induces Gram-negative colonization of the upper gut through mechanisms other than the failure of the gastric acid barrier. This observation concurs with other studies showing that gastric hypochlorhydria per se does not lead to Gram-negative colonization of the stomach [32, 34, 40, 44].

Surgery

Gastric surgery reducing acid secretion is associated with gastric bacterial overgrowth with URT flora corresponding to the degree of pH elevation, and in a proportion of patients also Gram-negative bacilli, depending on the type of surgery [16, 57, 58]. Greenlee et al. [59] carefully examined the influence of different types of gastric acid-reducing surgery on the microflora of the upper gut in dogs. Gastrectomy and truncal vagotomy resulted in 100–1,000 times higher concentrations of bacteria in the upper jejunum. After proximal gastric vagotomy, however, resulting in a similar elevation of gastric pH, no change of jejunal microflora was found [59]. The same

pattern is seen in clinical studies [16, 58, 60]. Changes of the anatomy and the parasympathetic innervation of the antroduodenal region after surgery may interfere with motility and clearance, and thus predispose to colonization with Gram-negative bacilli in the small bowel.

Consequences for the Gastric Microflora

Gastric Acidity and the Density of the Gastric Microflora

There is a close correlation between gastric acidity and the density of bacteria in the stomach. At fasting gastric pH < 3, gastric aspirate will be sterile or contain less than 10³–10⁴ CFU/ml [1, 14, 61–63]. With an elevation of gastric pH, bacterial counts increase to a plateau of about 10⁶–10⁸ CFU/ml at pH 6–7.5 [1, 62–64] (see fig. 3). This was recently reviewed in further detail by Yeomans et al. [65].

Gastric Acidity and the Composition of the Gastric Microflora

In healthy individuals URT flora multiplies in gastric aspirate during treatment with antisecretory compounds and in particular proton pump inhibitors [34, 40, 44]. This concerns viridans streptococci, coagulase-negative staphylococci, *Haemophilus* sp., diphtheroids, *Moraxella* sp., lactobacilli, and other streptococci, most of which are Gram-positive bacteria. With dedicated measures anaerobic species of oral origin are also recovered [66].

Gram-negative bacilli are in general not recovered or only occasionally and at low counts in studies of healthy individuals on acid inhibitors [34, 40, 43, 44] (table 2). This pattern has also been shown in healthy old people with hypochlorhydria secondary to chronic gastritis, of whom the great majority only harbored URT flora despite gastric pH > 6 [32].

In patient populations with gastric hypochlorhydria, as discussed above, Gram-negative bacilli are recovered in a minor proportion. This concerns 10–30% of patients on acid inhibitors, in particular proton pump inhibitors [36, 39, 67], 10–50% after gastric ulcer surgery depending on the type of surgery [16, 57, 58], and about 30% of patients with pernicious anemia (table 2). The Gram-negative bacilli most frequently reported are *Escherichia coli*, *Klebsiella* sp., and *Proteus* sp., belonging to the Enterobacteriaceae. This type of colonization is hard to explain only with increased gastric pH.

Patients with peptic ulcer disease have mucosal injury and may develop fibrosis in the antroduodenal region and

Table 2. Degree and cause of failure of the gastric acid barrier and gastric microflora in density and composition

Gastric pH	Cause	Gastric bacterial density	Gastric microflora
2–3 (4)	H ₂ blockers	No or mild increase <10 ^{3–5} CFU/ml	Sterile or URT (5–10% GNB in patients)
3–4	PPI in Hp– healthy subjects	Moderate increase 10 ^{4–6} CFU/ml	URT
	PPI in Hp– patients ^a		URT (10–25% GNB)
4–6	Moderate Hp gastritis ^b	Marked increase 10 ^{5–7} CFU/ml	URT
	Incomplete proximal vagotomy		URT (10% GNB)
	PPI in Hp+ healthy subjects		URT
	PPI in Hp+ patients		URT (10–30 % GNB)
6–7.5	Peptic ulcer surgery		URT (10–50% GNB) ^c
	Advanced Hp gastritis ^d	Maximum increase 10 ^{8–9} CFU/ml	URT
	Peptic ulcer surgery		URT (10–50% GNB)
	Autoimmune atrophic gastritis		URT (20–30% GNB)

GNB = Gram-negative bacilli; PPI = proton pump inhibitor; Hp = *H. pylori*.

^a Patients with peptic ulcer disease and reflux esophagitis.

^b Early stage of atrophic corpus gastritis of limited extension (less common).

^c The prevalence of Gram-negative bacilli colonization depends on the type of surgery (see text).

^d Atrophic corpus gastritis (prevalent in the elderly due to the high prevalence and duration of *H. pylori* colonization in this age cohort).

changes in mucosal defense and motility [68] that may contribute to a shift from URT flora to Gram-negative bacilli when on proton pump inhibitors. Moreover, 41% of patients with reflux disease have delayed gastric emptying [69], a delay that is considerable in some patients, suggesting an underlying motility disorder [70].

To predict the type of gastric microflora in patients with elevated gastric pH, the presence of local structural and functional changes that may result from diseases requiring acid inhibition [36, 39, 67], nutritional status [55], degree of illness [56], and concurrent diseases or drugs that may interfere with gastrointestinal motility [71] must be considered. It should be recalled that when such factors are present, acid inhibition may promote colonization with Gram-negative bacilli in the upper gut. In a detailed prospective study of patients with late radiation enteropathy, concurrent failure of the gastric acid barrier was found to aggravate significantly the bacterial overgrowth with Gram-negative bacilli resulting from failure of intestinal peristalsis [12]. Accordingly, jejunal bacterial overgrowth was promoted by concurrent hypochlorhydria in patients with progressive systemic sclerosis [72].

Summary of Failure of the Gastric Acid Barrier: Gastric Bacterial Overgrowth

Selective failure of the gastric acid barrier, as seen in otherwise healthy individuals on proton pump inhibitors or with *H. pylori*-induced corpus gastritis, results in gastric colonization of swallowed oropharyngeal bacteria. In otherwise healthy subjects this will be mainly Gram-positive bacteria belonging to the URT flora and strict anaerobic bacteria of oral origin.

Gastric acid is the main defense mechanism against gastric bacterial overgrowth, and the density of bacteria correlates to intragastric acidity, as shown in figure 3 and table 2, depending mainly on basal acid output. A significant increase in bacterial density is seen when fasting gastric acidity exceeds pH 3, the upper normal limit for pH in fasting gastric juice aspirated during endoscopy. Bacterial density peaks at 10⁸–10⁹ CFU/ml of gastric juice at pH 6–7.5.

H. pylori is now recognized as the main cause of selective gastric hypochlorhydria, which today is highly prevalent (more than 50%) in the normal elderly population of western countries and predominant in developing countries with prevalence often exceeding 90%. The in-

fluence of proton pump inhibitors on gastric pH and microflora is enhanced in the presence of *H. pylori* (fig. 2). H₂-receptor blockers have less effect on gastric acidity, remaining below pH 3, and thus on gastric microflora.

Concurrent colonization by Gram-negative bacilli occurs in some patients with failure of the gastric acid barrier, suggesting additional deficiencies of host defense: abnormal oral flora, malnutrition, general illness, or diseases or medication interfering with intestinal peristalsis and clearance. This type of microflora is also seen in 10–30% of patients on acid inhibitors, for which mucosal injury and functional changes related to peptic ulcer and reflux disease may be responsible.

Consequences for the Intestinal Microflora

The consequences of a failure of the gastric acid barrier for the intestinal microflora emerge from studies of healthy individuals and patient populations with other important defense mechanisms against bacterial colonization intact.

Intestinal Microflora in Healthy Individuals with Gastric Hypochlorhydria

Drug-Induced Inhibition of Acid Secretion

Shindo et al. [66] treated 19 healthy volunteers with omeprazole 20 mg, cultured gastric and jejunal aspirate, and determined gastric pH and bile acid metabolism. Although motility studies were not performed, it can be assumed that intestinal migrating motor complexes were normal [21] (fig. 4). Bacterial colonization was defined by species density exceeding 10⁵ CFU/0.5 ml, and only reported for those exceeding this limit.

Omeprazole resulted in an increase in URT flora, without a significant shift towards Gram-negative bacilli colonization. Two subjects had *E. coli* colonization in jejunal aspirates before treatment. Eleven showed colonization during treatment, all by a single species: *Bacteroides vulgatus* (n = 4) and *Bacteroides uniformis* (n = 1), *Eubacterium parvum* (n = 2) and *Eubacterium lentum* (n = 1), *Lactobacillus bifidus* (n = 2), and *Corynebacterium granulosum* (n = 1). These are anaerobic and aerobic bacteria that may colonize the oropharyngeal habitat. The *Bacteroides* spp. are, however, of the intestinal type, although they are not obligatorily intestinal as is *Bacteroides fragilis* [73]. It is notable that Shindo et al. [74] also reported significant jejunal colonization by intestinal types of anaerobes in healthy individuals during cimetidine treatment, which they explained by a shift to neutral pH in gastric juice

[74]. Significant jejunal colonization by *E. coli* was found in 7 of 53 individuals before and in 4 individuals only during treatment with H₂ blocker. The same species as reported during omeprazole treatment [66] were recovered [74], mostly bacteria of oropharyngeal origin.

Significant colonization by *E. coli* in 13% [74] and 21% [66] of the healthy subjects prior to treatment may suggest oral carriage for reasons unrelated to gastrointestinal structure and function. H₂-receptor blockers elevate gastric pH only modestly, regardless of *H. pylori*, and fasting gastric pH < 3 should be expected [36, 38], which does not lead to major changes of gastric or duodenal microflora in healthy individuals [34, 36, 38–40]. Moreover, colonization by strict anaerobic bacteria of intestinal type in the proximal small bowel has thus far been associated with stasis of the small bowel [12, 75] and co-colonization by coliforms (Enterobacteriaceae) at significant counts [12, 75]. Many standard identification schemes for *Bacteroides* spp. are designated for potentially pathogenic intestinal types and may misidentify isolates of oral origin [73].

Furthermore, similar glucose hydrogen breath tests in the elderly with and without omeprazole [76] and normal ¹⁴C-*d*-xylose breath test in healthy old people with acquired gastric hypochlorhydria (pH > 6) [32] counterindicate that H₂ blockers induce colonization with strict anaerobes of intestinal types (colonic flora) in the upper gut.

An important novel finding in these studies was the detection of bile acid metabolism during acid suppression in healthy volunteers [66, 74], presumably caused by gastric bacteria, in particular when gastric pH exceeds 4 [66, 74, 77]. In vitro experiments showed that most of the bacteria recovered, mainly of oropharyngeal origin, were able to metabolize ox bile [66, 74]. In contrast, the ¹⁴C glycocholic breath test was unchanged 6 weeks after omeprazole 40 mg and 26 weeks after 20 mg [78], and more studies of acid inhibition and microbial metabolism in the upper gut are thus needed.

The consequence of bacterial bile acid metabolism [66, 74, 77] is hardly clinically significant malabsorption [6] in otherwise healthy individuals [32, 79], but in predisposed individuals this may be different. Accordingly, omeprazole interferes with the absorption of vitamin B₁₂ [80–83] and protein assimilation [84]. The mechanism for altered vitamin B₁₂ absorption is prevention of its cleavage from dietary protein [83], for which the importance of the concurrent bacterial overgrowth has not yet been ruled out.

Shindo et al. [66, 74, 77] explain the presence of *Bacteroides* spp., presumably of the intestinal type, by migra-

tion from the ileum due to the change of pH in the small bowel. With a pH between 5 and 6 in the physiological state allowing bacterial colonization, the minor shift induced by cimetidine is unlikely to change significantly the microbial ecology of the small bowel. Accordingly, gastric pH did not correlate to Gram-negative colonization in jejunal aspirate [85].

Retrograde colonization is less likely in the absence of a widespread motility disorder or fistula [12, 75, 86]. When judged by defecatory intervals and stool form score, omeprazole was found to speed intestinal transit [87], which is comparable to experimental data showing that the predominant effect of commensal intestinal bacteria on physiological small bowel motility is the stimulation of myoelectric activity and transit [88, 89]. Elevated gastric pH will increase the load of bacteria that enter small intestine (fig. 3). Accordingly, in a recent thesis the combination of 40 mg of omeprazole twice daily and 300 mg of H₂ blocker at bedtime induced intestinal contractile activity during the fasting state by increasing phase II activity at the expense of phase I of migrating motor complex [90] (fig. 4).

In conclusion, total bacterial counts in the duodenum and the most proximal part of the jejunum of healthy subjects increase by about 2 log during standard proton pump inhibition with omeprazole 20 mg daily [87]. The bacterial species encountered are mainly of the URT flora. Gram-negative bacilli are occasionally recovered at low counts, the origin of which may be ingested food or oral carriage. There is disagreement concerning gastrointestinal bacterial metabolism during acid inhibition. Most studies have been negative [32, 76, 78, 87], but recent data [66, 72, 74, 77, 91] may indicate otherwise, at least for bile acids. Gastric overgrowth by URT flora, the ultimate result of elevated gastric pH, may thus not be as harmless as currently thought [52, 81, 83, 84]. Further studies are required [92] to clarify this important issue regarding the safety of pharmacological acid suppression in clinical practice.

Age-Associated *H. pylori*-Induced Hypochlorhydria

Healthy old people with fasting gastric hypochlorhydria and preserved intestinal motility [79] had normal ¹⁴C-*d*-xylose breath test, corresponding with gastric culture showing predominantly URT flora in >90% of the individuals [32]. Overgrowth with Gram-negative bacilli in the upper gut is thus not a consequence of failure of the gastric acid barrier per se [32]. This corresponds to the absence of Gram-negative bacilli in the upper gut of patients with normal migrating motor complex in proxi-

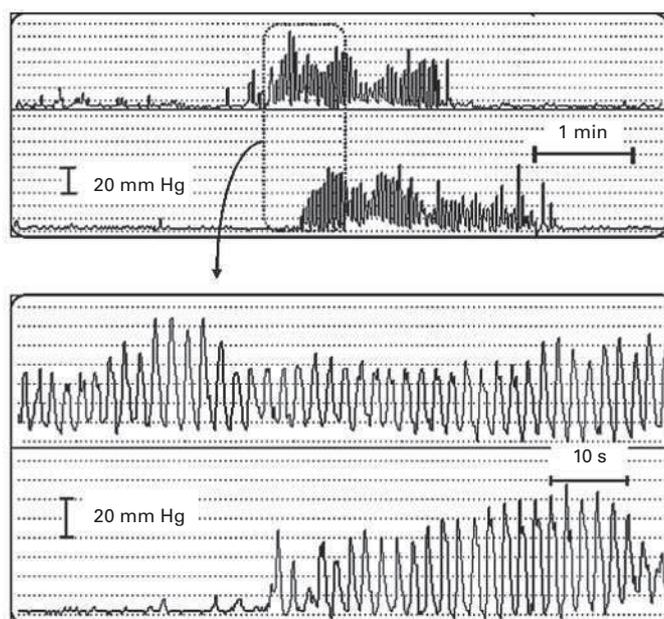


Fig. 4. The normal nocturnal migrating motor complex (MMC) recorded in the duodenum (upper tracing) and proximal jejunum (lower tracing) of a 91-year-old healthy woman. A short period is shown in high resolution in the lower panel. Phase III is preceded by phase II with some contractile activity, usually limited during sleep, and succeeded by contractile quiescence, phase I. The sequence of phase III-I-II-III constitutes one MMC cycle, and recurs during fasting (modified with permission from Husebye and Engedal [79]).

mal small bowel [12], and malnourished children during recovery when hypochlorhydria is still present [55].

Intestinal Microflora in Patients with Gastric Hypochlorhydria

Cregan et al. [93] showed that neither gastric hypochlorhydria nor the presence of a profuse gastric microflora necessarily lead to the development of a resident flora in the mid-small bowel: 'an antibacterial mechanism, distinct from that in the stomach, must operate in small intestine'. Accordingly, Frederiksen et al. [85] could not find any relationship between gastric secretory capacity and Gram-negative bacilli in jejunal aspirate in a large series of patients.

Of 41 patients with chronic abdominal complaints after previous successful abdominal radiotherapy for pelvic malignancy, 29 patients had preserved intestinal peristalsis and clearance evidenced by normal migrating motor complex activity during prolonged ambulatory intestinal manometry, and normal anatomy by small bowel follow-through [12] (fig. 5). Five of these 29 (18%) had gastric hypochlorhydria. Dense gastric bacterial colonization

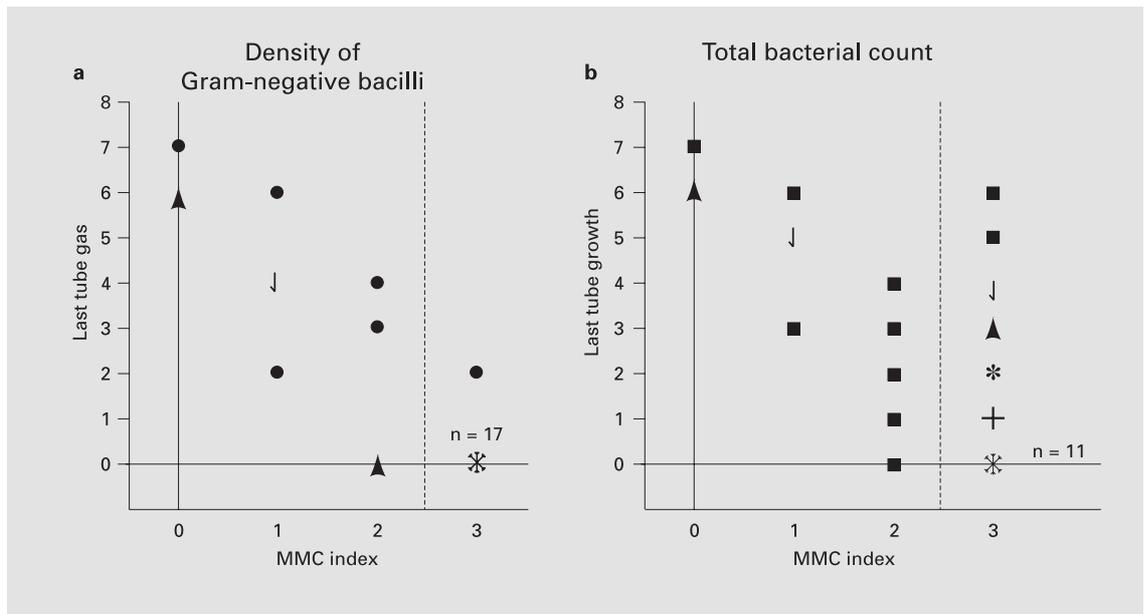


Fig. 5. Relationship between fasting intestinal motility [x-axis: migrating motor complex (MMC) index] and bacterial colonization of small bowel in 41 patients with late radiation enteropathy (LRE) is shown by two plots. Relationship to Gram-negative bacilli (a) and to total bacterial count (b) in the duodenum is shown. Note that no significant Gram-negative colonization was found in patients with normal MMC (index = 3). The vertical dotted lines show the normal limit for MMC index. Increased bacterial counts due to URT flora were found in some patients with normal MMC (b). Tied observations are indicated as follows: n = 1: ●, ■; n = 2: ↓; n = 3: ▲; n = 4: +; n = 6: *. For n > 6 number is given (with permission from Husebye et al. [12]). For total bacterial count 'last tube growth' indicates log₁₀ for CFU/ml. For 'last tube gas' see [12].

was found in all, consisting of only URT flora in 4 of 5 (80%). *E. coli* was recovered in only 1 patient and strict anaerobic bacteria of colonic origin were not detected. Despite dense colonization of the stomach, the duodenum was only moderately colonized [12] by principally the same bacterial species. This corresponds to the findings of Sherwood et al. [75], sampling from five sites along the small bowel. They showed that intestinal anaerobic overgrowth occurred in relation to local or general stasis in the small bowel. In their study group with previous partial gastrectomy, intestinal anaerobes were not recovered from any site of the small bowel, despite marked gastric hypochlorhydria and complementary gastric bacterial overgrowth [75].

A correspondence between gastric and duodenal microflora when the gastric acid barrier fails has also been shown in patients with pernicious anemia [17].

Summary of Failure of the Gastric Acid Barrier: Intestinal Bacterial Overgrowth

When the gastric acid barrier fails the bacterial counts in the most proximal part of small bowel increase. Stan-

dard proton pump inhibition by omeprazole 20 mg daily will increase bacterial density by about 2 log, because bacteria are continuously emptied from the colonized gastric reservoir. In the duodenum the species will be quite similar to those cultured from the stomach. Unless there are concurrent factors or conditions predisposing to colonization with intestinal Gram-negative bacilli, URT flora will predominate. Recent data suggest that this URT flora may cause bacterial metabolism of bile acids and alter the assimilation and proteins and vitamin B₁₂, the significance of which remains to be clarified. In patients with a failure of other defense mechanisms predisposing to colonization by Gram-negative bacilli, proton pump inhibition will augment this type of bacterial overgrowth, which may be clinically harmful.

When intestinal peristalsis and clearance are intact, the bacteria are rapidly transported aborally, and in the mid jejunum bacterial counts are in general low (normal) despite dense gastric colonization. Considerable evidence indicates that bacteria recovered from small bowel under such conditions are transient rather than resident.

Intestinal Clearance

Defining Intestinal Clearance

Intestinal clearance is henceforth defined as the ability of the small bowel to clear its lumen of bacteria. The known conditions of major clinical importance for intact intestinal clearance are (1) normal gastrointestinal anatomy, including the absence of intestinal diverticula and fistula, and (2) normal intestinal motility.

Secretion and the immune system also contribute to luminal clearance of bacteria, but dysfunction and abnormalities of clinical relevance for the development of bacterial overgrowth have so far been associated with the factors outlined above. Moreover, normal intestinal motility, tested by manometry, also indicates that the enteric neuroendocrine control of motility, secretion, absorption and circulation is intact [24, 25, 94]. To the extent that gastrointestinal secretion has been studied, failure does not seem to result in bacterial overgrowth [95–98]. Studies on the immune system are briefly discussed later. The failure to recognize the clinical importance of these factors in the present context may, however, also reflect current methodological and scientific limitations. Although a failure of the gastric acid barrier increases the bacterial load to the small intestine from the gastric reservoir, evidence does not indicate that this defense mechanism contributes significantly to intestinal clearance of bacteria.

Intestinal Motor Activity and Clearance of Bacteria

Rolly and Liebermeister [95] showed that bacteria introduced into the small bowel disappeared rapidly, without bile, pancreatic, and intestinal juices having antibacterial properties alone or mixed. Later studies, of which those by Dack and Petran [96], Dixon [99] and Dixon and Paulley [100] are of particular importance, provided considerable further evidence that intestinal peristalsis is the main line of defense against bacterial colonization of the small bowel. This was also concluded by Donaldson [101–103] when he reviewed host defense mechanisms in 1964. At that time, however, the insights into small bowel motility were confined to the reflex-mediated peristaltic behavior.

Bayliss and Starling [104] described the peristaltic reflex of small intestine in 1899. This enterically controlled reflex elicits a contraction oral to and a relaxation distal to a segmental distension, resulting in the movement of contents in the aboral direction [104]. The peristaltic reflex is funda-

mental for understanding the behavior of the small bowel during nutrient stimulation, and there is a revived interest in the control mechanisms involved [105].

Reflex behavior, however, does not fully explain intestinal motor activity. During the fasting state, the small bowel moves at intervals, apparently spontaneously, in the absence of nutrients. The enteric nervous system intermittently inhibits the intestinal smooth muscle cells, which would otherwise spontaneously contract at a regular rate, like the cardiac muscle, due to the intrinsic pacemaker properties [106, 107]. The fasting state thus shows periods of both silence and contractile activity, depending on the degree of enteric inhibitory control with a maximum contractile rate of 11/min in the duodenum, decreasing to 7–8/min in the ileum. Regular contractions at this frequency occur for time periods of about 5 min at intervals ranging from 20 min to hours in healthy individuals [21, 23]. This band of regular propagating contractions, called phase III of the migrating motor complex, migrates in the aboral direction (fig. 4). C.F. Code named the migrating motor complex the gastrointestinal housekeeper, due to its propulsive properties capable of clearing the lumen of contents during the fasting state [22].

Intestinal mechanical clearance thus consists of both reflex-mediated contractions (peristalsis) elicited by the stimulatory effect of luminal contents and of periods of spontaneous contractile activity (e.g. the migrating motor complex). During fasting about 50% of intestinal transit has been attributed to phase III of the migrating motor complex, the remaining mostly to the propulsive contractions and motor patterns during phase II [108]. Luminal flow can also occur in the absence of propagating contractions of the circular muscle layer, so far considered the motor event mainly responsible for flow in the small intestine.

The motility of the small bowel has been studied in great detail in experimental, physiological and clinical research [21, 71, 106, 107, 109], and the patterns are well defined in man [21, 23, 110]. Although a standard test of intestinal motor activity with regard to the efficiency of mechanical luminal clearance is not yet established for clinical use, means to evaluate this function have been proposed.

Testing Intestinal Clearance

Microbial Culture of Intestinal Contents

The absence of Gram-negative bacilli in the small bowel is a reliable indicator of preserved intestinal clearance [12, 75, 111]. Although significant colonization of Gram-

negative bacilli results from failure of intestinal clearance [12, 75, 111], oral carriage due to malnutrition [55], illness and reduced health [56], and other structural and functional changes [36, 39, 67] can also be the cause. The presence of Gram-negative bacilli in small bowel is therefore an indication, but no proof of failing intestinal clearance. The denser the colonization, however, the more likely there is a failure of clearance. When strict anaerobic bacteria of the intestinal type are present, advanced failure with stagnation is indicated [12, 75], unless there is a blind loop or a fistula [7].

Reference to the normal oropharyngeal microflora is required to distinguish the URT flora [112], for which α -hemolytic streptococci are predominant and the candidate marker. Among the Gram-negative bacilli of the intestinal type, Enterobacteriaceae are easy to recover by culture because they are facultative and their prevalence in bacterial overgrowth is high. *E. coli* is the predominant species, and therefore the candidate marker. The limit of 10^5 CFU/ml serves to distinguish transient Gram-negative bacilli that may be recovered in health [7]. When strict anaerobic species of the colonic type are recovered this limit may be too high, but this depends largely on the culturing technique. Standardization of culture for bacterial overgrowth is required to establish more specific quantitative limits at the species level. Microbiological expertise and control data are, therefore, required for an appropriate interpretation of cultures for the diagnosis of bacterial overgrowth. This also concerns the occasionally difficult distinction between strict anaerobic bacteria of oral and colonic types [73].

Testing the Intestinal Mechanical Clearance (Intestinal Motor Activity)

If anatomical abnormalities have been ruled out, testing of the small bowel motor activity is useful to elucidate the pathogenesis of bacterial overgrowth with Gram-negative bacilli (table 1). This choice is encouraged by the correlation between clinical disorders associated with bacterial overgrowth and disorders associated with dysmotility of the small bowel [113].

Manometry remains the gold standard test, because phasic contractions are generally lumen occlusive in small bowel and thus reliably detected by intraluminal pressure measurements. Transit tests are more convenient; nevertheless, they are time-consuming and do not provide the same detailed information about contractile activity [114, 115]. These methods are briefly discussed.

Small Bowel Manometry

Data on small bowel motility disorders have been obtained by using both stationary techniques [21, 71, 114, 116, 117] with external transducers and water-perfused catheters [117] and by the use of ambulatory techniques [21, 118]. The establishment [110] and further implementation [119–121] of ambulatory techniques allow prolonged recording throughout the day and night at home [118]. Testing of both the response to nutrient challenge and the fasting motility is required in the present context, which implies prolonged recordings. This favors the use of ambulatory techniques.

Stanghellini et al. [122] have carefully defined the most common abnormalities of phase III activity and other abnormal motility patterns that occur in patients with chronic intestinal pseudoobstruction, who often suffer from bacterial overgrowth [113]. This concerns phase III with abnormal migration (stationary or retrograde) and with abnormal isotonic component, abnormal burst activity, and a failure of the postprandial pattern.

Phase III of the migrating motor complex serves as a marker of intestinal motility for several reasons. When phase III fails, concurrent abnormalities of postprandial motility patterns and other propulsive patterns during fasting are common [12, 21, 71, 117, 122–124]. Normal occurrence of the migrating motor complex and absence of strictly abnormal motor patterns during prolonged recording, including both the fed and fasting states, are valid and reliable indicators of preserved intestinal mechanical clearance [21]. In a large series comparing prolonged ambulatory small bowel manometry and culture, failure of the migrating motor complex predicted colonization by Gram-negative bacilli in the small bowel [12]. A semiquantitative migrating motor complex index was, therefore, proposed [12]. Schemes to analyze and evaluate a small bowel manometric record have been proposed [21, 125] (fig. 5), and international consensus is pending.

Small Bowel Transit

A small bowel transit study can be used to evaluate intestinal propulsion and clearance, and the presence of Enterobacteriaceae (Gram-negative bacilli) in the small bowel indicates delayed transit [111]. The wide normal variability, however, makes transit tests rather insensitive, and thus less useful clinically [126, 127]. It is also a problem that accelerated and delayed transit may coexist in neuropathies and confuse the interpretation. Finally, as nutrients are mostly absorbed in the proximal small bowel, and the rate and pattern of transit vary along the intestine, segmental failure of transit is easily missed by global

transit measurements. Although easier to perform, the clinical utility is often limited unless the dysfunction is severe [127]. The most commonly used transit tests available are briefly discussed with reference to the current study.

Scintigraphy

Single- and dual-isotope techniques have been applied [126] with labeling of the liquid and solid phase by ^{99m}Tc , ^{111}In or ^{113}In , or ^{67}Ga . The difference between the half-emptying time of the stomach and the half-filling time of the cecum has usually been estimated. The more accurate approach, however, is to use the technique of deconvolving the profiles for gastric emptying and colonic filling to obtain a spectrum of transit times, and then to calculate the mean value [126]. By this technique there was no discrimination between transit of liquids and solids [126]. Transit time ranged from 1.5 to 6 h in healthy subjects after a mixed meal [126], reflecting the limitations for clinical use. Only marked acceleration or delay can be detected, which apply mainly to patients with intestinal pseudoobstruction. Modified and simplified scintigraphic tests have been developed [114, 128], the use of which should be encouraged if a transit test is chosen to evaluate intestinal peristalsis in the presence of bacterial overgrowth.

Breath Tests

Studies of small bowel transit time have demonstrated a great variability both within and between individuals. When the hydrogen breath test was performed under fasting conditions, using 10 ml of lactulose, the coefficient of variation amounted to 18%. Di Lorenzo et al. [129] showed that variations under fasting conditions are partly accounted for by the phase of the migrating motor complex at the intake of test solution. Moreover, when a lactose-containing meal was used, the coefficient of variation was reduced to 4% [130].

The main limitation of breath tests in this setting is the bias induced by the intestinal overgrowth flora, generating a breath signal that can be difficult to distinguish from the arrival of the substrate in the cecum. This is further hampered by the intermittent passage of the head of a meal into the colon, which may, also under normal conditions, generate multiple signal peaks before a more sustained signal is obtained.

Breath tests are, therefore, less useful for testing of intestinal transit in the presence of bacterial overgrowth.

Failure of Intestinal Clearance

Causes of Failing Intestinal Clearance

Abnormal Intestinal Anatomy

Anatomical changes can alter luminal flow into a surgically prepared blind loop, a diverticulum, or through a fistula. These anatomical abnormalities of relevance for the development of bacterial overgrowth have been carefully defined in previous literature [2, 7, 98].

When significant Gram-negative overgrowth is detected, anatomical abnormalities should be considered prior to studies of intestinal mechanical clearance. The anatomy is revealed by X-ray using small bowel follow-through, optionally supplied by new modalities of ultrasound, computer tomography and magnetic resonance imaging.

Failure of Intestinal Mechanical Clearance (Intestinal Motility)

Although hereditary neuropathies and myopathies affecting small intestinal motility are rare, the entire spectrum of diseases that can interfere with motility is wide, including for example diabetes mellitus, Crohn's disease, scleroderma, and postoperative and radiation sequelae [21, 71, 116, 123, 131].

Failure of intestinal motility can be severe leading to frank intestinal pseudoobstruction [122, 132] or mild to moderate depending on the underlying disease, its severity, and the degree of intestinal involvement.

In some patients believed to suffer from the irritable bowel syndrome, an underlying enteric neuromuscular disorder has later been identified [133]. The bridge to infectious diseases is also of interest, with several enterotropic viruses in focus, and reports of lymphocytic infiltration of enteric neural structures in patients with unexplained intestinal dysmotility require further studies.

Neuromuscular Diseases

Enteric Neuropathies. Different kinds of familial visceral neuropathies have been described: the dominant type 1 [134], the recessive type 2 [135] and a recessive form with calcified basal ganglia [134]. Furthermore, aganglionosis of the small bowel (Hirschsprung's disease) [136], hyperganglionosis (neurofibromatosis) [137], neuronal intestinal dysplasia [138] and Parkinson's disease [139] are neuropathies to consider. The recognition of the pacemaker cells of the small bowel, the interstitial cells of Cajal, has prompted studies to detect abnormalities of these cells, another possible cause of pseudoobstruction [140].

Extrinsic Neuropathies. Autonomic dysfunction [141], pandysautonomia [142, 143], Shy-Drager syndrome [144] and sympathetic dysfunction are conditions associated with intestinal dysmotility.

Vagal neuropathy in diabetes mellitus [145, 146] and truncal vagotomy [147] may markedly change intestinal motility, as do heart-lung transplantation [148]. Spinal cord lesions also alter gut function, but the outlet obstruction due to failure of the striated muscles involved in defecation is more important than the enteric smooth muscle effects [149].

Enteric Myopathies. The familial types include the dominant type 1 [150], the recessive type 2 with ophthalmoplegia [151] and the recessive type 3 [116]. The sporadic types include muscular dystrophies [152] including myotonic dystrophy [153] and Duchenne's dystrophy. Dysmotility has been associated with all these diseases.

Diseases and Injury of the Gut Wall

Radiation Injury. Late radiation enteropathy is associated with alterations of small intestinal motility [154], intestinal pseudoobstruction [154, 155] and Gram-negative colonization of the small bowel in patients with impaired small bowel motility [12]. In patients with severe injury, alterations in the motility and microflora are of main importance for the clinical symptoms [154].

Inflammation. Chronic inflammatory bowel disease affecting the small bowel can lead to disturbances of intestinal motility [146]. Potential mechanisms are previous surgery, development of fibrosis and strictures, malabsorption, and 'cross-talk' between inflammatory and enteric nerves [156, 157]. Patients with Crohn's disease are often included in aggregate studies of bacterial overgrowth [23, 75, 158], reflecting this link.

Connective Tissue Diseases. Scleroderma is the connective tissue disease most frequently associated with intestinal dysmotility and bacterial overgrowth [159, 160]. Although the motility of the esophagus is most frequently affected, and a prerequisite for the label CREST syndrome, small bowel involvement is seen in a proportion of these patients. When present, intestinal clearance is usually impaired because of shallow contractions resulting in ineffective peristalsis and clearance. This can lead to overgrowth with Gram-negative bacilli, in part responsible for the malabsorption [161].

The neuromuscular compartment of the bowel wall is also affected in certain types of the Ehler-Danlos syndrome [162], maybe in amyloidosis [163], and in the presence of diffuse lymphocytic infiltration [164].

Infectious Diseases

Chagas disease affects enteric ganglionic cells. This leads to altered motility with a reduced rate of migration for the migrating motor complex [165], a change associated with colonization with Gram-negative bacilli [12].

Dysmotility has been reported in Lyme disease [166] and in postviral syndromes associated with cytomegalovirus and herpes simplex virus [167]. Altered intestinal motility can also be part of infectious mononucleosis [168].

Metabolic and Endocrine Disorders

Thyroid Disease. Hypothyroidism (myxedema) [169] and hyperthyroidism [170] alter small bowel motility. Although today these diseases are usually recognized before such symptoms develop, thyroid function must be examined in unexplained intestinal pseudoobstruction.

Diabetes mellitus. Diabetes mellitus interferes with gastrointestinal motility through different mechanisms including blood sugar oscillations, extrinsic vagal neuropathy, vascular changes and enteric neural injury. Intestinal dysmotility [145, 146] is seen in a proportion of the diabetics, and intestinal pseudoobstruction associated with bacterial overgrowth can develop. When abdominal complaints are chronic and disabling, studies of intestinal microflora and clearance should be considered, in particular if nutritional problems occur.

Paraneoplastic Syndromes

Intestinal pseudoobstruction is also part of paraneoplastic syndromes. The anti-*hu* antibodies are useful to indicate this condition, as shown in bronchial small cell carcinoma [171]. In pheochromocytoma [172] and carcinoid [173] neuromediators affecting small bowel motility are produced by the tumor cells. Intestinal pseudoobstruction has also been reported in neuroblastoma [174].

Hepatic Disease

Patients with advanced liver cirrhosis often suffer from bacterial overgrowth [175]. Chang et al. [176] reported a reduced frequency and rate of migration for migrating motor complexes in patients with liver cirrhosis, alterations that predispose to colonization of the small bowel by Gram-negative bacilli [12].

Drug-Induced Dysmotility

The perhaps most important and easily ignored cause of secondary dysmotility is the drug-induced toxic type. Pharmaceuticals are important to consider, in particular those with anticholinergic and/or opioid properties [177]. In individuals with reduced reserve capacity of the gut, either due to concomitant disease or age, such drugs may

elicit pseudoobstruction. Although aging does not lead to clinically significant dysmotility, the reduction in the rate of migration for migrating motor complexes and increased prevalence of clustered contractions indicate reduced reserve capacity [21, 79].

Surgery

Although the gastrointestinal tract has great adaptive and reserve capacities, surgery can directly or indirectly through generation of fibrosis, adhesions and strictures interfere with small intestinal motility [178, 179]. Vagal injury will be of importance in particular for the motor response to feeding, whereas direct injury or modifications of intestinal loops are usually present if pseudoobstruction results. The Billroth type II resection of the stomach and the Roux-en-Y anastomosis result in chronic dysmotility, likely to be of importance when postoperative abdominal complaints occur [178].

Consequences for the Gastric Microflora

Gastric emptying is delayed in patients with intestinal pseudoobstruction [180]. Intestinogastric reflexes including the duodenal and ileal brakes are candidate mechanisms for this effect. Recent data indicate that delayed gastric emptying per se does not interfere significantly with gastric microflora, when the gastric acid barrier is maintained [181].

A certain proportion of short-reaching retroperistaltic waves at the gastroduodenal junction is physiological [182], but in severe functional dyspepsia both segmental spread and the contribution of retroperistalsis in the duodenum was increased [183]. Paradoxical gastric colonization by Gram-negative bacilli despite the presence of a normal gastric acid barrier has been reported in some patients with severe late radiation enteropathy associated with marked intestinal dysmotility [12]. Giant retrogradely migrating contractions, observed in these patients, may also reflux intestinal contents into the gastric lumen.

In conclusion, gastric microflora is altered in patients with severe forms of intestinal pseudoobstruction due to frequent duodenogastric reflux episodes caused by abnormal retrogradely propagating contractions.

Consequences for the Intestinal Microflora

Pharmacological suppression of intestinal peristalsis in experimental animals leads to bacterial colonization of

small bowel by all types of bacteria present in the gut, including Gram-negative bacilli [100, 184, 185]. Similar studies cannot be performed in man, but patients taking opioids regularly show changes of intestinal motor activity with a slowing of peristalsis and transit resulting in constipation.

Intestinal Microflora in Patients with Failure of Intestinal Peristalsis

Vantrappen et al. [23] for the first time showed the relevance of phase III of the migrating motor complex in the current context, when reporting its absence in 5 of 12 patients with bacterial overgrowth detected by the bile acid breath test and response to antibiotics.

The consequences of altered intestinal motility patterns for the microflora of the small bowel have later been addressed in detail [12]. Forty-one patients with varying degrees of dysmotility due to previous successful abdominal radiotherapy for malignancy were studied. Impaired phase III of the migrating motor complex was invariably associated with intestinal colonization by Gram-negative bacilli, whereas normal phase III reliably predicted the absence of such microflora [12, 21] (fig. 5). Significant URT flora was detected in the small bowel of patients with normal motility patterns and failure of the gastric acid barrier [12]. The underlying pathophysiology could thus be established, considering the type of overgrowth flora [12].

Further analyses showed that not only the presence of phase III, but also its migration velocity determined clearance. Slow migration velocity was independently associated with Gram-negative bacilli colonization [12]. The migration velocity, the duration of each phase III activity, and the overall occurrence of phase III during prolonged recording in the fasting state were summarized by a migrating motor complex index. It was then possible to predict semiquantitatively the failure of intestinal clearance, as evidenced by Gram-negative bacilli in the small bowel, with a high sensitivity, superior to the qualitative evaluation of the presence or absence of phase III of the migrating motor complex [12, 21] (fig. 4, 5). The sensitivity and specificity of MMC index for the detection of Gram-negative bacilli in the duodenum were 91% and 90%, respectively [12].

There are also distinctly abnormal patterns of motility [21] that are independently associated with Gram-negative bacilli overgrowth, such as prolonged isolated irregular bursts and giant migrating contractions [12]. Accordingly, in enteric neuropathies uncoordinated contractile activity can cause temporal stagnation and even retroperistalsis [122–124].

Moreover, certain enteropathogenic microorganisms [186, 187] and commensal bacteria colonizing the small bowel in experimental bacterial overgrowth [188] induce giant migrating contractions that do not occur in healthy subjects [21]. Giant migrating contractions cause rapid intestinal clearance [189], and have been reported in patients with severe Gram-negative bacilli overgrowth with strict anaerobic bacteria of the intestinal type [12].

These data on the motility patterns of the small bowel and clearance concur with a recent study showing that Gram-negative bacilli (Enterobacteriaceae) in the small bowel are associated with delayed small bowel transit [111], and the early pioneer study on bacterial overgrowth showing the association between local stasis and Gram-negative colonization including strict anaerobes [75]. Moreover, the absence of phase III in a subset of patients with bacterial overgrowth has been reconfirmed [190].

Summary of Consequences for Intestinal Microflora

Failure of intestinal clearance caused by impaired motor activity or local stagnation for anatomical reasons results in Gram-negative colonization of the small bowel. Small bowel aspirate, mucosal brush, or biopsies are optional samples for culture, which is still the gold standard for detecting this type of overgrowth.

The absence of Gram-negative bacilli is a reliable and valid indication of preserved intestinal clearance, which precludes a significant failure of intestinal motility and anatomical abnormalities inducing stasis or recycling of contents from the lower gastrointestinal tract.

The presence of Gram-negative bacilli, however, can also be due to alterations in oral and gastric microflora, or to the general effects of illness and malnutrition. Further diagnostic workup to elucidate the pathogenesis is thus encouraged when bacterial overgrowth of Gram-negative bacilli is found in patients with clinically significant gastrointestinal symptoms to detect anatomical abnormalities or intestinal dysmotility.

The Significance of Changes in Local Mucosal and Systemic Immunity

Systemic Immunity

Humoral Immunity

Blood donors with selective IgA deficiency have a normal gastrointestinal microflora without evidence of bacterial overgrowth [191]. In patients with complex immunodeficiency increased bacterial colonization is seen in

the upper gut, predominantly by URT flora, which may be related to concurrent gastric hypo- or achlorhydria [54, 191]. Similar findings have been made in a limited study of jejunal flora in children [192].

Cellular Immunity

Patients with HIV display different degrees of failure of cell-mediated immunity. The prevalence of URT flora in the upper gut was in line with what was expected, and did not change with the clinical severity of the disease [193]. Colonization by Gram-negative bacilli was frequently found in HIV patients with diarrhea, regardless of its cause, but not in those with normal stools [193]. The cause and the consequences of Gram-negative bacilli in the upper gut of HIV patients remain unclear, but the degree of malnutrition [55] and illness [56] are likely to contribute. Children with a T cell defect have URT flora in the upper gut, but the prevalence is hardly significantly increased [192]. Duodenal microflora was examined in 32 patients with HIV infection [193]. Those with and without increased density of bacteria in the small bowel ($>10^5$ CFU/ml) had gastric pH 4.0 and 2.8 (nonsignificant), respectively. Gastric pH values for patients with and without Gram-negative bacilli flora in the duodenum were 3.3 and 3.2, respectively. This study could not establish a link between HIV infection and bacterial overgrowth, and provided further evidence that factors other than gastric hypochlorhydria explain the presence of Gram-negative bacilli in the small bowel.

Intestinal Mucosal Immunity

An identified specific defect of local mucosal immunity that results in bacterial overgrowth with URT flora or Gram-negative bacilli has not yet been detected, but there is evidence indicating that the mucosal immune system responds to a resident overgrowth flora with Gram-negative bacilli in the small intestine. The number of IgA2 immunocytes was increased in the jejunum, whereas the number of IgM immunocytes was reduced [194]. The increase in IgA2 may enhance mucosal protection and probably reflects immunomodulation caused by lipopolysaccharides of Gram-negative bacilli [194]. Accordingly, stimulated production of luminal IgA was recently reported in elderly patients with bacterial overgrowth with Gram-negative bacilli [195].

Moreover, bacterial overgrowth flora with Gram-negative bacilli, but not with URT flora, is associated with an increase in intraepithelial lymphocytes, reflecting an im-

Table 3. Algorithm for clinical management of bacterial overgrowth based on stratification by pathogenesis

GNB overgrowth	anatomical disorder ruled out or unsuitable for surgery	URT overgrowth no further diagnostic measures
<p>anatomical disorder possible</p> <p>Diagnostic workup Search for fistula, large or multiple small intestinal diverticula, and an enlarged surgical blind loop (X-ray) or confined segment of intestine</p> <p>Management If significant abnormality is detected, <i>consider surgical correction</i></p>	<p>Diagnostic workup Consider the presence of diseases and conditions associated with impairment of small bowel motility</p> <p>If present and clinical symptoms are significant, <i>test small bowel motility</i>^a</p> <p>Management If significantly abnormal, avoid drugs interfering with small bowel motility and/or intestinal microflora; give dietary advice to avoid nutrients requiring major grinding and mixing, and provide vitamin D, calcium, iron, and vitamin B₁₂ as indicated; optimize treatment of underlying disorder, and <i>provide drugs promoting small bowel motility</i></p>	<p>If patients are on PPI and nutritional deficiencies occur, the indication and further prescription should be reconsidered, in particular in the elderly</p>
<p>If none of the above measures are relevant or sufficiently effective <i>Provide antibiotics</i>, by preference poorly absorbed types, efficient against Enterobacteriaceae and strict anaerobic bacteria; give intermittent trials and cycle different antibiotics to reduce the risk of resistance <i>Avoid drugs</i> suppressing gastric acid secretion <i>Monitor effects</i> by symptomatic improvement, gain of body weight and improved blood tests as indicated: hemoglobin, calcium, albumin, iron, B₁₂ and folic acid</p>		

GNB = Gram-negative bacilli; PPI = proton pump inhibitors.

^a See text for testing of small bowel motility. If testing is not available, the management advice can be followed provided that significant colonization by Gram-negative bacilli is present in small bowel (see text for testing of bacterial overgrowth).

mune response in the small intestine [196]. These findings emphasize corresponding differences in the pathophysiology for the two types of bacterial overgrowth defined by pathogenesis.

From Pathogenesis to Clinical Management

Knowing the cause of the problem can facilitate and improve clinical management. In table 3 a clinical algorithm for dealing with each type of bacterial overgrowth is proposed based on the insight of the pathogenesis as discussed in the present review.

Failure of the gastric acid barrier and URT overgrowth are 'benign' alterations of gut function, as opposed to bacterial overgrowth with Gram-negative bacilli associated

with a wide spectrum of potential clinical problems. This difference should not be attributed solely to the bacteria, but rather to the underlying defect of the host. Failure of intestinal motility, for example, can lead to problems of digestion and transport independent of the Gram-negative bacilli in the lumen. The overgrowth flora reflects microbial adaptation that may produce additional clinical symptoms, depending on density and composition.

The clear distinctions in the pathogenesis, microbial flora and pathophysiology of bacterial overgrowth with URT flora and Gram-negative bacilli, respectively, encourage a classification based on the pathogenesis (table 1). As shown in table 3, corresponding distinctions can be made in the diagnostic workup and further clinical management.

Findings and Insights of Particular Interest

Suppression of the Gastric Acid Barrier

Reduced protein and vitamin B₁₂ assimilation, in particular in the elderly, is a recently recognized risk. Malabsorption due to acid deficiency, microbial metabolism of bile acids by the URT flora, and reduced reserve capacity of the gut in the elderly are candidate mechanisms, confined or combined. The increased risk of gastrointestinal infections is well established and more relevant nowadays with the marked increase in traveling between continents. This also involves elderly people, more often suffering from a failure of the gastric acid barrier due to *H. pylori*-induced gastritis or using proton pump inhibitors. Although no clear link exists between URT overgrowth in the upper gut and cancer, this issue has not been fully explored. Furthermore, precipitation or aggravation of bacterial overgrowth with Gram-negative bacilli in predisposed individuals has been demonstrated, and may be a problem with the more liberal use of long-term proton pump inhibitor treatment beyond study audit. Clinical data are still too sparse to justify management guidelines

for these issues, prompting clinical awareness and further research.

Failure of Intestinal Clearance

By recognizing significant anatomical abnormalities and intestinal dysmotility, attempts to restore the causal problems are encouraged. This is important, because the Gram-negative bacilli can be innocent bystanders, a clinically silent consequence of the underlying problem rather than the cause of the symptoms. Attempts to modulate the abnormal microflora by anti-, pre-, or probiotics are justified when treatment of the underlying problem is impossible or ineffective. This will often be the case, and antibiotics usually have temporary effect. Studies comparing antibiotics are now emerging [197, 198]. Drugs with effects on intestinal anaerobic and facultative bacteria are in general effective, and poorly absorbed antibiotics, like rifaximin [198], are good candidates because of limited systemic effects and antimicrobial action along the entire length of the small intestine. This rifamycin derivative indeed proved to be one of the most effective antimicrobials in the treatment of bacterial overgrowth [199].

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Note Added in Proof

After submission of this paper I came across the interesting review by Singh and Toskes [1] where the pathophysiology of bacterial overgrowth is carefully presented. In the most recent article by Lin [2] the possible role of bacterial overgrowth in the pathogenesis of irritable bowel syndrome is discussed, paying particular attention to the high prevalence of bloating in this syndrome.

The high prevalence of bacterial overgrowth in patients with chronic renal failure is a novel finding [3], and a condition to add to the list in the current review. Interestingly, concurrent abnormalities of small bowel motility were detected, of which increased prevalence of retrograde pressure waves in duodenum is of particular relevance in the present context [12, 183]. Castiglione et al. [4] further indicate the usefulness of both metronidazole and ciprofloxacin in the treatment of bacterial overgrowth associated with Crohn's disease. Indeed both clinical improvement and normalization of the lactose and glucose breath tests occurred after antibiotic treatment. The symbiotic modulation of the gut flora [5] is an interesting new approach for the management of minimal hepatic encephalopathy, emphasizing the importance of understanding the role of changes in gut flora.

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Pathophysiology and Impact of Enteric Bacterial and Protozoal Infections: New Approaches to Therapy

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Key Words

Diarrhea · Enteric infections · *Escherichia coli* · *Cryptosporidium* · *Shigella* · *Vibrio cholerae* · *Clostridium difficile* · *Salmonella*

Abstract

Despite numerous scientific advances in the past few years regarding the pathogenesis, diagnostic tools and treatment of infectious enteritis, enteric infections remain a serious threat to health worldwide. With globalization of the food supply, the increase in travel, mass food processing and antibiotic resistance, infectious diarrhea has become a critical concern for both developing and developed countries. Oral rehydration therapy has been cited as the most important medical discovery of the century due to the millions of lives that have been saved. However, statistics concerning diarrhea-induced mortality and the highly underestimated morbidity continue to demonstrate the severity of the problem. A more complete understanding of the pathogenesis of infectious diarrhea and potential new vaccines and effective treatments are badly needed. In addition, public health preventive actions, such as early detection of outbreaks, care with food, water and sanitation and, where relevant,

immunization, should be considered a priority. This article provides an overview of the epidemiological impact, pathogenesis and new approaches to the management of enteric infections.

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Introduction

Enteric infections are among the leading causes of mortality worldwide. They are estimated to cause 2.5 million deaths per year, which translates to over 6,800 children who die each day from diarrheal diseases, and the huge morbidity from diarrhea has not been reduced but may actually be increasing [1]. Even more impressive is the long-term developmental impact of repeated childhood diarrheal disease [2]. Diarrhea further aggravates malnutrition, especially in developing countries. Studies done in Brazil show that diarrhea in the first 2 years of life is associated with an approximate 4% decrement in physical fitness, a growth shortfall of 3.6 cm and impaired cognition and school performance at 6–9 years of age [3–6]. In addition to acute morbidity and mortality, some causes of infectious diarrheal disease result in serious long-term sequelae such as hemolytic-uremic syndrome with renal

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failure following enterohemorrhagic *Escherichia coli* (EHEC) infection (a risk that may be increased by treatment with certain antimicrobial agents such as sulfa-trimethoprim or quinolones). Other examples include Guillain-Barré syndrome following *Campylobacter jejuni* infection and malnutrition with or without diarrhea following infection with enteroaggregative *E. coli* (EAggEC), *Cryptosporidium* species or other enteric infections [7–10].

With the globalization of our food supply and increasing international travel, enteric infection is now also a serious threat to industrialized countries, as demonstrated in recent years by diarrheal outbreaks in North America due to *Cyclospora* following ingestion of imported Guatemalan raspberries [11]. Other examples include water-borne outbreaks caused by *Cryptosporidium* and food-borne outbreaks caused by EHEC. The economic cost of infectious diarrheal diseases is also considerable. In the United States, an estimated USD 6 billion each year is spent on medical care and loss of productivity due to food-borne diseases, most of which cause diarrhea [12, 13]. The exploding developing world's population, the disparity between the rich and the poor, and emerging antibiotic-resistant infections make enteric infections a critical global health concern.

This article provides an overview of the epidemiological impact, pathogenesis and new approaches to the management of enteric infections. Although several enteric viruses are important causes of diarrhea in both developed and developing country, we will focus this overview on bacterial and selected parasitic pathogens.

Epidemiology

EHEC, *Salmonella*, *Shigella*, *Vibrio cholerae*, *Cyclospora*, *Cryptosporidium*, *Giardia*, *C. jejuni*, *Clostridium difficile*, caliciviruses and other viruses such as rotavirus, astrovirus and torovirus are the main causes of diarrhea worldwide, and cause more than 211–375 million cases of diarrheal illnesses in the United States each year [14, 15]. In addition, in the last 2 or 3 decades, other enteric pathogens have been recognized as emerging causes of enteric infections. There are now several types of *E. coli* enteropathogens in addition to the classical enteropathogenic *E. coli* (EPEC), including enterotoxigenic *E. coli* (ETEC), which produces a cholera-like heat-labile toxin (LT) or heat-stable toxins STa or STb, EHEC, which produces a Shiga-like toxin (SLT), enteroinvasive *E. coli* (EIEC) and EAggEC, which is associated with persistent diarrhea in

developing and developed countries. Among the parasitic protozoa, microsporidia can also be included as emerging infectious pathogens especially in immunocompromised hosts, as well as *Cyclospora* and *Cryptosporidium* [16, 17].

Many of these organisms are easily transmitted through food and water or by human contact. Thus, prevention by avoiding the ingestion of raw or undercooked meat, seafood or unpasteurized milk products, and the selective use of available vaccines are the key to the control of infectious diarrhea.

In the United States alone, episodes of diarrheal illness result in 73 million physician consultations, 1.8 million hospitalizations and 3,100 deaths each year. Food-borne illnesses alone account for 76 million illnesses and 350,000 hospitalizations each year [15, 18, 19].

Traveler's diarrhea is a common problem that occurs in 20–50% of the 35 million people who cross international borders from the developed to tropical or semitropical developing countries every year, resulting in more than 7 million cases [20–23]. The etiological agents of traveler's diarrhea depend on the geographical location, standards of food, hygiene, sanitation, water supply and season. The most common causes of traveler's diarrhea in adults in developed countries include *E. coli*, specially ETEC, *Shigella* spp., *Salmonella* spp., *Campylobacter* spp., *Vibrio parahaemolyticus* (in Asia), rotavirus (in Latin America) and protozoa (*Giardia*, *Cryptosporidium* and *Cyclospora* spp., and *Entamoeba histolytica*) [24, 25].

Pathogenesis

There are several mechanisms by which enteric pathogens can cause diarrhea (table 1). Recent progress in understanding of the pathogenesis at the molecular level opens new perspectives on the treatment of infectious diarrhea. Furthermore, different microorganisms often share common pathogenic pathways. Microbes must first adhere to the mucosa in order to elicit disease. Thereafter, some microorganisms such as *V. cholerae* or ETEC produce toxins that can subvert ion transport across the intestinal epithelium. Other microorganisms such as *Shigella* and *Salmonella* species can invade the mucosa causing inflammation. In extreme cases, microorganisms can also invade the bloodstream [26]. Other organisms such as *C. difficile* produce enterocytotoxin, which causes intense disruption of the intestinal mucosa [27].

Aside from the features of the microorganisms cited above, the host defenses also play an important role in the

Table 1. Clinical, epidemiological and pathogenic features of enteric infections

Pathogenic agent	Epidemiology	Incubation period	Diarrhea	Virulence determinant/mechanism
<i>V. cholerae</i>	all ages in developing world/related to poor sanitation	18–40 h but can be as short as 12 h or as long as 72 h	profuse watery diarrhea causing severe dehydration	cholera toxin → G _s protein → adenylate cyclase → secretion → prostaglandin → secretion → enteroendocrine cells → endogenous secretagogues → secretion
ETEC	young children > adults in developing world/travelers to tropics	10–72 h	acute watery	CFA-I-IV → colonization LT-I and -II → adenylate cyclase → secretion STa → guanylate cyclase → secretion STb → cyclic nucleotide-independent HCO ₃ ⁻ secretion
EPEC	infants in developing world	as short as 9–12 h	acute → persistent watery	not fully understood, possibilities are increase in mucosal permeability and loss of microvilli leading to malabsorption
EHEC	all ages/primarily in US, Canada, Europe, South America and Japan	12–60 h	acute bloody (hemorrhagic colitis in 31–61%); occasionally nonbloody diarrhea	SLT-I and -II → bloodstream → inhibition of protein synthesis → endothelial cell damage → microvascular thrombosis → hemolytic-uremic syndrome
EAggEC	children in the developing world	20–48 h	persistent	FliC → inflammation EAST-1 → guanylate cyclase → secretion heat-labile toxin → Ca ²⁺ -dependent actin phosphorylation; cytoskeletal damage Pet → histopathologic effects on human intestinal mucosa
EIEC	all ages/primarily in the developing regions; occasional outbreaks in industrialized countries	as short as 10–18 h	acute watery diarrhea followed by dysentery	cell invasion → spread → inflammation
<i>C. difficile</i>	history of antibiotic use, advanced age, underlying illness	5–10 days of anti-bacteria treatment (range 1st day to 10 weeks of antibiotics)	mild to severe inflammatory diarrhea	toxins A and B → monoglucosylation of Rho protein → disruption of actin cytoskeleton → mucosal disruption. → COX-2 → prostaglandin E ₂ → synthesis of inflammatory cytokines
<i>Cryptosporidium</i>	all ages/children in developing areas/immunocompromised adults/outbreaks in developed areas	7–10 days (range 5–28 days)	intermittent and scant to continuous and watery	prostaglandins → cAMP-mediated apical chloride secretion and inhibition of electroneutral sodium chloride and water absorption release of IL-1, IL-8 and TNF- α
<i>Salmonella</i>	all ages/travelers to tropics	6–48 h	moderate volume, and usually without blood	mucosal invasion via M cells or enterocytes → macrophages and lymphocytes in Peyer's patches and other lymphoid tissue → bloodstream
<i>Shigella</i>	incidence highest in children 1–5 years of age	24–72 h	watery at the onset and may evolve to bloody diarrhea or dysentery	invasion and destruction of the distal ileal and colonic mucosa → release of cytokines → PMN mucosal infiltration

FliC = Flagellin sequence in EAggEC responsible for IL-8 induction [64].

acquisition of enteric infection. Host defenses include normal gastric acidity, intestinal mucus, cellular and humoral immunity, motility and intestinal microbial flora.

A bacterial enteric infection may manifest as diarrhea or may also remain asymptomatic. Recently, it was recognized that even asymptomatic enteric infections by *Cryptosporidium*, EAggEC and *Giardia lamblia* may be associated with nutritional shortfalls, even in the absence of overt diarrheal illness [17].

Intestinal infections that cause persistent diarrhea normally result in histopathological changes to the intestine including villus blunting, crypt hypertrophy and inflammatory infiltrate in the lamina propria. These histopathological disarrangements are seen in *Cryptosporidium*, *Cyclospora* and microsporidial infections [28]. Furthermore, it has been documented that there are substantial disruptions of intestinal barrier function as measured by lactulose:mannitol permeability ratios in patients with AIDS

and in children with diarrhea in northeast Brazil [29, 30].

Among functional alterations in patients with infectious diarrhea are increased secretion, failure of barrier function and reduction of absorptive function causing dehydration and nutritional deficiency. An understanding of the molecular pathogenesis with regard to each enteric pathogen will likely lead to a quicker diagnosis, more effective treatment and prevention of enteric infections.

Vibrio cholerae

V. cholerae (01 and 0139) pathogenesis has been extensively studied. This pathogen causes a devastating diarrhea characterized by severe dehydration without mucosal disruption or invasion. The microbe interacts with the host cell mainly in the proximal small intestine where the motile vibrios penetrate the mucus and bind to the enterocytes via toxin-coregulated pili, producing several toxins including cholera toxin [31]. Cholera toxin binds to the membrane of enterocytes and is subsequently internalized, thus causing activation of the catalytic unit of the stimulatory G protein (G_s). The activation of G_s protein results in uncontrolled production of cyclic AMP (cAMP), which inhibits sodium absorption and induces chloride secretion [32–34]. For decades, this was the only mechanism that explained the large loss of liquid associated with cholera-induced diarrhea. However, there is now evidence that prostaglandins are also involved in the secretion induced by cholera toxin [35]. Additionally, it has been shown that cholera toxin interacts with enteroendocrine cells, stimulating the release of endogenous secretagogues. Cholera toxin also interacts with the enteric nervous system, altering electrolyte transport and motility [36].

Escherichia coli

Several types of *E. coli* have been recognized, each with its own pathogenesis. ETEC is a major cause of dehydrating infant diarrhea in the developing world. It is also the most common cause of travelers' diarrhea [31, 37]. Like *V. cholerae*, ETEC causes an acute, watery diarrhea following the ingestion of contaminated water or food. The incubation period has been found to be 10–72 h. The organism attaches via the fimbrial colonization factor antigens (CFAs), multiplies in the proximal small intestine and produces one or more enterotoxins [38, 39]. Of the four known enterotoxins (LT-I, LT-II, STa, STb) produced by ETEC, LT-I and STa are well established in the literature as important human secretagogues. LT-I is simi-

lar to cholera toxin with respect to structure and mechanism. After binding to a GM_1 ganglioside receptor, LT-I activates adenylate cyclase, resulting in an increase of the intracellular levels of cAMP, which ultimately stimulates chloride secretion and inhibits sodium absorption [40, 41]. The ST toxin family bears significant homology to the endogenous intestinal peptide guanylin. STa binds to an extracellular domain of particulate guanylate cyclase, resulting in increased intracellular levels of cyclic guanosine monophosphate (cGMP), which leads to decreased absorption of sodium and increased chloride secretion [42]. Protective immunity to ETEC appears to be mediated by secretory IgA antibodies directed against fimbriae and LT. One of the most promising vaccine candidates, now in a phase III clinical trial, is an oral ETEC vaccine containing recombinant cholera B subunit in combination with five different formalin-inactivated *E. coli* strains expressing common fimbrial CFA-I and coli surface antigen 1–6 [31].

EPEC causes a degeneration of the microvillus brush border, with 'cupping and pedestal' formation of the plasma membrane at the sites of bacterial attachment and reorganization of cytoskeletal proteins [43, 44]. Invasion has been observed in some clinical specimens, but the mechanism of how this bacteria produces diarrhea is not fully understood. Some possibilities include an increase in permeability and loss in microvilli leading to malabsorption.

With mass food processing and fast food practices, EHEC, also called Shiga toxin-producing *E. coli*, has emerged as an important bacterial pathogen in industrialized countries [45]. Like EPEC, EHEC causes filamentous actin accumulation at the site of attachment in association with 'cup and pedestal' formation [46]. The toxins of EHEC bear both structural and functional similarity with Shiga toxin and are named SLTs or verotoxins, reflecting their cytotoxic effect in Vero cells. There are at least two immunologically different forms of SLT (SLT-I and SLT-II). These toxins are capable of inducing secretion and mucosal injury in animal models [47]. In severe disease, especially when bloody diarrhea is present, it is thought that these toxins gain access to the bloodstream and are involved in the pathogenesis of hemolytic-uremic syndrome. It is proposed that SLT binds to receptors on host cells named Gb_3 (glycolipid globotriaosylceramide) [48]. The variability in surface expression of this receptor determines the cell susceptibility to damage induced by these toxins. In addition, the proliferation rate and tissue origin of endothelial cells influence their susceptibility to the cytotoxicity of these toxins [49, 50]. For example,

human renal and intestinal endothelial cells are very sensitive to SLTs [48, 51, 52], whereas human brain endothelial cells and endothelial cells derived from large vessels such as saphenous vein or human umbilical vein are relatively resistant [50, 53, 54]. Evidence shows that cocultivation of endothelial cell culture with proinflammatory cytokines, such as interleukin (IL)-1 and tumor necrosis factor (TNF)- α , stimulates the expression of Gb₃ and markedly increases the cytotoxicity of the toxins towards endothelial cells [50, 53]. The damage of endothelial cells stimulates the expression of adhesion molecules, leading to leukocyte recruitment [55]. Activation of adherent leukocytes would result in the release of leukocyte products, such as reactive oxygen metabolites and proteases which exacerbate the endothelial damage. The detachment of endothelial cells as a result of direct and indirect effects of SLTs expose the basement membrane and underlying matrix, initiating the coagulation characteristic of hemolytic-uremic syndrome [56–58].

EAggEC include a heterogeneous group of organisms, some strains exhibiting no virulence. The characteristic HEp-2 adherence occurs via a flexible bundle-forming fimbrial structure, aggregative adherence fimbriae I [59, 60]. EAggEC also secretes an enterotoxin named EAST-1, which bears homology to domains of guanylin and ST, sharing with them the capacity to increase cGMP and induce secretion [61]. However, the role of EAST-1 in EAggEC-induced diarrhea is questionable given the lack of diarrhea in volunteers challenged with EAST-1-producing EAggEC strains that colonized the intestine at high levels [62]. EAggEC is also able to produce a heat-labile toxin which increases the intracellular level of calcium and stimulates calcium-dependent phosphorylation [63], but no *in vivo* effect of this protein has been shown. It was also shown that a product from EAggEC induces secretion of IL-8 by intestinal epithelial cells *in vitro* [8], and this could contribute to the intestinal inflammation detected in children with EAggEC infection [8]. This IL-8-releasing factor from EAggEC has been cloned, sequenced and expressed as a unique flagellin [64]. In addition, a 104-kD protein termed Pet (plasmid-encoded toxin), secreted by some strains of EAggEC, has been cloned and sequenced and bears homology to a class of serine protease auto-transporter proteins from *E. coli* and *Shigella* spp. [65]. Pet raises the transepithelial short-circuit current, decreases the electrical resistance of rat jejunum mounted in an Ussing chamber, causes contraction of cytoskeleton and loss of actin stress fibers, and is required for the histopathologic effects of EAggEC on human intestinal mucosa [65–67]. A definitive role of these virulent factors in the

pathogenesis of EAggEC diarrhea remains to be established.

EIEC invades and multiplies within colonic epithelial cells, causing cell death and inducing inflammation. This inflammatory response, along with necrosis and ulceration of the large bowel, leads to a bloody and mucoid diarrhea. Among the virulence factors, a 140-MD plasmid has been described to encode the genes responsible for outer membrane proteins important for invasion [68]. In addition, some strains produce enterotoxins capable of inducing secretion in Ussing chambers that might play a role in the watery diarrhea seen after an incubation period as short as 10–18 h [69, 70].

Clostridium difficile

C. difficile colonization and infection occur in the setting of altered intestinal microflora, usually precipitated by antibiotic exposure. Colitis and diarrhea are mediated by large exotoxins, *C. difficile* toxin A and toxin B. These toxins are produced intraluminally, bind to specific epithelial surface receptors and are internalized [71, 72]. Once in an intracellular location, both toxins monoglucosylate small GTP-binding proteins. Modification and inactivation of small GTPases (Rho, Rac, Cdc42) cause disruption of the actin cytoskeleton [73, 74]. This leads to loosening of tight junctions and eventually mucosal disruption. Interestingly, toxin A also appears to alter the morphology of neutrophils and adversely affect nondirected and directed migration induced by FMLP (f-met-leu-phe) through inactivation of Rho [75]. Toxin A-negative/toxin B-positive strains have been documented to cause disease, even nosocomial outbreaks [76]. Although both toxins can cause clinical disease, many of the secretory and inflammatory effects of *C. difficile* infection are attributed to toxin A, while the cytopathic effect is more prominent with toxin B. Toxin A has been demonstrated to cause the release of proinflammatory cytokines in an animal model [77]. Indeed, upregulation of IL-8 transcription [78] and generation of prostaglandin E₂ by inducing cyclooxygenase-2 (COX-2) expression have been recently reported along with blockade of toxin A-induced secretion and inflammatory damage by COX-2 inhibition [79]. Chemotaxis of polymorphonuclear cells and monocytes and recruitment of mast cells further contribute to the intense inflammatory reaction [80, 81]. Activation of the enteric nervous system as evidenced by increased substance P in intestinal macrophages and dorsal root ganglia in toxin A-induced enteritis in rats has also been demonstrated [82]. Disruption of the epithelial barrier, release of proinflammatory cytokines and recruitment of immune

and inflammatory cells all contribute to fluid accumulation and mucosal injury.

Cryptosporidium

The *Cryptosporidium* parasite attaches to the host's intestinal epithelium, becomes intracellular but remains extracytoplasmic. In vitro studies suggest that attachment is mediated by a *Cryptosporidium parvum* sporozoite ligand and an intestinal epithelial cell surface protein interaction [83, 84].

Although infection with *C. parvum* is considered predominantly secretory, histopathologic studies have revealed varying degrees of villous atrophy and infiltration of inflammatory cells beneath the epithelial mucosa [85, 86]. Prostaglandins, which are known to induce cAMP-mediated apical chloride secretion and inhibit electroneutral sodium chloride and water absorption in enterocytes, have been demonstrated to be elevated in a porcine model of cryptosporidiosis [87]. Inflammatory cytokines such as IL-1, IL-8 and TNF- α are induced in intestinal epithelial cell lines infected with *Cryptosporidium* and in animal models of cryptosporidiosis and have been postulated to play a role in pathogenesis [88, 89]. Expression of TNF- α and IL-1 mRNA in the majority of jejunal biopsies of adult volunteers after experimental infection were also observed, although this did not correlate with the enteric symptoms [90].

Lactoferrin, a protein found in secondary granules of polymorphonuclear cells, was observed to be mildly to moderately elevated in the stools of children with endemic cryptosporidiosis [91] and healthy adult volunteers with experimental infection [92]. Indeed, in another study of malnourished children in Haiti, cryptosporidiosis was noted to stimulate an inflammatory response, as evidenced by elevated IL-8, TNF- α , lactoferrin, IL-13 and IL-10 [93]. Further studies are needed to elucidate the role of inflammatory mediators in the development of prolonged diarrhea, malabsorption and malnutrition in immunocompromised hosts and children in endemic areas.

Shigella

Shigella is the most common etiological agent of dysentery. Initially, this pathogen produces a watery diarrhea, followed by the onset of dysentery that is characterized by scanty stools of blood and mucus. The pathogen invades the mucosa of the distal ileum and colon via the M cells overlying the gut-associated lymphoid tissue [94, 95]. Invasion plasmid antigens, which are secreted by the bacteria on contact with M cells or epithelial cells, lead to reorganization of the cytoskeleton through activation of

small GTPases of the Rho family and recruitment of the protooncogene *c-src*, resulting in internalization of the bacterium by macropinocytosis [95–97]. The internalized bacterium lyses its phagocytotic vacuole and initiates intracytoplasmic movement, resulting from polar assembly of actin filaments caused by a bacterial surface protein, VirG (also called IcsA), which binds and activates neuronal Wiskoff-Aldrich syndrome protein, thus inducing actin nucleation [98–101]. Actin-driven motility promotes efficient colonization of the host cell cytoplasm and rapid cell-to-cell spread via protrusions that are engulfed by adjacent cells in a cadherin-dependent process [102]. Bacterial invasion causes an intense proinflammatory response from invaded cells through activation of nuclear factor- κ B [103]. A major consequence is IL-8 production, which attracts polymorphonuclear leukocytes (PMN) [104]. On transmigration, PMN disrupt the permeability of this epithelium and promote its invasion by *Shigella*, leading to mucosal ulceration and microabscess formation [31]. Subsequent apoptotic killing of macrophages in a caspase 1-dependent process causes the release of IL-1 β and IL-18, which accounts for the initial steps of inflammation [105–107]. There are four species or groups of *Shigella*: *S. dysenteriae*, *S. flexneri*, *S. boydii* and *S. sonnei*. All include multiple serotypes, complicating vaccine development strategies. One approach that is being followed is to prepare conjugate vaccines for parenteral administration by covalently linking O polysaccharides of the most prevalent *Shigella* serotypes to carrier proteins [108]. Another approach that has been studied is that of attenuated strains. Investigators have attempted to apply tools of biotechnology to develop modern attenuated strains of *Shigella* that can serve as live oral vaccines. One of these prototype vaccines contain a strain that harbors a mutation in a plasmid virulence gene *icsA* (i.e. *virG*) that limits the intra- and intercellular spread of the bacteria, combining with other mutations. Proteosomes are outer-membrane proteins of meningococci that are highly hydrophobic and assemble into membranous vesicles and can combine with antigens to form a competent antigen delivery system. One of the most successful uses of proteosomes has been to prepare complexes with the lipopolysaccharides of *S. sonnei* and *S. flexneri* [31, 109]. Clinical trials of these candidate vaccines are currently under way.

Salmonella

Salmonella species are a major source of food-borne disease throughout the developing and the developed countries [110]. This pathogen invades the mucosa

through the M cells or through enterocytes, resulting in the extrusion of infected epithelial cells into the intestinal lumen with consequent villus blunting and loss of absorptive surfaces. *Salmonella* also elicit a PMN influx into infected mucosa and induce watery diarrhea, which may contain blood [111]. Wallis and Galyov [111] reviewed and proposed a sequence of events occurring during the pathogenesis of *Salmonella*-induced enteritis: (1) *Salmonella* interacts with enterocytes and delivers *Salmonella* outer proteins (Sops) into the cell cytoplasm via a TTSS-1 (TTSS are secreted virulence-associated effector proteins) and *Salmonella* invasion protein (Sip)-dependent pathway. (2) Sips, SopE and possibly other Sops induce enterocyte membrane ruffling promoting bacterial invasion. (3) Intracellular bacteria reside within membrane-bound vesicles and possibly continue translocation of TTSS-1 secreted effectors. The replication of *Salmonella* within the vesicles is promoted by TTSS-2. (4) The intracellular SopB protein affects inositol phosphate signaling events, causing a transient increase in the concentration of Ins(1,4,5,6)P₁, which in turn can antagonize the closure of chloride channels, influencing net electrolyte transport and thus fluid secretion. (5) *Salmonella*-infected epithelial cells secrete chemokines and prostaglandins that act to recruit inflammatory cells to foci of infection. The release of at least some chemokines and prostaglandins is probably affected by the intracellular activity of Sops. (6) *Salmonella* interacts with inflammatory cells and stimulates the release of proinflammatory cytokines that enhance the inflammatory response. *Salmonella*-infected epithelial cells release pathogen-elicited epithelial chemoattractant across the apical membrane, which stimulates PMN transepithelial migration between the enterocytes. (8) Infiltrating inflammatory cells phagocytose *Salmonella*. (9) *Salmonella*-infected enterocytes become extruded from the villus surface, leading to shedding of infected cells into the intestinal lumen and resulting in villus blunting and loss of absorptive surfaces. (10) Some of the infected cells migrate to the draining lymphatics, carrying *Salmonella* to systemic sites.

Campylobacter

Although not reviewed in detail here, *C. jejuni* and *C. coli* are another major cause of inflammatory colitis that may be complicated by Guillain-Barré syndrome or reactive arthritis. In addition, their resistance to antimicrobials (particularly to quinolones) is increasing. In the United States, fluoroquinolone resistance of *C. jejuni* rose from 13% in 1997 to 18% in 1999 [112].

Management of Enteric Infections

Because the most common risks of diarrhea are dehydration and malnutrition, the critical initial treatment must be rehydration. Thus, the first approach to patients with enteritis should be the evaluation of their hydration status by checking mucosal hydration, skin turgor and orthostatic changes in pulse and blood pressure. Oral or intravenous rehydration therapy should precede any search for etiological diagnosis. Although both methods are life-saving procedures, oral rehydration is better tolerated, safer and more inexpensive than intravenous fluid administration. Some patients with mild diarrhea can compensate for water loss in the stool by ingesting more dietary liquids such as soups, juices, etc. However, patients with severe diarrhea may need additional rehydration. Rehydration can be accomplished by providing the patient with an oral solution containing electrolytes and glucose. The concentrations recommended by the World Health Organization are as follows: glucose 111 mM, Na 90 mM, K 20 mM, Cl 80 mM and HCO₃ 30 mM [113]. The principle behind the use of this solution is that nutrients such as glucose and amino acids are transported across the apical membrane of the enterocyte by a carrier that cotransports sodium [114]. Unlike apical sodium-hydrogen exchange, nutrient-sodium cotransport is not impeded by elevated intracellular cAMP levels [115]. Recently, it has been shown that glutamine and especially its stable derivative alanyl-glutamine may not only increase sodium absorption but also improve the repair of intestinal epithelium after damage [116–120]. Alanyl-glutamine is more advantageous than glutamine due to its much greater solubility and its stability in solution and in acidic conditions such as the stomach. Other advantages include its ability to be heat sterilized and capacity for long term storage (US patent No. 5,561,111).

The second step in the management of the patient with enteritis is the collection of a detailed clinical history including the epidemiological features. Relevant clinical information includes symptomatic onset, stool characteristics (watery, bloody, mucous, purulent, greasy, etc.), frequency of bowel movements, quantity of stool produced, presence of dysenteric symptoms (fever, tenesmus, blood and/or pus in the stool), symptoms of volume depletion (thirst, tachycardia, orthostasis, decreased urination, lethargy, decreased skin turgor) and associated symptoms (nausea, vomiting, abdominal pain, cramps, headache, myalgias, altered sensorium). In addition, all patients must be asked about potential epidemiological risks such as travel to endemic areas, day care center attendance or

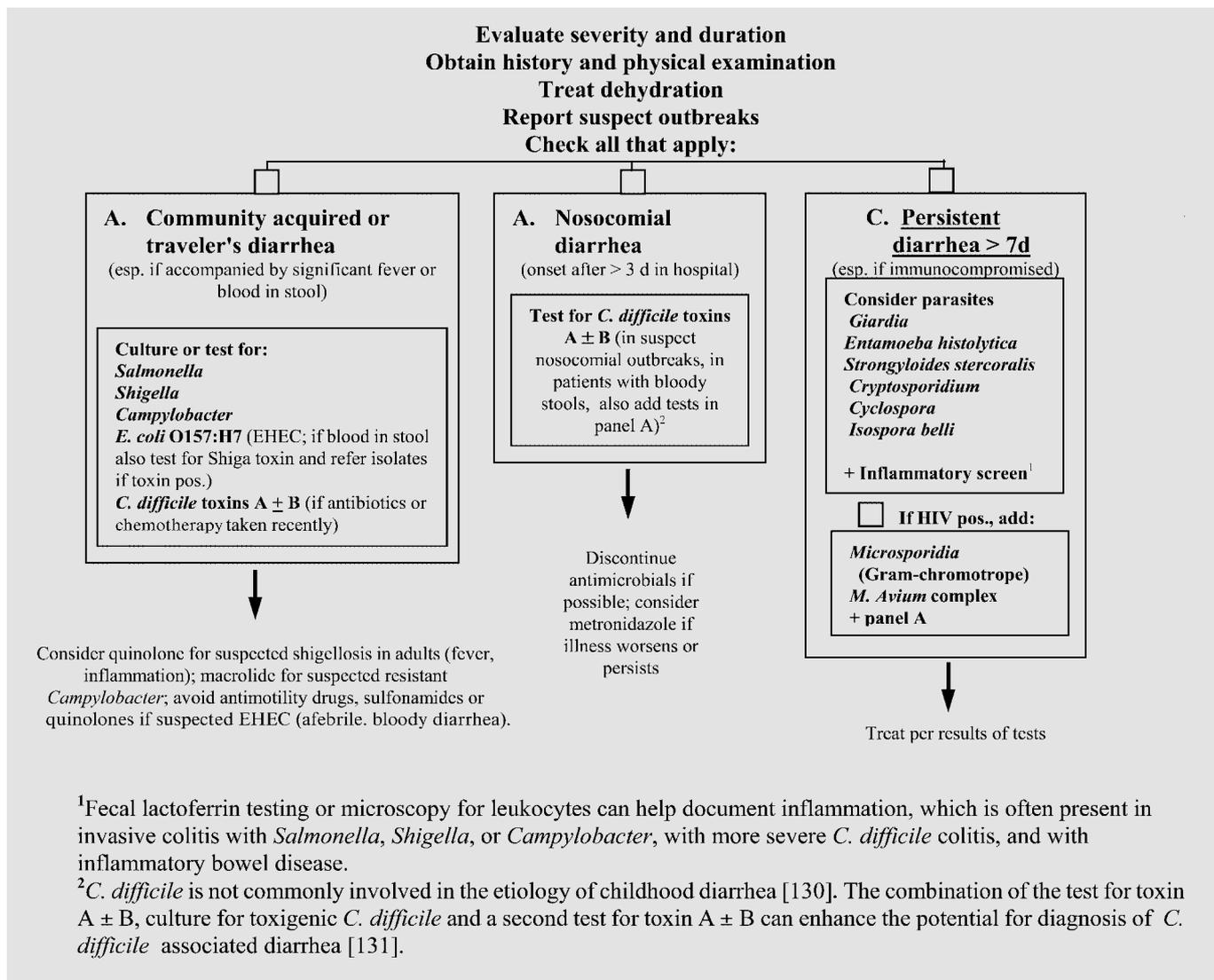


Fig. 1. Recommendations for the diagnosis and management of enteric infections. Adapted from Guerrant et al. [113], Infectious Diseases Society of America Practice Guidelines for the Management of Infectious Diarrhea.

employment, consumption of unsafe foods (raw meats, seafood, unpasteurized milk or juices), contact with pets with diarrhea, use of antibiotics and underlying medical conditions (AIDS, immunosuppressive medication, etc.).

Physical examination is essential for evaluation of signs of hydration status. In addition, it is important to screen for the presence of fever, and to evaluate diagnostic findings that may indicate another etiology.

Combining clinical and epidemiological features with fecal analysis gives important clues to the etiological diagnosis. For example, any patients with diarrheal illness lasting more than 1 day, accompanied by fever, bloody

stools, systemic illness, signs of serious dehydration and recent use of antibiotics, day care attendance and hospitalization should have a fecal sample specimen sent for evaluation. Additional laboratory exams may be necessary for selected cases. The Infectious Diseases Society of America Practice Guidelines for the Management of Infectious Diarrhea recommend a selective approach such as that shown in figure 1 [113].

With the increasing appearance of antibiotic-resistant infections, the side effects of antibiotics and superinfection as a consequence of the disturbance of the intestinal microflora, the immediate decision to use antibiotics

should be reconsidered. Although most forms of traveler's diarrhea can be managed effectively with symptomatic treatment alone, with agents such as loperamide or bismuth preparations, empirical antibiotics are commonly recommended. Treatment with fluoroquinolone or, in children, trimethoprim-sulfamethoxazole (TMP-SMZ) can reduce the duration of the diarrhea from 3–5 days to less than 1–2 days [24, 113]. Some also consider empirical treatment of diarrhea that lasts longer than 10–14 days for suspected giardiasis, if other evaluations are negative and, especially, if the patient's history of travel or water exposure is suggestive [113]. For patients with febrile diarrheal illnesses, especially those believed to have moderate to severe invasive disease, empirical treatment should be considered after a fecal specimen is obtained for performance of the studies noted in figure 1. This empirical treatment can be with an agent such as a quinolone antibiotic or, for children, TMP-SMZ [113]. The increasing worldwide resistance to TMP-SMZ and the more recently reported resistance to fluoroquinolones [121] is a driving force for the development of new antimicrobial agents, such as rifaximin and azithromycin. Rifaximin (a rifamycin derivative) is poorly absorbed when administered orally, but it has been shown to be safe and effective in comparison with TMP-SMX and ciprofloxacin [122, 123]. A study performed with adult students from United States in Mexico or international tourists in Jamaica showed that treatment for 3 days with rifaximin (400 mg twice a day) was as effective as ciprofloxacin (500 mg twice a day) with regard to the duration of disease, clinical improvement and microbiological cure. The incidence of adverse events was low and similar in each group [123]. Another study suggested that rifaximin (600 mg, 3 times a day, for 14 days) may improve clinical symptoms and clearing of protozoan infections in HIV-1-infected patients with $CD4 \geq 200/mm^3$ who presented with *Cryptosporidium* or *Blastocystis* associated with bacteria [124]. Additionally, the effect of rifaximin was compared with neomycin plus bacitracin in children with bacterial diarrhea in which the etiologic agents were *Salmonella* spp. and EPEC. Rifaximin yielded bacteriological cure in 12 out of 14 children, the reference drug in 13 of 17. With both drugs, the stool number per day fell after 1 day; within 2 days, stool consistency shifted to normal [125].

Because the development of antibiotic resistance will continue to be a problem, the development of effective alternative treatments is imperative. Immunization, probiotics, antisecretory agents, improved oral rehydration and nutrition therapy and nonabsorbable antibiotics are being considered by clinicians and researchers. Novel

therapeutic agents, other than antimicrobials, include the 5-hydroxytryptamine-2 and -3 receptor antagonists [126, 127], calcium-calmodulin antagonists, zaldaride maleate and σ -receptor agonist igmesine [126]. An enkephalinase inhibitor named racecadotril has also been developed, based on the antisecretory role of the neurotransmitter enkephalin, and it has been reported to have good efficacy and tolerability in clinical trials [128].

Prevention

Education, simple rules of personal hygiene and safe food preparation can prevent many diarrheal diseases. Hand washing with soap is an effective step in preventing spread of illness. Human feces must always be considered potentially hazardous. Immunocompromised persons, alcoholics, persons with chronic liver disease and pregnant women may require additional attention, and health care providers can play an important role in providing information about food safety. These populations should avoid undercooked meat, raw shellfish, raw dairy products, French-style cheeses and unheated deli meats [114].

Several bacterial pathogens have been targeted as a priority for the development of new or improved vaccines, such as *V. cholerae*, *Shigella*, *E. coli* and *Salmonella*. Substantial progress in molecular biology, bacterial pathogenesis, and immunology make possible the development of new candidate vaccines, but the evaluation of these candidates is a long and expensive process [31, 129]. In the developing countries, where the incidence of diarrhea is greater, financial resources are scarce and few countries have incorporated immunization for enteric pathogens into their immunization program. In the USA, only cholera and typhoid fever vaccines are commercially available. Immunization is recommended for typhoid fever (types Vi, Ty21a or the heat-phenol-inactivated vaccine for those under 2 years of age) in individuals living in or traveling to high-risk areas. Two modern oral cholera vaccines have been licensed by regulatory authorities in a number of countries. One is a nonliving vaccine consisting of inactivated *V. cholerae* O1 administered in combination with the B subunit of cholera toxin, so-called B subunit whole-cell cholera vaccine. The other vaccine is a genetically engineered attenuated strain of *V. cholerae* O1, CVD 103-HgR, which is used as a single-dose live oral vaccine [31]. Older cholera vaccines are not recommended in the US because of their limited efficacy and the low risk of cholera to the traveler [114].

Conclusion

Emerging infectious pathogens, increasing antimicrobial resistance, recognition of the long-term impact of diarrheal diseases and the appearance of diseases that decrease the host defense have heightened the necessity to develop new and more specific treatments and further

clarify the pathogenesis of diarrheal illnesses. New antibiotics, vaccines and micronutrients that improve mucosal recovery and host defenses are currently being tested. Additionally, it is critical to prevent enteric infections by increasing vaccination and improving sanitary conditions and the availability of safe drinking water.

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Rifaximin, a Poorly Absorbed Antibiotic: Pharmacology and Clinical Potential

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Key Words

Rifaximin · Rifamycin · Antibiotic · Gut bacteria · Enteric infection · Diarrhea, infectious · Hepatic encephalopathy · Small intestine bacterial overgrowth · Inflammatory bowel disease · Colonic diverticular disease · Irritable bowel syndrome · Constipation · *Clostridium difficile* infection · *Helicobacter pylori* infection · Colorectal surgery · Bowel decontamination, selective · Pancreatitis, acute · Bacterial peritonitis, spontaneous · Nonsteroidal anti-inflammatory drug enteropathy

Abstract

Rifaximin (4-deoxy-4'-methylpyrido[1',2'-1,2]imidazo[5,4-c]-rifamycin SV) is a synthetic antibiotic designed to modify the parent compound, rifamycin, in order to achieve low gastrointestinal (GI) absorption while retaining good antibacterial activity. Both experimental and clinical pharmacology clearly show that this compound is a nonsystemic antibiotic with a broad spectrum of antibacterial action covering Gram-positive and Gram-negative organisms, both aerobes and anaerobes. Being virtually nonabsorbed, its bioavailability within the GI tract is rather high with intraluminal and fecal drug concentrations that largely exceed the minimal inhibitory concen-

tration values observed in vitro against a wide range of pathogenic organisms. The GI tract represents, therefore, the primary therapeutic target and GI infections the main indication. The appreciation of the pathogenic role of gut bacteria in several organic and functional GI diseases has increasingly broadened its clinical use, which is now extended to hepatic encephalopathy, small intestine bacterial overgrowth, inflammatory bowel disease and colonic diverticular disease. Potential indications include the irritable bowel syndrome and chronic constipation, *Clostridium difficile* infection and bowel preparation before colorectal surgery. Because of its antibacterial activity against the microorganism and the lack of strains with primary resistance, some preliminary studies have explored the rifaximin potential for *Helicobacter pylori* eradication. Oral administration of this drug, by getting rid of enteric bacteria, could also be employed to achieve selective bowel decontamination in acute pancreatitis, liver cirrhosis (thus preventing spontaneous bacterial peritonitis) and nonsteroidal anti-inflammatory drug (NSAID) use (lessening in that way NSAID enteropathy). This antibiotic has, therefore, little value outside the enteric area and this will minimize both antimicrobial resistance and systemic adverse events. Indeed, the drug proved to be safe in all patient populations, including young children. Although rifaximin has stood the test

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of time, it still attracts the attention of both basic scientists and clinicians. As a matter of fact, with the advancement of the knowledge on microbial-gut interactions in health and disease novel indications and new drug regimens are being explored. Besides widening the clinical use, the research on rifaximin is also focused on the synthesis of new derivatives and on the development of original formulations designed to expand the spectrum of its clinical use.

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Introduction

Hundreds of bacterial species make up the human gut flora. The intestine has at least 400 different species of bacteria totaling over 1,012 organisms. Of these, 99% are anaerobic bacteria. Although anaerobes are part of the normal commensal flora, they can become opportunistic pathogens, causing serious, sometimes fatal infections if they escape from the colonic milieu. Most often, this escape occurs as a result of perforation, surgery, diverticulitis or cancer [1]. Pathogens range from highly virulent organisms, which infect people with well-functioning immune systems as well as people with poorly functioning immune systems, to opportunistic organisms, which infect only those with impaired immune systems (e.g. HIV-infected patients or transplant and oncology patients taking immunosuppressive drugs) [2]. In these subjects infection can be particularly severe, debilitating, and difficult to treat.

The host gastrointestinal (GI) tract is exposed to countless numbers of foreign antigens and has embedded a unique and complex network of immunological and non-immunological mechanisms, often termed the GI 'mucosal barrier', to protect the host from potentially harmful pathogens while at the same time 'tolerating' other resident microbes to allow absorption and utilization of nutrients. Of the many important roles of this barrier, it is the distinct responsibility of the mucosal immune system to sample and discriminate between harmful and beneficial antigens and to prevent entry of food-borne pathogens through the GI tract. This system comprises an immunological network termed the gut-associated lymphoid tissue (GALT) that consists of unique arrangements of B cells, T cells and phagocytes which sample luminal antigens through specialized epithelia termed the follicle-associated epithelia (FAE) and orchestrate coordinated molecular responses between immune cells and other components of the mucosal barrier [3].

Certain pathogens have developed ways to bypass and/or withstand defense by the mucosal immune system to establish disease in the host. Some 'opportunistic' pathogens (such as *Clostridium difficile*) take advantage of host or other factors (diet, stress, antibiotic use) which may alter or weaken the response of the immune system. Other pathogens have developed mechanisms for invading the GI epithelium and evading phagocytosis/destruction by immune system defenses [4]. Once cellular invasion occurs, host responses are activated to limit local mucosal damage and repel the foreign influence. Some pathogens (*Shigella* spp., parasites and viruses) primarily establish localized disease while others (*Salmonella*, *Yersinia*, *Listeria*) use the lymphatic system to enter organs or the bloodstream and cause more systemic illness. In some cases, pathogens (*Helicobacter pylori* and *Salmonella typhi*) colonize the GI tract or associated lymphoid structures for extended periods of time and these persistent pathogens may also be potential triggers for other chronic or inflammatory diseases, including inflammatory bowel disease (IBD) and malignancies [5]. The ability of certain pathogens to avoid or withstand the host's immune assault and/or utilize these host responses to their own advantage (i.e. enhance further colonization) will dictate the pathogen's success in promoting illness and furthering its own survival [4].

Emerging infectious pathogens, increasing antimicrobial resistance (mediated primarily through horizontal transfer of a plethora of mobile DNA transfer factors) and the appearance of diseases that decrease the host defense have increased the need for more effective and safe treatments [6]. Antibiotics have an important place in the management of GI diseases [7–9]. Antibiotic use in gastroenterology falls into three general settings [8]: (1) GI infections (e.g. bacterial diarrhea, cholangitis, diverticulitis), (2) GI diseases that may involve infectious agents but are not 'classic' infectious diseases (e.g. *H. pylori*-positive peptic ulcer, Whipple's disease, IBD), and (3) antibiotic prophylaxis for GI procedures.

The proliferation of antibacterial agents has made the choice of antibiotics increasingly complex. General considerations in selecting antibiotic therapy include (1) the identity and susceptibility pattern of the infecting organisms, (2) the anatomic localization of the infection, (3) the antimicrobial spectrum of the drug, and (4) its pharmacokinetic properties. Other important considerations include the possible selection of resistant organisms, interactions with other drugs, toxicity and cost [8].

The anatomic location of the GI infection influences the selection of the antimicrobial agent and the route of

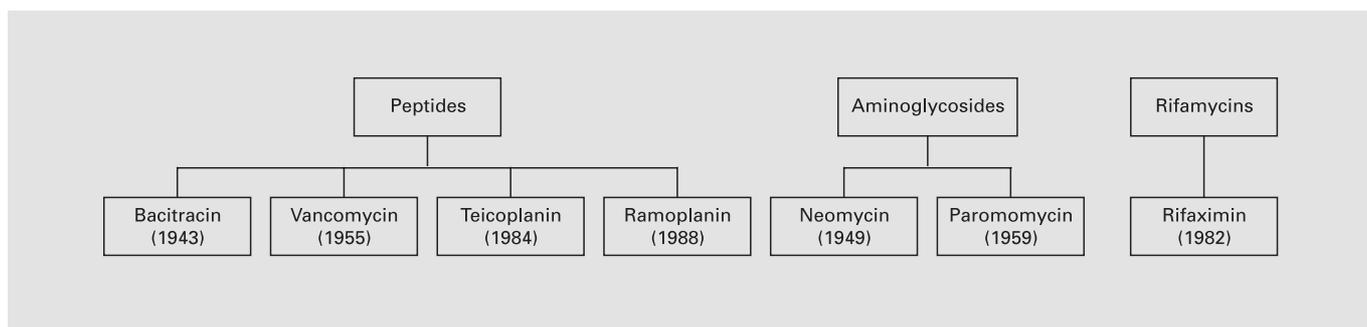


Fig. 1. Poorly absorbed antibiotics currently used in the treatment of GI infections. The date in parentheses refers to the first full description of the chemical synthesis of each compound.

administration. For instance, oral administration of a poorly absorbable antibiotic may be used for the eradication of noninvasive enteric pathogens [10]. Although the importance of attaining high biliary concentrations of antimicrobial agents in treating patients with cholangitis is still debated, it has been suggested that agents undergoing biliary secretion have a higher efficacy in the treatment of these infections [11].

It is well known that the dynamic bacterial community lining the gut exerts many physiological functions [12, 13]. These include metabolic activities that result in the salvage of energy and absorbable nutrients, important trophic effects on intestinal epithelia and on immune structure and function, and protection of the colonized host against invasion by alien microbes [13]. Oral administration of antibiotics can cause 'ecological' disturbances in the normal intestinal microflora [12]. Suppression of the normal microflora may lead to reduced colonization resistance with subsequent overgrowth of preexisting, naturally resistant microorganisms, such as yeasts and *C. difficile*. Although the incidence varies among antibiotics, the occurrence of pseudomembranous colitis has been associated with virtually every antibiotic [14]. New colonization by resistant potential pathogens may also occur and may spread within the body or to other patients and cause severe infections

Nonabsorbed oral antibiotic therapy, unlike systemically available antibiotics, allows localized enteric targeting of pathogens and is associated with a minimal risk of systemic toxicity or side effects [15]. Provided that nonabsorbed antibiotics are as effective as systemically absorbed drugs for the target illness, their safety and tolerability profiles may render them more appropriate for certain patient groups, such as young children, pregnant or lactating women, and the elderly, among whom side

effects are a particular concern. The restricted use of non-absorbed oral antibiotics only for enteric infections should also reduce the development of widespread resistance, a major limitation of current antibiotics for enteric infections [15].

Compared to systemic drugs, the number of poorly absorbed antimicrobials that would best target the GI tract is relatively small and almost completely limited to aminoglycosides (fig. 1). Indeed, oral vancomycin [16], teicoplanin [17], and bacitracin [18] are confined to the treatment of *C. difficile* infection [19–21]. Ramoplanin, a glycolipodepsipeptide antibiotic [22], is being developed for the treatment of *C. difficile*-associated diarrhea [23] and vancomycin-resistant enterococcal infection in high-risk patients [24]. Paromomycin and neomycin represent therefore the most widely used compounds [25, 26]. Neomycin is often associated with bacitracin, which is highly active against Gram-positive microorganisms, in order to extend its antibacterial activity. However, even poorly absorbed aminoglycosides are not completely devoid of untoward effects. Indeed, both ototoxicity [27–29] and nephrotoxicity [30] have been reported after oral neomycin especially in patients with renal dysfunction. Such patients can in fact accumulate toxic levels of the antimicrobial since the kidneys represent the major route of drug excretion [31]. Ototoxicity has actually been reported after ototopic (i.e. ear drops) aminoglycoside administration [32].

In order to overcome the limitations of the above drugs, a series of rifamycin derivatives with improved pharmacokinetic (i.e. virtually absence of GI absorption) and pharmacodynamic (i.e. with broad spectrum of antibacterial activity) properties have been synthesized at Alfa Wassermann laboratories [33]. Amongst the different molecules, the compound marked L-105 and later

named rifaximin was selected for further development. The antibiotic was first marketed in Italy and subsequently introduced in other European countries. Rifaximin was also licensed in some Northern African and Asian areas as well as in Mexico. The compound has recently been approved by the US FDA for the treatment of infectious diarrhea in the traveler (TD) [34].

The aim of this review is to summarize the available pharmacology and safety data on this nonsystemic antibiotic as well to outline its current and potential clinical use.

Rifaximin: Structure and Physicochemical Properties

Rifamycin is a clinically useful macrolide antibiotic produced by the Gram-positive bacterium *Amycolatopsis mediterranei* (originally classified as *Streptomyces mediterranei*). Rifamycin B, the compound originally isolated, has no antibacterial activity, but it is oxidized to the very active derivative rifamycin S, which inhibits the growth of Gram-positive bacteria. This antibiotic is primarily used against *Mycobacterium tuberculosis* and *Mycobacterium leprae*, causative agents of tuberculosis and leprosy, respectively. In these bacteria, rifamycin treatment specifically inhibits the initiation of RNA synthesis by binding to the β subunit of RNA polymerase. Apart from its activity against the bacteria, rifamycin has also been reported to inhibit reverse transcriptase (RT) of certain RNA viruses. Rifamycin derivatives have also been discovered that are effective against *Mycobacterium avium*, which is associated with the AIDS complex. Consequently, the importance of and demand for rifamycin have increased tremendously worldwide [35]. The rifamycin antibiotics, namely rifampicin (called rifampin in the US), rifabutin and rifapentine, are uniquely potent in the treatment of tuberculosis and chronic staphylococcal infections. Intestinal absorption of these drugs does occur and it is affected by the presence of food [36].

Rifaximin (4-deoxy-4'-methylpyrido[1',2'-1,2]imidazo[5,4-c]rifamycin SV, fig. 2) is a synthetic product designed to modify the parent compound, rifamycin, in order to achieve low GI absorption while retaining good antibacterial activity [37]. It is a rifamycin SV derivative, prepared by condensing 2-aminopyridine derivatives to 3-bromorifamycin S (fig. 3) [37–39]. This pyridoimidazo rifamycin SV derivative, which proved to be stable in gastric juice for 24 h, displays a zwitterionic nature at physiological pH [38].

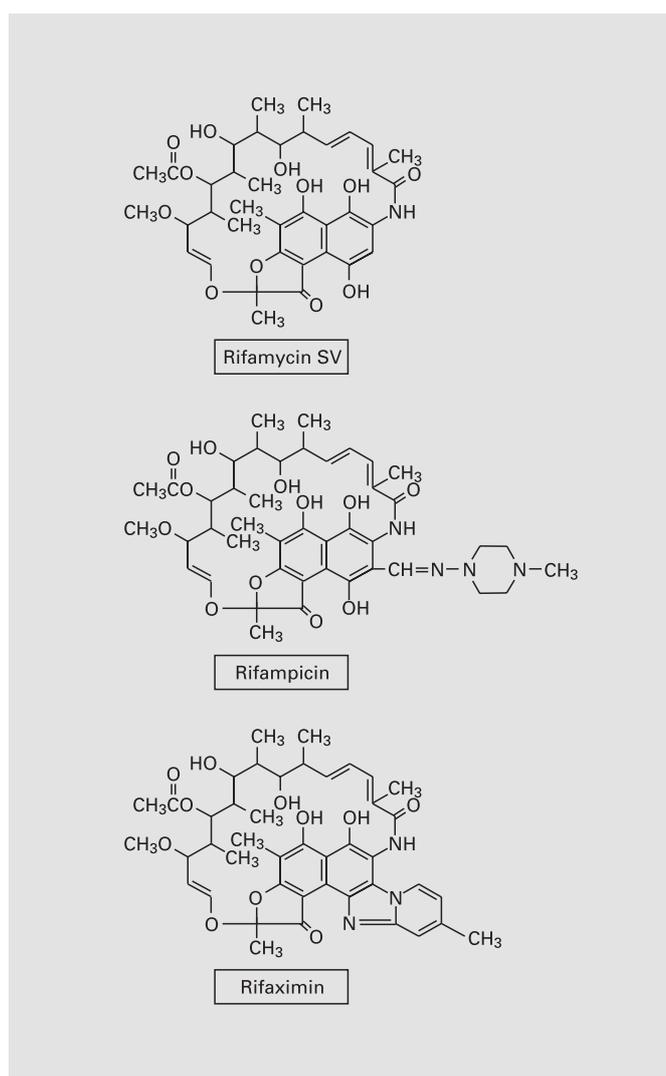


Fig. 2. Chemical structures of rifampicin and rifaximin as well as of their parent compound, rifamycin SV. The empirical formula of rifaximin is $C_{43}H_{51}N_3O_{11}$ and its molecular weight 785.9 daltons.

A solid-state X-ray study [40] did confirm the structure proposed on the basis of 1H -NMR studies in solution and showed that the compound is in a mesomeric betaine form, the pyrido nitrogen being positively charged and the imidazo nitrogen being negatively charged, a feature most likely responsible for the pharmacokinetic behavior of these new drugs. Indeed, since rifamycins are generally absorbed by passive diffusion, the presence of the two opposite charged nitrogens, together with the presence of the phenolic hydroxyls, leads to a molecule ionized at all the pH values encountered along the GI tract, which thus prevents its absorption. Rifaximin also displays a strong

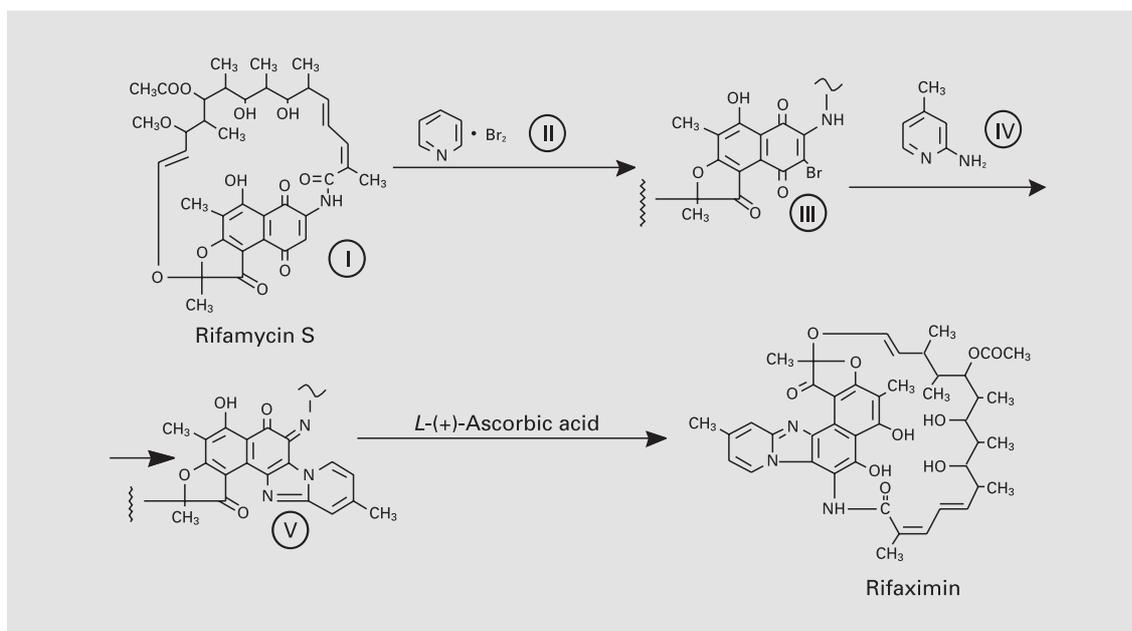


Fig. 3. Stepwise synthesis of rifaximin (from De Angelis [39]). The reaction of rifamycin S (I) with pyridine perbromide (II) in 2-propanol/chloroform (70/30) mixture at 0 °C gives 3-bromorifamycin S (III), which is then condensed with 2-amino-4-methyl-pyridine (IV) at 10 °C. The *o*-quinonimic compound (V) is then obtained. This compound is finally reduced with ascorbic acid to rifaximin.

tendency to self-associate both in solution and in the solid state, and the increase in molecular size may also play a role in preventing its absorption [41].

Antimicrobial Activity

In vitro Activity

The *in vitro* antibacterial activity of rifaximin has been determined by using minimum inhibitory concentrations (MIC) against bacteria from clinical isolates or stock culture collections. It should be pointed out that – in the absence of known GI concentrations – the interpretation of MICs is difficult. It is likely, however, that the drug concentration achieved at the desired site of action, i.e. the GI tract, will largely exceed the reported MIC values. For instance, fecal levels after oral administration of the antibiotic range between 4,000 and 8,000 µg/g of stool, which is 160–250 times higher than the MIC₉₀ for the various bacterial enteropathogens [42].

Several *in vitro* studies, summarized by Gillis and Brogden [33] and Jiang and DuPont [43], have shown that – like rifampicin – rifaximin displays an inhibitory activity against Gram-positive and Gram-negative, aero-

bic and anaerobic bacteria. Sharing the properties of the rifamycin family, the drug was shown to be active against the H37RV *M. tuberculosis* [44] as well as five *Mycobacterium* isolates from patients with tuberculosis [45]. In general, the activity of rifaximin proved to be greater against Gram-positive than Gram-negative bacteria. Amongst Gram-positive microorganisms, the susceptibility of oxacillin-resistant strains of *Staphylococcus aureus* [46, 47] and of *C. difficile* [47, 48] is particularly interesting. Of the Gram-negative rods, identified in patients with GI disease, those belonging to the Enterobacteriaceae family appear to be the most sensitive ones [49, 50]. Interestingly enough, rifaximin was found to inhibit the growth of *H. pylori* [51–54] with MIC values (table 1) intermediate between those of amoxicillin and colloidal bismuth subcitrate [55]. In contrast to metronidazole, no strain tested exhibited primary resistance [52, 53]. Furthermore, the activity of rifaximin was only slightly affected by lowering the pH of the medium, conversely from what is currently observed with other antibiotics [56]. Last but not least, rifaximin was shown to be active against clarithromycin-resistant strains [54].

Finally, a recent study [57] found rifaximin active against 408 clinical strains of *Vibrio cholerae* isolated

Table 1. MICs ($\mu\text{g/ml}$) of rifaximin and other antimicrobials against 40 strains (39 clinical isolates and the NCTC 11638 strain) of *H. pylori* at pH 7.0 (from Holton et al. [53])

	MIC ₅₀	MIC ₉₀	Range
Rifaximin	4	8	4–6
Ampicillin	0.03	0.25	0.03–0.5
Metronidazole	0.5	4	0.12–4
Omeprazole	32	>128	32->128

from different geographical areas and with different antimicrobial resistance patterns, the MIC values ranging from 0.5 to 4 $\mu\text{g/ml}$ for all strains. These findings, together with the pharmacokinetic properties of the drug, suggest that rifaximin could be an attractive antimicrobial agent for cholera.

Mechanism of Action

To establish whether rifaximin, like the other members of the rifamycin family [36, 58], specifically inhibits bacterial RNA synthesis the effect of this antibiotic as well as that of rifampicin and chloramphenicol on RNA (via ³H-uridine incorporation), DNA (via ³H-thymidine incorporation) and protein (via ³⁵S-methionine incorporation) synthesis was studied in growing cultures of *Escherichia coli* [59]. While chloramphenicol reduced protein synthesis, both rifaximin and rifampicin inhibited RNA synthesis in a concentration-dependent fashion. In contrast, none of them affected ³H-thymidine incorporation into DNA. These data suggest that rifaximin, like rifampicin, inhibits RNA synthesis by binding the β subunit of the bacterial DNA-dependent RNA polymerase [60].

In vitro Antimicrobial Resistance

Despite much effort, antibiotic resistance continues to increase [61]. Looking back, it is clear that this was an inevitable consequence of antibiotic use [62]. Antibiotic resistance, which has been recognized to be an important clinical problem, varies in prevalence from one country to another and among the pathogens themselves. This has great clinical, economic, political and environmental implications worldwide [63]. Strict adherence to the ongoing measures of infection control, education and antibiotic policy should minimize antibiotic resistance [64].

Development of resistance to rifaximin may be similar to that of rifampicin, which is primarily due to a chromosomal single-step alteration in the drug target, the DNA-dependent RNA polymerase [65, 66]. This differs from

the plasmid-mediated resistance commonly acquired by bacteria to aminoglycoside antibiotics, such as neomycin or bacitracin [67]. The spread of resistance due to the chromosomal mechanism is less frequent than that due to plasmid-mediated transfer [63, 66].

The development of resistance to rifaximin was studied in detail on several aerobic (Gram-negative and Gram-positive) and anaerobic strains using two different methods, i.e. the agar dilution and broth dilution methods [68, 69]. In the agar dilution method bacterial inocula were spread onto plates containing antibiotics at concentrations 2, 4 or 8 times the MIC value. After incubation under aerobic and anaerobic conditions for the appropriate species of bacteria, surviving colonies were counted, purified and the frequency of spontaneous resistant mutants was evaluated. Incubation in an anaerobic atmosphere was used to mimic the situation of the human GI tract environment (prevalently anaerobic). In the broth dilution method bacteria were inoculated in test tubes containing subinhibitory concentrations of the drug tested and incubated at 37°C. Aliquots of these cultures were transferred from tubes containing the highest drug concentration that permitted bacterial growth to another series of tubes containing 2-, 4-, 8- and 16-fold the concentration of the antimicrobial agent. The experiments were ended when the test bacteria were able to grow in media containing at least 100 $\mu\text{g/ml}$ of the drug tested. Thanks to the well-known ability of suboptimal concentrations of antibiotics to promote this phenomenon [70] it is obviously easier to select resistant mutants with this second method, where exposure to subinhibitory concentrations of antimicrobials is allowed.

As expected [71], spontaneous selection of resistance was rare for the anaerobic bacteria; in fact, among these anaerobes only a few species showed spontaneous emergence of resistant mutants [46, 48].

The selection of rifaximin-resistant strains was also investigated on five different isolates of *H. pylori*. None of the strains exhibited primary resistance to rifaximin, but, after exposure to subinhibitory concentrations of the antibiotic, all five strains became resistant. The mutation frequency was similar to that observed with macrolides and quinolones, but was less frequent than that observed with metronidazole [52, 53].

Rifaximin selected resistant Gram-positive cocci mutants more easily under aerobic conditions than in an anaerobic atmosphere [46, 48]. In comparison with Gram-positive microorganisms, drug-resistant Gram-negative bacilli were rarely detected [46, 48].

Table 2. Therapeutic activity of rifaximin and other antibiotics on experimentally induced infection in mice (from Venturini and Marchi [74])

Route of administration	ED ₅₀ , mg/kg		
	rifaximin	rifampicin	gentamycin
Oral	>10	0.15	>10
Subcutaneous	0.46	ND	ND

Swiss mice were infected by intraperitoneal injection of 0.25 ml of *S. aureus* Colliva (clinical isolate). The bacterial suspension contained a sufficient number of organisms to kill all the animals within 72 h. Antibiotics were given orally or subcutaneously 1 h after bacterial inoculation. ND = Not determined.

Table 3. Effect of rifaximin and rifampicin on experimentally induced tuberculosis in guinea pigs (from Lucchesi et al. [76])

Treatment	Feldman's index	Pathological findings
Saline	54.11	extensive tuberculosis
Rifaximin (60 mg/kg daily)	54.47	extensive tuberculosis
Rifampicin (30 mg/kg daily)	11.00	very limited infection

Albino male guinea pigs were infected by subcutaneous injection of 0.001 mg of bacterial (*M. tuberculosis*) coating and treated with rifamycins by oral route immediately after the infection. The antibiotic treatment lasted 4 months, after which the animals were sacrificed and inguinal lymph nodes, spleen, liver and lungs examined to quantify the severity of the disease according to Feldman [75].

Spontaneously resistant mutants were more easily selected after preincubation of the test bacteria with subinhibitory concentrations of rifaximin rather than after exposing the microorganisms to high levels of the antibiotic. Taking into account that rifaximin is poorly absorbed (see below) the high amounts of the drug available within the digestive lumen compare better with supra- rather than with subinhibitory concentrations of the drug. Furthermore, since the anaerobic atmosphere did hinder the selection of rifaximin-resistant enterobacteria, it is expected that – during antibiotic therapy with this drug – the selection of resistant mutants in the GI tract (a prevalently anaerobic environment) is very low. In summary, thanks to the high drug bioavailability in an oxygen-deficient milieu, the in vivo occurrence of bacterial resistance should with rifaximin be an infrequent phenomenon. The

constant therapeutic efficacy of the antibiotic in the management of various GI infections [33, 71] clearly suggests that this is the case.

In vivo Activity

The antibacterial activity of rifaximin has been confirmed by in vivo studies performed in both laboratory animals and humans.

Two studies compared the effect of rifaximin to that of neomycin and/or of rifampicin [72, 73] on the fecal flora in rats. In the first study [72] the antibiotic (1, 10, 30 and 100 mg/kg orally for 7 days) did inhibit both aerobic (especially coliforms and cocci) and anaerobic bacterial growth. Its activity was similar to that of neomycin and stronger than that of rifampicin. In the second investigation [73] the antibiotic effect on aerobic microorganisms was specifically investigated. Oral rifaximin treatment (50 mg/kg for 3 days) caused a marked reduction in the number of total aerobic bacteria and salmonellae, while neomycin led only to a decrease in salmonella counts, but did not cause statistically significant changes in the total aerobic bacterial population.

The in vivo protective activity of rifaximin was studied in mice, infected experimentally by intraperitoneal inoculation of *S. aureus* Colliva and compared to that of rifampicin (a systemic rifamycin) and gentamicin (a poorly absorbed aminoglycoside) [74]. After oral administration, only rifampicin was effective whereas the other two compounds were inactive at doses up to 10 mg/kg. However, when injected subcutaneously, rifaximin displayed a good therapeutic efficacy (table 2). While confirming its antibacterial activity, these results clearly indicate that rifaximin, like gentamycin, is poorly absorbed after oral administration.

The lack of absorption and, consequently, the lack of 'systemic' therapeutic activity was also demonstrated in a model of experimental tuberculosis in the guinea pig [75]. Here again, oral rifampicin but not oral rifaximin did protect the animals from the development of the infection (table 3) [76].

A large number of human studies [71, 77–80] performed in patients with infectious diarrhea or other GI diseases (e.g. hepatic encephalopathy, small bowel bacterial overgrowth, IBD, colonic diverticular disease) have confirmed the antibacterial activity of rifaximin demonstrated in vitro and in laboratory animals.

In approximately 50% of patients with infectious diarrhea enrolled in clinical trials the most common organism isolated and presumed causative was *E. coli*. Treatment with the antibiotic led to clearance of the bacterium in

Table 4. Changes in fecal bacterial population after oral rifaximin administration in healthy volunteers (from Testa et al. [81])

Organisms		Weeks					
		0	1	2	4	8	12
<i>E. coli</i>	× 10 ⁸	2.9	0.46	2.5	2.7	3.0	3.0
Other enterobacters	× 10 ⁷	1.0	0.09	1.0	1.2	1.1	1.2
Enterococci	× 10 ⁷	5.6	0.08	3.1	5.7	5.6	4.9
<i>Bacteroides</i> spp.	× 10 ⁹	6.0	0.10	5.4	6.1	6.2	5.6
<i>Clostridium</i> spp.	× 10 ⁸	1.1	0.04	1.0	1.1	1.0	0.9
Anaerobic cocci	× 10 ⁷	6.1	0.02	5.6	6.0	6.2	5.8

Rifaximin (800 mg) was given in two daily doses for 5 days after the first stool collection.

65–80% of the patients [33, 71]. Eradication of or decrease in microorganisms count in the feces was also reported for other Enterobacteriaceae after short-term rifaximin treatment [33, 71].

On the basis of log bacterial survival rates, the antibacterial activity of rifaximin was greater than that of paromomycin against *Enterococcus* spp., anaerobic cocci, *Bacteroides* spp. and *Clostridium* spp. isolated in fecal samples from 20 patients with subclinical hepatic encephalopathy (fig. 4) [81]. On the other hand, *E. coli* and *Klebsiella* spp. appeared more susceptible to paromomycin while both antibiotics showed equal potency against *Proteus* spp. [81]. Here again it should be pointed out that stool concentrations of rifaximin are 250–500 times higher than the MIC₉₀ values [71], which makes the in vitro differences of activity between this and other antimicrobials meaningless from a clinical standpoint.

Repeated oral administration of an antibiotic that reaches very high concentrations within the GI lumen could have profound effects on intestinal flora [12, 13]. As expected, rifaximin markedly reduced fecal bacterial counts during oral intake but the effect was short-lasting since the bacterial population recovered within 1–2 weeks from the end of treatment (table 4) [82]. Most importantly, fungal colonization occurred very rarely. Indeed, *Candida albicans*, which has been implicated in the pathogenesis of antibiotic-associated diarrhea [82, 83], was isolated from the fecal samples of only 2 out of 10 patients given 1,200 mg of rifaximin daily [81] and in none of the volunteers taking 800 mg daily [82].

It is worthwhile mentioning that treatment with high-dose (600 mg, 3 times a day, for 14 days) rifaximin was also efficacious in resolving the clinical symptoms and clearing protozoan infections in HIV-1-infected patients with a CD4 count ≥200/mm³, who presented enteric and sys-

temic symptoms due to *Cryptosporidium parvum* or *Blas-tocystis hominis* associated with enteropathogens [84].

In vivo Antimicrobial Resistance

Antimicrobial resistance to rifamycins develops rapidly both in vitro and in vivo [65, 85, 86]. As a consequence, all the three members of the family (i.e. rifampicin, rifabutin and rifapentine) are used clinically as components of combination therapies [65, 87]. Being structurally related, rifaximin could share this potential. And indeed resistance rates, recorded in fecal strains of Enterobacteriaceae, *Enterococcus*, *Bacteroides*, *Clostridium* and anaerobic cocci, ranged between 30 and 90% after short-term (5 days) antibiotic (800 mg daily) treatment [82]. A similar pattern was observed in 10 patients with hepatic encephalopathy after treatment with rifaximin 1,200 mg/day for 5 days [80].

Nevertheless, a rapid disappearance of resistant bacteria was observed after stopping the antibiotic treatment (fig. 5). Different kinetics of disappearance were, however, observed. The aerobic species showed a more rapid return to the baseline 'sensitive' status whereas the anaerobic bacteria, especially the Gram-negative rods, regained sensitivity to rifaximin more slowly. In any case, 3 months after the end of treatment resistant strains were no longer detectable in the feces [82]. These results support the cyclic use of rifaximin that has been adopted by the investigators in the treatment of hepatic encephalopathy [77] and colonic diverticular disease [79].

However, DuPont and Jiang [88], by studying changes in the susceptibility of intestinal flora during a 3-day rifaximin course among US students with TD, failed to document the emergence of drug-resistant Gram-positive (e.g. enterococci) and Gram-negative (*E. coli*) organisms during treatment.

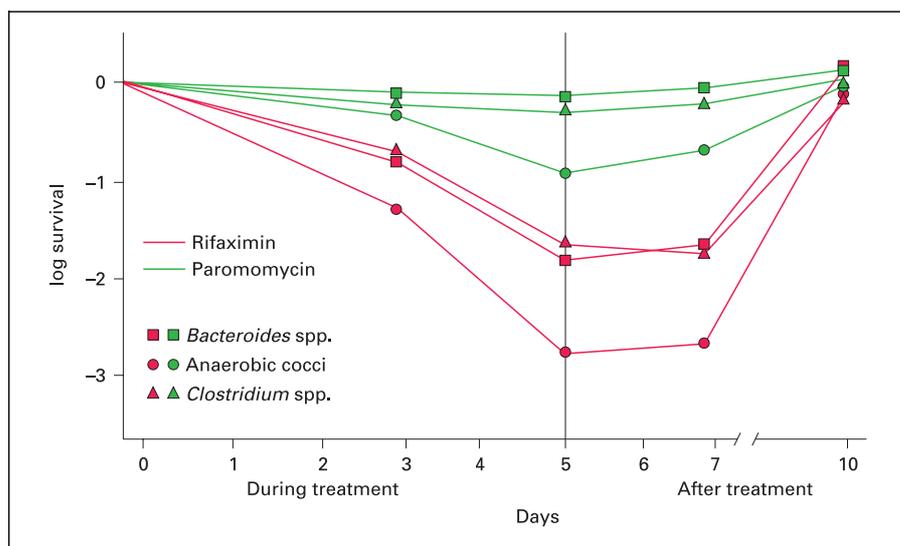


Fig. 4. Changes in anaerobic flora population following oral administration of rifaximin or paromomycin in patients with hepatic encephalopathy (from Testa et al. [81]).

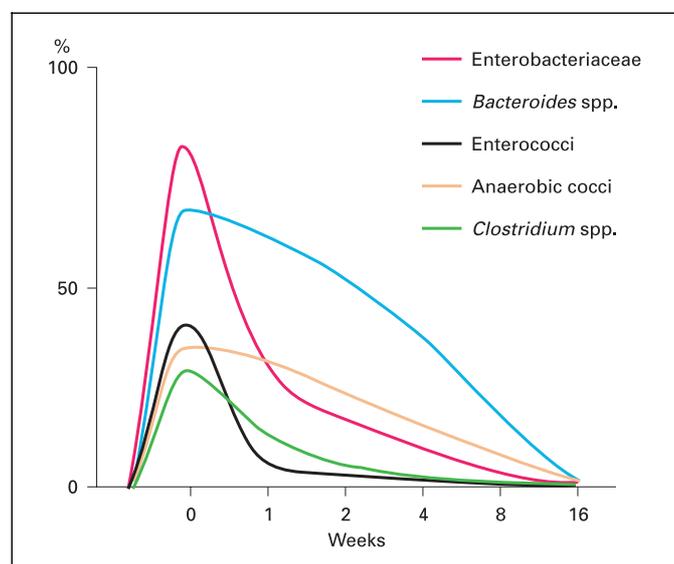


Fig. 5. Disappearance of rifaximin-resistant bacteria from the human intestine after stopping the antibiotic treatment (week 0) (from De Leo et al. [82]).

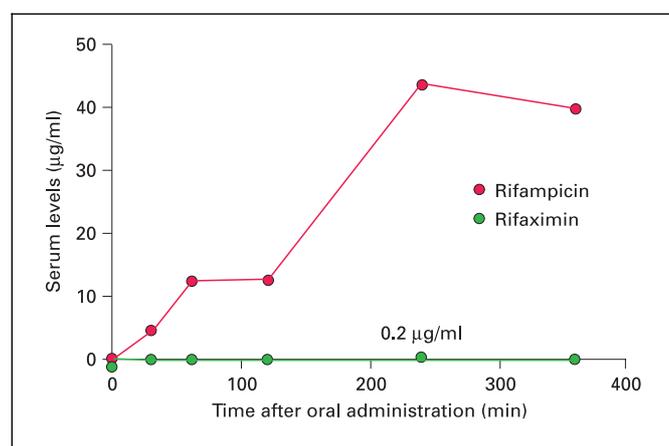


Fig. 6. Drug serum levels after oral administration of rifampicin (100 mg/kg) or rifaximin (100 mg/kg) to fed rats (from data in Venturini [97]).

Since rifamycins are important drugs for the treatment of *M. tuberculosis* infection [36, 86, 87] the activity of rifaximin on and interference with this bacterium have been carefully studied. Indeed, a potential problem of the treatment with this antibiotic is represented by the possibility that even very low blood levels achieved by oral administration might be able to select mutants, cross-resistant to rifamycins [85], in patients treated for GI infections and harboring *M. tuberculosis*.

A study in the guinea pig with experimentally induced tuberculosis [44] did show that sensitivity to both rifampicin and rifaximin of the *Mycobacterium* (H37RV strain) remained unchanged after 3 months of drug administration, the MIC values being virtually the same before and after treatment. The development of resistance to rifaximin of different strains of *M. tuberculosis*, isolated from patients with pulmonary and renal tuberculosis, was studied under extremely stringent conditions [45].

The drug was tested at very high concentrations (up to 270 ng/ml), largely exceeding those expected from intestinal absorption during an average course of therapy for enteric infections. These experiments showed a concentration-dependent subinhibitory effect of the antibiotic on *Mycobacterium* growth. One can, therefore, speculate that the circulating, albeit very low, levels of rifaximin could represent a stimulus for the selection of resistant mycobacterial strains [89]. However, measurement of both rifaximin and rifampicin MICs before and after successive exposure to rifaximin gave the same value in all the *Mycobacterium* strains examined [45].

All these data, taken together, do suggest that therapy with rifaximin in GI patients harboring *M. tuberculosis* should not represent a hazard to the treatment of pulmonary as well as extrapulmonary tuberculosis. A recent 10-year survey [90] in Italy showed that – in the period 1990–2000, during which rifaximin has been largely used – mycobacterial resistance to rifampicin has been quite stable. This finding is consistent with the idea that the likelihood of rifampicin resistance emerging in infected, but otherwise asymptomatic, individuals receiving this antibiotic is extremely low.

Rifampicin is currently employed prophylactically to eliminate pharyngeal carriage of *Neisseria meningitidis*, thereby lowering the potential risk of meningitis [91–93]. The selection of resistant *Neisseria* mutants could theoretically be the consequence of rifaximin use for GI infections. Here again, a recent study performed in Italy [94] found that all the meningococci isolated from asymptomatic carriers were susceptible to rifampicin, thus ruling out this possibility.

Pharmacokinetics and Drug Interactions

Animal Pharmacokinetics

The pharmacokinetic investigations on rifaximin were performed in the animal species (namely the rat and dog) in which the short- and long-term toxicological studies had been performed according to the OECD (Organization for Economic Co-Operation and Development) guidelines. Experimental studies were performed by using unlabeled and labeled rifaximin that was purportedly synthesized [95] to allow the quantitation of minute amounts of the antibiotic within the body. Rifaximin was given to fed and fasted animals in order to ascertain whether the GI absorption, if any, of rifaximin could be modified by food intake. Rifampicin, another systemic rifamycin derivative, whose pharmacokinetic properties

are well known [36, 96], was used as a reference compound.

The first study was performed by Venturini [97, 98] in both rats and dogs by using a microbiological assay (i.e. agar diffusion test and *S. aureus* 209 P FDA as test organism). Conversely from rifampicin, whose serum levels were already detectable 30 min after the administration and still measurable after 48 h, only trace amounts (i.e. 0.2 µg/ml) of rifaximin were detected in serum of fed rats 4 h later (fig. 6). The amount of detectable antibiotic was reduced by 50% in fasted animals. Similar results have been obtained in dogs after oral administration of 25 mg/kg of both rifamycin derivatives [97, 98]. No detectable amount of rifaximin was found in serum at any time.

The negligible intestinal absorption of rifaximin was subsequently confirmed with the use of the labeled drug. After oral administration in rats the total radioactivity present in plasma was found to be no more than 0.1 and 2% of the administered dose after the intake of ³H-labeled [99] and ¹⁴C-labeled [59] compound, respectively.

Although theoretically safe, poorly absorbed antimicrobials could become ‘absorbable’ in the presence of mucosal inflammatory or ulcerative changes [100], like those occurring in IBD or when invasive bacteria colonize the intestine. To verify whether the presence of intestinal lesions would affect rifaximin absorption, the drug was given to rats with experimentally induced colitis [101]. The indomethacin-induced enteropathy did not affect intestinal absorption of rifaximin. However, under the same experimental conditions, systemic bioavailability of neomycin did increase [101].

The labeled rifaximin molecules were also employed to better evaluate the fecal and urinary excretion of the antibiotic, originally studied with the unlabeled drug [97]. The major route of excretion was in feces with all rats excreting more than 96 or 86% of the dose of the ³H-labeled [99] and ¹⁴C-labeled [59] rifaximin, respectively. In contrast, urinary recovery was found to be very low, i.e. less than 1%, thus confirming the insignificant systemic absorption of the drug. A small, sex-dependent recovery of radioactivity (1.72% in male and 0.5% in female) was recorded in bile [59], thereby proving evidence of an, albeit negligible, enterohepatic circulation of the drug. When the fecal and urinary counts were added to the radioactivity found after carcass digestion, the total radioactivity recovered over 7 days ranged between 95 and 100% [59, 99]. Similar findings have been reported in dogs [59].

The studies with radiolabeled rifaximin also showed that the greatest concentration of radioactivity is found in

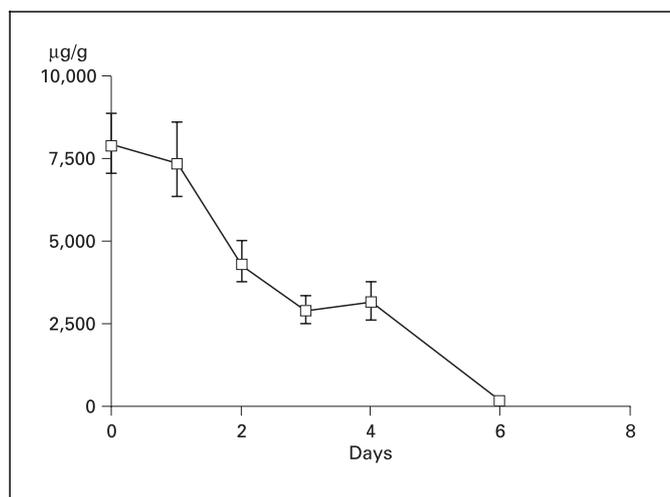


Fig. 7. Fecal concentration of rifaximin in patients with TD after treatment with 800 mg/day of the drug for 3 consecutive days. Each square refers to the mean of the values obtained from 39 subjects. Vertical bars are standard errors (from Jiang et al. [42]).

Table 5. Effect of food on the mean pharmacokinetic parameters following administration of a single rifaximin dose (400 mg) in 14 healthy volunteers (from FDA rifaximin label [34])

Parameter	Fasting	Fed
C_{max} , ng/ml	3.8 ± 1.32	9.63 ± 5.93
T_{max} , h	1.21 ± 0.47	1.90 ± 1.52
Half-life, h	5.85 ± 4.34	5.95 ± 1.88
AUC, ng·h/ml	18.35 ± 9.48	34.70 ± 9.23
Excreted in urine, %	0.023 ± 0.009	0.051 ± 0.017

* Each value represents the mean \pm SD.

the GI tract [59, 99] that represents the therapeutic target organ. The radioactivity peak was reached at 0.5 h in the stomach, at 2 h in the small intestine and at 7 h in the cecum and large intestine. Other than in the GI tract, radioactivity counts were generally low and only the liver and kidneys contained more than 0.01% of the dose administered, a finding consistent with the results obtained with unlabeled rifaximin measured by a microbiological assay [97].

Specific studies directed to elucidate the metabolism of rifaximin have not been done since the compound does not reach the systemic circulation. Theoretically, it may

be assumed that the small amount of circulating rifaximin is metabolized like the other rifamycin derivatives [36, 96], that is via the liver.

Human Pharmacokinetics

The pharmacokinetics of rifaximin after oral administration has been studied in healthy volunteers and patients with intestinal infections or IBD. The aim of these studies was to confirm the low, if any, systemic absorption of the drug; metabolism and excretion data are scant. In all these investigations a sensitive high-pressure liquid chromatographic (HPLC) method was used to measure rifaximin in body fluids.

After oral administration of 400 mg of rifaximin to fasted healthy volunteers blood drug concentration was found to be lower than the detection limit of the analytical method (i.e. 2.5 ng/ml) in half of them [102]. In the remaining subjects very low amounts were detected at some of the time intervals during the first 4 h after intake. Along the same lines, the urinary concentrations of the drug were very low and often undetectable. The effect of food on the absorption of the antibiotic was also evaluated [34] and a significant, albeit not clinically relevant, increase of bioavailability was observed after a high-fat breakfast (table 5).

^{14}C -labeled rifaximin was administered as a single dose to 4 healthy male subjects [34]. The mean overall recovery of radioactivity in the urine and feces of 3 subjects during the 168 h after administration was $96.94 \pm 5.64\%$ of the dose. Radioactivity was excreted almost exclusively in the feces with only a small proportion of the dose (i.e. 0.32%) excreted in urine. Analysis of fecal extracts indicated that rifaximin was being excreted as unchanged drug [34]. The amount of radioactivity in urine (<0.4% of the dose) is consistent with animal data and shows that rifaximin is poorly absorbed from the GI tract and is almost exclusively and completely excreted in feces as unchanged drug.

Systemic absorption of rifaximin (200 mg 3 times daily) was also evaluated in 13 patients with shigellosis on days 1 and 3 of a 3-day course of treatment [34]. Rifaximin plasma concentrations were low and variable. There was no evidence of drug accumulation following repeated administration for 3 days (9 doses). Peak plasma rifaximin concentrations after 3 and 9 consecutive doses ranged from 0.81 to 3.4 ng/ml on day 1 and 0.68 to 2.26 ng/ml on day 3 [34]. Fecal excretion of the drug was also assessed in 39 patients with acute diarrhea after administration of 400 mg every 12 h for 3 consecutive days [42]. As shown in figure 7, post-therapy stool rifaximin levels were high, decreasing gradually over a 5-day period. It is

Table 6. Plasma concentrations and urinary excretion of rifaximin in patients with UC given 400 mg of the drug (from Rizzello et al. [103])

Patient No.	Plasma concentrations, ng/ml					Urinary excretion							
	0 h	1 h	4 h	8 h	24 h	0–4 h		4–8 h		8–24 h		total amount	
						urine ml	rifaximin ng·ml ⁻¹	urine ml	rifaximin ng·ml ⁻¹	urine ml	rifaximin ng·ml ⁻¹	ng	total excretion, %
1	–	–	–	–	–	600	48.4	450	10.9	800	13.0	44,322	0.011
2						750	31.7	600	15.4	950	7.8	40,412	0.010
3	–	–	–	22.4	–	300	38.5	450	32.5	900	20.4	44,485	0.011
4				2.38		150	46.3	400	62.1	850	8.8	39,215	0.010
5	–	4.64	4.63	4.39	–	200	–	150	42.7	1,000	11.1	17,525	0.004
6			3.03			100	412.0	50	66.8	800	12.3	54,409	0.014
7	–	2.00	–	–	–	180	33.6	200	12.2	750	6.7	13,483	0.003
8	–	–	–	–	–	100	277.4	0	–	600	8.0	32,518	0.008
9				13.4		700	3.8	200	12.6	1,000	43.2	48,306	0.012
10	–	–	BLQ	–	–	200	6.5	200	8.5	450	4.6	5,053	0.001
11	–	4.38	BLQ	BLQ	–	600	59.9	100	44.0	1,700	26.6	85,534	0.021
12				BLQ		700	5.2	600	5.6	1,200	5.3	13,350	0.003
Mean												36,551	0.009
SD												22,244	0.006

BLQ = Below the limit of quantitation; – = not detected.

worthwhile mentioning that fecal drug concentrations largely exceeded the MIC values for the bacterial isolates obtained from patients with TD [42].

IBD being one of the therapeutic indications of rifaximin, its absorption was carefully studied in patients with mild-to-moderate ulcerative colitis (UC), after the administration of two tablets (i.e. 400 mg) orally [103]. As shown in table 6, in most plasma samples rifaximin concentrations were below the detection limits. Only in few patients was the drug detected during the first 8 h after administration. The total rifaximin amount recovered in the urine was only 0.009% of the dose. This figure fits well with the corresponding value (i.e. 0.007%) observed in healthy volunteers [102]. No correlation between disease activity and urinary concentrations was found after repeated drug administration [104]. It is worthwhile mentioning that, even after 15 days of therapy of patients with resistant pouchitis with high dose (2 g daily) of rifaximin, together with ciprofloxacin (1 g daily), no plasma level of the antibiotic was detectable in any patient [105].

Since animal data indicated the presence of an enterohepatic circulation [59], the biliary concentration of rifaximin after oral preoperative administration of the drug (400 mg every 12 h) was evaluated in bile samples taken from patients undergoing cholecystectomy [106]. In 7 out

of 13 subjects only trace amounts of the antibiotic were detected while the remaining ones had biliary concentrations ranging from 4.5 to 15.6 µg/ml. These figures are extremely low especially if compared to the biliary rifampicin concentrations (i.e. >150 µg/ml) observed after a single 450-mg oral dose [107]. Taking into account the bile volume, the highest biliary excretion was less than 0.2% [106].

Although the pharmacokinetics of rifaximin in patients with renal insufficiency has not been specifically studied, its very low renal excretion makes any dose adjustment unnecessary. The same holds true for patients with hepatic insufficiency. In fact, the mean peak drug plasma concentrations (i.e. 13.5 ng/ml) detected in subjects with hepatic encephalopathy patients given rifaximin 800 mg 3 times daily for 7 days [34, 108] were not dissimilar to those found in healthy subjects [102] and patients with IBD [98]. Indeed, in all the trials performed in this condition the drug has been well tolerated [33, 77].

Finally, drug absorption and excretion have not been evaluated in pediatric or geriatric populations. However, here again the tolerability of rifaximin in childhood and in the elderly has found to be extremely good [33].

Drug-to-Drug Interactions

As new classes of antimicrobial drugs have become available, pharmacokinetic drug interactions with antimicrobials have become more common. Macrolides, fluoroquinolones, rifamycins, azoles and other agents can interact adversely with commonly used drugs, usually by altering their hepatic metabolism [109]. The mechanisms by which antimicrobial agents alter the biotransformation of other drugs are increasingly understood to reflect inhibition or induction of specific cytochrome P450 enzymes. Rifampicin and rifabutin induce several cytochromes P450, including CYP3A4, and can therefore enhance the metabolism of many other drugs [109].

By using *in vitro* preparations of human enzymes it is possible to predict those antibiotics that will adversely affect the metabolism of other drugs [110]. Such studies have shown that rifaximin, at concentrations ranging from 2 to 200 ng/ml, did not inhibit human hepatic cytochrome P450 isoenzymes: 1A2, 2A6, 2B6, 2C9, 2C19, 2D6, 2E1 and 3A4 [34]. In an *in vitro* hepatocyte induction model, rifaximin was shown to induce cytochrome P450 3A4 (CYP3A4) [34], an isoenzyme which rifampicin is known to induce [109].

Since rifampicin impairs the effectiveness of oral contraceptives (OCs) and pregnancies have been reported in women taking OCs and antibiotics [111], the interaction between an OC containing ethinyl estradiol and norgestimate and rifaximin was studied in 28 healthy female subjects given a short course of the drug [34]. Results of this study showed that the pharmacokinetics of single doses of ethinyl estradiol and norgestimate were not altered by concomitant antibiotic administration.

Midazolam and other benzodiazepines (e.g. alprazolam and triazolam) are selective substrates of CYP3A4 [110] and the concomitant administration of potent metabolic CYP3A4 inducers results in statistically significant pharmacokinetic changes and a consequent loss of therapeutic efficacy [112]. To evaluate the possible midazolam-rifaximin interaction an open-label, randomized, crossover trial was designed to assess the effect of oral rifaximin (200 mg thrice daily for 3 or 7 days) on the pharmacokinetics of a single dose of midazolam, administered either intravenously (2 mg) or orally (6 mg) [34]. No significant difference was observed in all the pharmacokinetic parameters of midazolam or its major metabolite, 1'-hydroxymidazolam, with or without simultaneous antibiotic therapy. These results, therefore, show that rifaximin does not significantly affect intestinal or hepatic CYP3A4 activity.

The lack of any significant *in vivo* interaction between rifaximin and human cytochrome P450 is also consistent with the absence of any significant induction of drug-metabolizing enzymes in the liver and the GI tract of rats given the antibiotic orally for 6 months [113]. When given for prevention or treatment of TD [70], rifaximin should, therefore, not affect the pharmacokinetics (and consequently pharmacodynamics) of other prophylactic drugs (e.g. antimalarials) [113].

General Pharmacology

The GI tract being the main therapeutic target of rifaximin, its potential effects on gastric secretion and GI motility have been investigated in rats and mice [59]. The antibiotic was given intraduodenally to pylorus-ligated rats (Shay rat) at doses ranging from 10 to 500 mg/kg, that is up to 50 times the therapeutic daily dose. No effect on pH and volume of gastric juice as well as on acid output and pepsin activity was observed.

Gastric emptying was studied in rats by means of a liquid meal labeled with phenol red [Scarpignato, unpubl. observations] while intestinal transit was evaluated in mice by means of the charcoal test meal [59]. Here again rifaximin was unable to influence either the emptying rate or intestinal motility. The drug could, however, be capable of correcting the GI motility derangement often observed in the presence of small intestinal bacterial overgrowth (SIBO) [114]. This is the case of patients with diabetes [115] or Crohn's disease [116], in whom the delayed intestinal transit, detected together with SIBO, was accelerated by a short-course treatment with rifaximin.

The effect of rifaximin on cardiovascular (CV) and respiratory systems was investigated in anesthetized rats and guinea pigs, respectively [59]. Rifaximin was given intraduodenally at doses up to 100 mg/kg and carotid pressure and flow as well as heart rate were continuously measured in rats while respiration amplitude and frequency were monitored in guinea pigs. The rifamycin derivative did not affect any of the measured parameters at any time after its administration.

Clinical pharmacological studies to specifically address the effect of rifaximin on GI or CV and respiratory functions have not been performed. However, while the most frequently reported, albeit few, adverse events associated with rifaximin administration were gastrointestinal in nature, no untoward reactions involving the CV or respiratory systems have been described [33, 117].

Clinical Use and Therapeutic Potential

Data from both experimental and clinical pharmacology clearly show that rifaximin is a nonsystemic antibiotic with a broad spectrum of antibacterial action covering Gram-positive and Gram-negative organisms, both aerobes and anaerobes. Being virtually nonabsorbed, its bioavailability within the GI tract is rather high with intraluminal and fecal drug concentrations that largely exceed the MIC values observed in vitro against a wide range of pathogenic organisms. The GI tract represents, therefore, the primary therapeutic target and GI infections the main indication [33, 71]. Since gut bacteria play a pathogenic role in several GI disorders (like for instance IBD or irritable bowel syndrome, IBS), the broad antimicrobial activity of rifaximin is also of value in these clinical conditions [33]. Thanks to the lack of transcutaneous absorption pointed out in both animal [118] and human [119] studies, its topical use in skin infections has also been investigated [120]. Finally, since the rifaximin spectrum of antibacterial action includes many organisms (e.g. *Bacteroides bivius-disiens*, *Gardnerella vaginalis*, *Haemophilus ducreyi*) causing genital infections [46], including *Trichomonas vaginalis* [46] and *Chlamydia trachomatis* [121], an attempt has been made to apply it locally in the treatment of bacterial vaginosis [120]. The growing list of therapeutic applications of rifaximin, for which there are published clinical studies, is shown in table 7. Amongst them, some are established indications for which the clinical trials so far performed have provided evidence for a substantial benefit of rifaximin. These include infectious diarrhea [71], hepatic encephalopathy [77], SIBO [78], colonic diverticular disease [79] and IBD [80]. For each of these indications a summary of scientific rationale and of available data, which the reader is referred to, has recently been published [71, 77–80]. In this section, therefore, only the potential clinical GI use of rifaximin is discussed in detail.

IBS and Chronic Constipation

A thoughtful review of Lin [122] has put forward the hypothesis that SIBO could represent a framework to understand IBS. The possibility that SIBO may explain bloating, which is present in 92% of patients with IBS, is supported by a greater total hydrogen excretion after lactulose ingestion, by a correlation between the pattern of bowel movement and the type of excreted gas, a prevalence of abnormal lactulose breath tests in 84% of IBS patients, and a 75% improvement of IBS symptoms after eradication of SIBO. Altered GI motility and sensation,

Table 7. Established and potential clinical indications for rifaximin

<i>Established indications</i>
Infectious diarrhea (including TD)
Hepatic encephalopathy
SIBO
IBD
Colonic diverticular disease
<i>Potential indications</i>
IBS and chronic constipation
<i>C. difficile</i> infection
Bowel preparation before colorectal surgery
<i>H. pylori</i> infection
Selective bowel decontamination in acute pancreatitis
Prevention of SBP in cirrhosis
Prevention of NSAID intestinal injury
Extra-GI indications
Skin infections
Bacterial vaginosis
Periodontal disease

changed activity of the central nervous system, and increased sympathetic drive and immune activation may be understood as consequences of the host response to SIBO [115]. To further support the suggestion that abnormal enteric flora may be a contributing cause of IBS, two recent studies have shown that metronidazole [123] and neomycin [124] are both able to cause a significant symptomatic improvement in this condition.

Besides constipation-related IBS, several studies have also suggested abnormalities of colonic bacterial composition in chronic idiopathic constipation [125]. Here again antibiotic treatment with vancomycin [126, 127], rovamycin (in combination with diphetarsona, an amebicidal agent) [128, 129] or erythromycin [130], which, however, displays a prokinetic activity [131, 132], proved to be capable of reversing long-lasting constipation. Furthermore, the efficacy in both clinical conditions of probiotics [133–135] lends further support to the pathogenic role of bowel flora.

Since remarkable symptomatic improvement can be achieved in most patients, antibiotic therapy is obviously the cornerstone of the treatment of SIBO [136]. Ideally, the choice of an antimicrobial agent should be based on in vitro susceptibility testing of the bacteria in the small bowel of the individual patient. However, because it is impractical to obtain this information in most cases, the choice of the antibiotic is largely empiric and based on results of published series involving small intestinal cultures [137]. Whereas most patients with SIBO have aro-

Table 8. Activity of rifaximin, vancomycin and metronidazole against 93 clinical isolates of *C. difficile* (from Marchese et al. [48])

Antibiotic	Range mg/l	MIC ₅₀ mg/l	MIC ₉₀ mg/l	Susceptible ^a %
Rifaximin	0.004–128	0.004	128	74.1
Vancomycin	0.25–4	1	2	100
Metronidazole	0.06–0.25	0.125	0.25	100

^a The breakpoint for susceptibility of rifaximin was assumed to be equal to that of rifampicin.

bic and anaerobic overgrowth, in others malabsorption has been associated with overgrowth of purely aerobic flora. Therefore, the most effective antibiotic regimens generally include one or more drugs with activity against aerobic and anaerobic bacteria. The majority of drugs which proved to be effective, with the exception of neomycin and rifaximin, are systemic antimicrobials [78, 136] whose adverse events may limit their usefulness, especially in the long term. Benefits of therapy must indeed be weighed against the risks of long-term antibiotic use, such as diarrhea, *C. difficile* colitis, patient intolerance and bacterial resistance. Thanks to its efficacy (the highest amongst the tested antimicrobials, [78]) and safety, rifaximin currently represents the best pharmacological option for SIBO and associated GI disorders. A randomized, placebo-controlled, double-blind trial on the effect of rifaximin on SIBO and symptom relief in IBS is ongoing in the US.

C. difficile Infection

Progress in defining new treatments for *C. difficile* infection has been hindered by the heterogeneous nature of hospital-acquired diarrhea, and in particular by whether colitis and/or pseudomembranous colitis is present in individual cases. Study groups have usually been poorly defined in this context, and given the spontaneous resolution of symptoms in a proportion of cases the true efficacy of treatment approaches often remains uncertain. Enthusiasm to explore new treatment possibilities for *C. difficile* has been largely fuelled by the apparently high relapse rate of conventional (metronidazole or vancomycin) treatment [138].

There is a consensus amongst published recommendations for the management of *C. difficile* infection [19–21]. The most important first step in the treatment is cessation of the precipitating agent, most commonly antibiotics, if this is deemed to be medically appropriate. In mild disease, this is often sufficient for full recovery. In more

severe disease, antimicrobial therapy directed against *C. difficile* is required. Oral metronidazole is recommended as the initial treatment of choice. Vancomycin, which has a comparable efficacy, represents the second-line therapy. Given the higher cost of oral vancomycin therapy and concern about the selection for vancomycin-resistant enterococci, metronidazole is preferred as the initial agent of choice. Vancomycin is appropriate for patients with contraindications or intolerance to metronidazole or for those who fail to respond to metronidazole. Other antibiotics such as glycopeptides [17, 22, 23] should be reserved for patients who cannot tolerate metronidazole or vancomycin or who are nonresponders [139, 140].

Rifaximin displays good antibacterial activity against *C. difficile* in vitro [47, 48], with MIC values lower than those of metronidazole and vancomycin (table 8). Furthermore, the microorganism showed a particularly low incidence of spontaneously resistant mutants ($<1 \cdot 10^{-9}$). The low incidence of resistant subpopulations selected by a concentration 8 times the MIC value suggests that the high levels of the drug that will be reached within the GI lumen may further prevent the selection of mutants [48]. Data from trials with antibiotics in TD and preoperative bowel decontamination suggest that poorly absorbed antimicrobials might have a decreased risk of causing antibiotic-associated diarrhea and *C. difficile*-associated disease [14]. In this connection rifaximin, which displays an intrinsic activity against this microorganism, should carry less of a risk.

A randomized open trial, performed in patients with *C. difficile* pseudomembranous colitis, compared rifaximin (200 mg 3 times daily) to vancomycin (500 mg 2 times daily) and found the two drugs similarly effective [141]. The clearance of bacterial toxins was, however, more rapid with vancomycin. Further large double-blind clinical studies are needed to better define the role of rifaximin in the treatment of *C. difficile* infection.

Table 9. Effect of short-term preoperative treatment with rifaximin or placebo on fecal bacterial counts in 118 patients submitted to colonic open surgery (from Gruttadauria et al. [148])

Bacterial counts	Placebo		Rifaximin	
	before	after	before	after
Aerobic, CFU × 10 ⁶	3.9 ± 0.7	3.4 ± 0.4	4.2 ± 0.53	1.58 ± 2.5*
Anaerobic, CFU × 10 ⁹	89.1 ± 15.3	78.2 ± 21.3	70.2 ± 12.1	14.5 ± 0.1*

Rifaximin (200 mg thrice daily) or placebo were given in a double-blind, double-dummy fashion for 3 consecutive days in addition to mechanical bowel preparation, which can account for the decrease, albeit not significant, of anaerobic flora in the placebo group.

* $p < 0.001$ versus the pretreatment value.

Bowel Preparation for Colorectal Surgery

Effective management of intra-abdominal infections requires a combination of preoperative mechanical bowel preparation (MBP), antibiotic prophylaxis and appropriate surgical technique [142]. However, a recent meta-analysis of randomized clinical trials [143] concluded that there is limited evidence in the literature to support the use of mechanical bowel preparation in patients undergoing elective colorectal surgery. Although drugs are not a substitute for attention to detail and meticulous surgical technique, the judicious use of antibiotic prophylaxis can decrease the overall risk of infection, especially following clean-contaminated and contaminated operations [144]. The benefit of administering antimicrobial agents as a prophylaxis against postoperative infections has long been debated. Because of the high density of bacteria in the large intestine and rectum, there appears to be general agreement on the need for prophylactic antibacterial treatment for surgical procedures involving this area [145–147]. It has been estimated that, without prophylaxis or with inappropriate prophylaxis, the postoperative complication rate following colorectal surgery can range from 30 to 60% compared with less than 10% with suitable antibiotic prophylaxis [146]. However, the choice of an ideal agent(s) and route of administration remains controversial. It should provide coverage of all likely pathogens, including aerobic and anaerobic organisms [142, 146]. Both oral and/or parenteral antibiotics have been used [for a review, see 147]. Two oral regimens are currently employed: (1) an aminoglycoside agent with erythromycin base and (2) an aminoglycoside agent with metronidazole. The regimen most often chosen in the United States is neomycin and erythromycin base. In Europe and Australia, however, physicians prefer kanamycin and metronidazole or neomycin and metronidazole [147].

Due to its pharmacokinetic and pharmacodynamic properties, rifaximin could be a suitable single oral agent for antibiotic prophylaxis in colorectal surgery. Indeed preoperative administration of 600 mg/daily [148] or 800 mg/daily [149] for 3 days significantly reduces fecal counts of both aerobic and anaerobic bacteria (table 9). In two studies rifaximin was compared to parenteral gentamycin [148] or to oral paromomycin [149] and proved to be equally effective in the prevention of infectious complications of colonic surgery. An additional trial [150] evaluated the effect of adding rifaximin to intravenous cefotaxime (3 g daily for 5 days) in the prevention of bacterial infections after major colic surgery. Compared to the cephalosporin alone, the antibiotic combination was more effective. In all studies, rifaximin was tolerated extremely well. Nasogastric administration of rifaximin suspension (800 mg/day) for 5 postoperative days after colon surgery was attempted at our university hospital [151] and appeared to be as effective as preoperative oral administration in preventing infectious complications.

In summary, 3-day preoperative treatment with rifaximin appears to be an effective antibiotic prophylaxis for patients submitted to colorectal surgery. However, this kind of prophylaxis should be compared with the convenience and efficacy of perioperative regimens [152, 153], such as those with an aminoglycoside or a cephalosporin, before gaining wider acceptance. Being perioperative antibiotics systemic drugs whose administration might be accompanied by adverse events, including life-threatening reactions [154], the use of a virtually unabsorbed antimicrobial would be the winner in terms of tolerability and safety [15]. A large, double-blind clinical trial comparing different prophylactic regimens in the same population of patients is eagerly waited to definitely assess the potential of rifaximin in colorectal surgery.

H. pylori Infection

H. pylori infection is a long-lasting, transmissible disease that has spread worldwide causing significant morbidity and mortality with a relevant economic impact. Since Warren and Marshall [155] first described the infectious etiology of peptic ulcer disease in 1984, a great deal of evidence has accumulated to suggest that *H. pylori* eradication therapy cures peptic ulcer disease [156] and can also be beneficial to other *H. pylori*-related diseases [157]. Having been classified as a definite 'type I carcinogen' [158], this Gram-negative, microaerophile bacterium needs to be eradicated from the host anyway [159, 160] since this can prevent the development of dyspeptic symptoms and peptic ulcer disease in healthy asymptomatic subjects [161] and that of gastric cancer in dyspeptic patients [162]. The survival capabilities of the *H. pylori* organism within the stomach make it, however, difficult to eradicate. The organism is able to survive over a wide pH spectrum. It is found within the gastric mucus layer, deep within the mucus-secreting glands of the antrum, attached to cells, and even within cells [56]. The organism must be eradicated from each of these potential niches and this is a daunting task for any single antibiotic. Initial attempts to cure the infection showed that the presence of antibiotic susceptibility in vitro did not necessarily correlate with successful treatment. It was rapidly recognized that therapy with a single antibiotic led to a poor cure rate and various antimicrobial mixtures were tried resulting in several effective combinations of antibiotics, bismuth, and antisecretory drugs [163].

Several European Guidelines [for a review, see 163] suggest the use of a 7-day triple therapy, comprising a proton pump inhibitor (or ranitidine bismuth citrate), clarithromycin and amoxicillin or metronidazole, as first-line therapy, whilst a 7-day quadruple therapy (proton pump inhibitor, bismuth salts, tetracycline, and metronidazole) is indicated for patients with eradication failure. However, increasing evidence suggests that the success rate following such regimens is decreasing in several countries. Indeed, some systematic reviews showed that standard triple therapies fail to eradicate *H. pylori* in up to 20% of patients [163]. Moreover, even lower cure rates have been observed in primary medical care settings, bacterial eradication being achieved in only 61–76% of patients [164]. As a consequence, new therapeutic combinations to cure *H. pylori* infection have been pioneered in the last few years. In addition, some effective rescue therapies have been developed in order to overcome treatment failures [165–167].

Table 10. Genes affected by point mutations or other genetic events leading to antibiotic resistance in *H. pylori* and the frequency of resistance (from Mégraud and Lamouliatte [166])

Antibiotic group	Genes affected	Frequency of resistance, %
Macrolides	23S rRNA	0–20
Metronidazole	<i>rdxA</i> , <i>fixA</i>	10–90
Quinolones	<i>gyrA</i>	0–10
Rifamycins	<i>rpoB</i>	0–5
Amoxicillin	PBP-1A	few cases described
Tetracycline	16S rRNA	few cases described

Although there are several reasons accounting for the lack of efficacy of eradication regimens, the main reason was found to be *H. pylori* resistance to antimicrobials (especially clarithromycin and metronidazole). The microorganism can become resistant to most antibiotics by chromosome mutation. This is the essential resistance mechanism found in this bacterial species, although genetic exchanges, especially transformation, have also been documented [166]. The genes affected by point mutations together with the frequency of resistance for the commonly used classes of antimicrobials are shown table 10.

Rifamycin derivatives (such as rifampicin, rifabutin and rifaximin) display antibacterial activity against *H. pylori* [51–54, 168, 169]. Rifabutin, whose oral bioavailability is rather low (i.e. 20%), is being increasingly used in some 'rescue' therapies after failed eradication attempts [170–172]. The prevalence of *H. pylori* resistance to this group of antibiotics is not exactly known but is probably extremely low as these drugs have – until recently – only been used in a limited number of patients to treat mycobacterial infections. On these grounds, some new rifamycin derivatives that display a potent bactericidal activity against *H. pylori* have been patented by different pharmaceutical companies. More interestingly, these compounds seem to be also active against those bacterial strains resistant to both metronidazole and clarithromycin [173].

Because of its antibacterial activity against microorganisms [51–54] and the lack of strains with primary resistance [52, 53], some preliminary studies [174–176] have explored the potential of rifaximin for *H. pylori* eradication. A first single-blind randomized study [174] evaluated the efficacy of rifaximin alone or in combination in *H. pylori*-positive patients with antral gastritis and

Table 11. Eradication rate and incidence of adverse events following administration of rifaximin and different rifaximin-based eradication regimens in *H. pylori*-positive patients (from Pretolani et al. [174])

Regimen	Eradication rate %	Adverse events %
Rifaximin 400 mg b.i.d.	40 (4/10)	0 (0/10)
Rifaximin + CBS 120 mg b.i.d.	50 (5/10)	20 (2/10)
Rifaximin + clarithromycin 500 mg b.i.d.	73 (8/11)	18 (2/11)
Rifaximin + metronidazole 250 mg t.i.d.	60 (6/10)	30 (3/10)

Drugs were given for 2 weeks and eradication checked 1 month after stopping treatment with rapid urease test and histology on biopsy samples taken from the antrum and the corpus. CBS = Colloidal bismuth subcitrate.

found the dual therapy with clarithromycin particularly effective (table 11).

On the basis of these results, a triple therapy including rifaximin, amoxicillin and omeprazole was attempted [175]. However, results were disappointing since the eradication rate (i.e. 60%) did not differ from that observed with dual therapy. However, in this study high doses (600 mg 3 times daily) of rifaximin suspension (2%) were used, which proved in a subsequent study [176] to be less effective than the tablet formulation. In addition, large volumes of the suspension had to be taken, which might have lowered patients' compliance and, consequently, the eradication rate [175]. Moreover, in the above study, drugs were given on an empty stomach. Since a meal markedly prolongs gastric residence time of the drug and improves its intragastric distribution to the body and fundus [177], postprandial dosing seems a more suitable strategy for improving topical delivery and mixing (thanks to increased antral motility) of antimicrobials [177, 178], provided that binding to or inactivation by food does not occur. An additional finding that would suggest postprandial dosing is that eating is associated with desquamation of gastric surface cells and discharge of mucus [179], possibly exposing the organisms to higher concentrations of the antimicrobial agent, or exposing a higher percentage of the organisms to it.

In conclusion, rifaximin-based eradication regimens are promising but new antimicrobial combinations (with and without proton pump inhibitors) need to be explored in well-designed clinical trials including a large cohort of *H. pylori*-infected patients.

Selective Bowel Decontamination in Acute Pancreatitis

Acute necrotizing pancreatitis (ANP) still remains a life-threatening disease despite several improvements in diagnosis, prevention and treatment. Infectious complica-

tions are the most frequent and severe complications of acute pancreatitis with a mortality rate up to 80%. Bacterial infections, frequently caused by microorganisms of enteric origin, are often seen during the progression of severe ANP, concomitant with the potential development of multiple organ dysfunction [180]. The role of antibiotics in reducing infectious morbidity and mortality has been debated for decades because of a lack of supportive clinical data. Research completed over the past decade has helped to define the microbiology, establish the risk factors, and improve the understanding of the pathogenesis of infectious complications in patients with ANP [181]. These patients are at the greatest risk of developing an infection with enteric Gram-negative or Gram-positive bacteria 'translocated' from the bowel lumen into the necrotic pancreatic tissue [181, 182].

Bacterial translocation is defined as the passage of viable indigenous bacteria from the GI tract to extraintestinal sites, such as the mesenteric lymph node complex, liver, spleen and bloodstream [183]. Three major mechanisms promote bacterial translocation: intestinal bacterial overgrowth, deficiencies in host immune defenses and increased permeability or damage to the intestinal mucosal barrier [184]. These mechanisms can act in concert to promote synergistically the systemic spread of indigenous translocating bacteria to cause lethal sepsis.

Animal and human studies support the use of antibiotics for the prevention of infectious morbidity and mortality in severe ANP. The most effective antimicrobial agents are the fluoroquinolones, imipenem-cilastatin, and metronidazole, which achieve adequate penetration into pancreatic juice and necrotic tissue and inhibit the growth of enteric bacteria. Although a recent meta-analysis [185] suggested that prophylactic antibiotic administration reduces sepsis and mortality and this approach has been recommended by recent guidelines and consensus state-

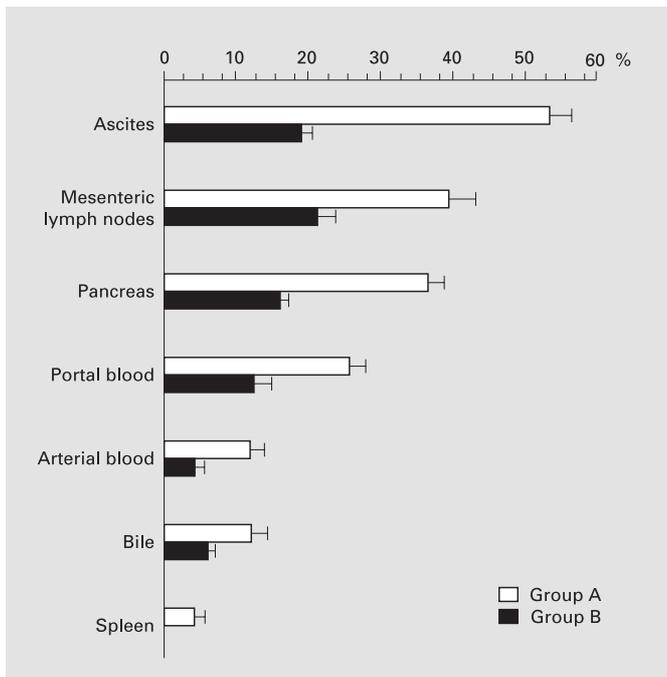


Fig. 8. Inhibition of bacterial translocation by rifaximin in the rat. Results of bacteriological culture (for aerobic and anaerobic bacteria) of tissues and biological fluids obtained from rats with taurocholate-induced ANP. Bars represent the percentage of positive biological samples. Horizontal lines are standard errors. Animals of group A received no treatment, while those of group B were given an enema of 30 ml warmed saline containing rifaximin (20 mg/kg) 1 h prior the induction of ANP (from Marotta et al. [196]).

ments [186–191] on the treatment of ANP, the first double-blind, placebo-controlled trial [192] found no benefit of antibiotic prophylaxis with respect to the risk of developing infected pancreatic necrosis.

The route of antibiotic administration might be crucial. Animal studies [193, 194] have shown that enteral administration (either by oral or rectal route) of antimicrobials reduces the rate of bacterial translocation and early mortality in rats or mice with experimentally induced pancreatitis. Indeed, in patients with ANP, selective bowel decontamination with oral and rectal antibiotics decreased the infection rate [195].

Investigations performed in rats with experimental acute pancreatitis [196] or ulcerative colitis [197] have shown that both rectal and oral administration of rifaximin decreased colonic bacterial translocation towards mesenteric lymph nodes. In the model of ANP [196] not only was the intra-abdominal spread of enteric bacteria (fig. 8) significantly reduced but also the pancreatic damage was lessened by rifaximin treatment.

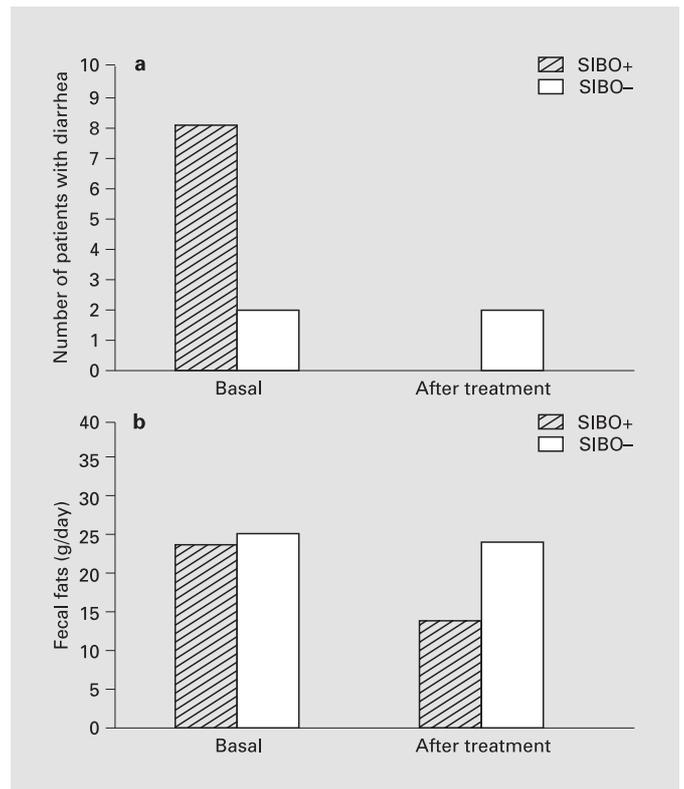


Fig. 9. Effect of short-term rifaximin administration (400 mg t.i.d. for 7 days) on diarrhea (a) and fecal fat excretion (b) in patients with chronic pancreatitis with and without SIBO (from Trespi and Ferreri [201]).

In addition to ANP where it is associated with GI dysmotility [198, 199], SIBO is present in a significant proportion of patients with chronic pancreatitis [200, 201]. Short-term rifaximin therapy was able to normalize the hydrogen breath test and improve symptoms (i.e. diarrhea and fecal fat excretion) in all patients studied (fig. 9) [201]. Bowel decontamination via administration of this ‘topical’ antibiotic could, therefore, be beneficial in both acute and chronic pancreatitis. Double-blind, placebo-controlled studies are to be performed to explore the rifaximin potential in this indication.

Prevention of Spontaneous Bacterial Peritonitis in Cirrhosis

Spontaneous bacterial peritonitis (SBP) is a serious complication of cirrhotic ascites, arising most frequently in those with advanced liver disease. Its development leads to a further reduction in the effective arterial blood volume, and it has a mortality rate equivalent to that of a variceal bleed [202]. Since hepatic blood flow and func-

tion are age-dependent [203], SBP could be particularly severe in the elderly.

SBP is considered a bacterial infection of a previously sterile ascitic fluid in the absence of any intra-abdominal, surgically treatable source of infection [204]. Independent predictors of the development of bacterial infections in hospitalized cirrhotic patients are poor liver synthetic function and admission for GI hemorrhage [205]. Diagnosis of SBP is established by a polymorphonuclear cell count in ascitic fluid ≥ 250 cells/mm³. The organism responsible for the infection is isolated in 60–70% of the cases. The remaining cases are considered to be a variant of SBP (culture-negative SBP) and are treated in the same way as those with a positive culture. The SBP resolution rate ranges between 70 and 90%, and hospital survival between 50 and 70% [206]. An early diagnosis and the use of a more adequate antibiotic therapy are the most probable reasons for the improvement in the prognosis for SBP in recent decades [205, 206]. Despite the resolution of the infection, SBP may trigger severe complications such as renal impairment, GI bleeding and accentuation of hepatic insufficiency, which are responsible for the associated mortality. Patients recovering from an episode of SBP should actually be considered as potential candidates for liver transplantation [206].

An *in vitro* [207] study on 124 aerobic bacterial pathogens obtained from patients with SBP showed that most isolates were Gram-negative organisms, and that *E. coli* and *Klebsiella pneumoniae* were responsible for 63% of the episodes evaluated. The fluoroquinolones (ciprofloxacin and ofloxacin) and the ‘fourth-generation’ cephalosporin cefpirome were the most active agents against the Gram-negative bacteria. Although the International Ascites Club (IAC) [208] suggested the use of intravenous cefotaxim or oral ofloxacin as the antimicrobial treatment of choice, a recent systematic review [209, 210] found no convincing evidence concerning efficacy of antibiotics in the treatment of SBP, and identified several gaps that warrant future research. Since bacterial translocation, depressed activity of the reticuloendothelial phagocytic system and decreased antibacterial capacity of ascitic fluid seem to be the main steps in the pathogenesis of ascitic fluid infection [211], antimicrobial prophylaxis seems, however, indicated.

It is now well established that the gut is the source of most of the bacteria that eventually cause SBP [212]. As cirrhosis develops in animals, Gram-negative bacteria increase in numbers in the gut [213]. Furthermore, the gut of animals and patients with advanced cirrhosis is more permeable to bacteria than the normal gut and more

permeable than the gut in less advanced cirrhosis [214, 215]. Once bacteria reach a critical concentration in the gut lumen, they ‘spill over’, and escape from the gut, ‘translocating’ to mesenteric lymph nodes. Then, they can enter lymph, blood, and eventually ascitic fluid [216]. If the ability of the ascitic fluid to assist macrophages and neutrophils in killing the errant bacteria is deficient, uncontrolled growth occurs [214]. Bacterial translocation is facilitated by SIBO that in turn is facilitated by disorders in intestinal motility. Bacterial overgrowth and alterations in intestinal motility have both been shown to be more frequent in cirrhosis, particularly in patients with the most severe liver disease [217, 218]. In summary, SBP is the result of a failure of the gut to contain bacteria and a failure of the immune system to kill the virulent bacteria once they have escaped from the gut. Treating SIBO should, therefore, prevent bacterial translocation and, consequently, SBP in cirrhosis. Indeed, selective intestinal decontamination with antibiotics reduces gut bacterial counts, reduces translocation rates, can prevent SBP in high-risk subgroups, and can improve the hyperdynamic circulatory state of these patients [208, 219, 220]. Selective intestinal decontamination can even improve the survival of rats with cirrhosis and ascites [221]. Although several guidelines [222] including those of the IAC [208] suggest long-term prophylaxis with oral norfloxacin, the growing microbial resistance to quinolones [223] and the well-known adverse effects of systemic antimicrobials [154], often amplified in patients with an impairment of the liver function, call for alternative effective and safe treatment. Thanks to its activity against the microorganisms often responsible for SBP and its safety profile, rifaximin could represent a suitable antibiotic for long-term prophylaxis in cirrhotics. In this connection, the efficacy of this rifamycin derivative, given for a week, monthly over a period of 12 months, in the prevention of SBP was prospectively evaluated in patients with alcoholic cirrhosis [224]. The incidence of peritonitis was significantly reduced by the antibiotic treatment. Rifaximin did correct SIBO in 91% of patients and in all of them the occurrence of SBP was prevented (table 12). Since these are the promising results of an open trial, a larger, double-blind, controlled study is needed to confirm these data and select the best dose regimen and duration.

Prevention of Nonsteroidal Anti-Inflammatory Drug-Induced Enteropathy

Nonsteroidal anti-inflammatory drugs (NSAIDs) are widely prescribed medicines. Although these compounds represent a very effective class of drugs, their use is associ-

Table 12. Prophylaxis of SBP with rifaximin in patients with alcoholic cirrhosis (from Trespi et al. [224])

	Cumulative incidence		
	rifaximin (n = 27)	controls (n = 25)	p value
Cirrhotic patients developing SBP	3/27 (11.1)	10/25 (40.0)	0.0246
Cirrhotic patients with SIBO developing SBP	1/11 (9.0)	8/9 (88.9)	0.0009

Figures in parentheses represent percentage.

ated with a broad spectrum of untoward reactions in the liver, kidney, skin and gut. GI problems constitute a wide range of different clinical pictures, ranging from symptoms such as dyspepsia, heartburn and abdominal discomfort to more serious events, like peptic ulcer and its life-threatening complications, bleeding and perforation [225]. Although the upper GI toxicity of NSAIDs is well documented, the fact that NSAID damage extends beyond the duodenum is less well recognized. Upper GI effects are indeed the most known and most feared adverse events, but it is evident that NSAID therapy is also associated with lower GI complications: 10–15% of NSAID users experience diarrhea [226]. Furthermore, in patients taking long-term NSAIDs [227–230] there is an increased risk of intestinal erosion, ulceration and perforation. The hitherto small number of reports of toxic effects of NSAIDs to the small bowel may reflect primarily a lack of diagnostic tools. Indeed, push enteroscopy [231] and endoscopic capsule [232] have only recently been available to clinicians. Nevertheless, a host of small bowel manifestations have now been documented, ranging from strictures causing dramatic small-bowel obstruction and severe bleeding to low-grade NSAID ‘enteropathy’, a syndrome comprising increased intestinal permeability and low-grade inflammation with blood and protein loss. The enteropathy, although not dramatic, may add to existing GI problems, especially in rheumatic patients, and contribute to iron deficiency anemia or hypoalbuminemia. NSAID use has also been associated with onset or relapse of IBD [227–230].

Animal studies indicate that the pathogenesis of NSAID small intestinal toxicity involves multiple interactions dependent on enterohepatic circulation, epithelial permeability, neutrophil infiltration and bacterial infection [233]. Several investigations [234–238] have suggested that bacterial flora may play a role in the pathogenesis of NSAID bowel injury and Robert and Asano [239] did show more than 25 years ago that germ-free rats are resis-

tant to indomethacin-induced intestinal lesions. A recent paper [238] found an unbalanced growth of Gram-negative bacteria in the ileum of NSAID-treated animals and showed that heat-killed *E. coli* cells and their purified lipopolysaccharide caused a deterioration of NSAID-induced ulcers. Additional studies demonstrated that antimicrobials, such as tetracycline [234], kanamycin [236, 237], metronidazole [235, 237] and neomycin plus bacitracin [237], attenuate NSAID enteropathy, thus giving further support to the pathogenic role of enteric bacteria. The evidence from animal experiments has been confirmed in human studies, showing that metronidazole, an antimicrobial mainly targeted against anaerobic organisms [240], significantly prevented indomethacin-induced increase of intestinal permeability in healthy volunteers [241] and reduced inflammation and blood loss in rheumatic patients taking NSAIDs (table 13) [242]. Although metronidazole is able to protect against mitochondrial uncoupling induced by indomethacin [243], it does not possess any ‘intrinsic’ anti-inflammatory activity [244]. Its beneficial effect on NSAID enteropathy is, therefore, likely to be due to the antibacterial action. In this connection, an elegant study [245] with microbiological cultures of luminal aspirates was able to show that small intestinal permeability is increased in subjects with SIBO. This finding can easily explain how antibacterial agents, by correcting SIBO, could counterbalance intestinal permeability changes which set in motion a series of pathophysiological events leading to gross lesion formation.

An indirect proof of the role exerted by gut bacteria in the pathogenesis of NSAID enteropathy is represented by the similarities between indomethacin-induced intestinal damage and Crohn’s disease [234, 246]. Not only are these lesions anatomically (both macro- and microscopically) similar [246], but also sensitive to the same drugs, e.g. sulfasalazine [234, 247], steroids [234, 247], immunosuppressive compounds [248], and antibiotics [234–237].

Table 13. Reduction of intestinal inflammation and blood loss by metronidazole in patients on NSAIDs (from Bjarnason et al. [242])

Parameter	Metronidazole 800 mg/day		Significance p
	before	after	
Fecal excretion of ¹¹¹ In-labeled neutrophils, %	4.7 ± 1.30	1.5 ± 0.36	<0.0001
Fecal excretion of ⁵¹ Cr-labeled red cells, ml	2.6 ± 0.44	0.9 ± 0.13	<0.01
5-hour urinary excretion ratio ⁵¹ Cr-EDTA/L-rhamnose	0.133 ± 0.012	0.1154 ± 0.017	NS

Intestinal inflammation was assessed by fecal excretion of ¹¹¹In-labeled neutrophils while blood loss was quantitated via fecal excretion of ⁵¹Cr-labeled red cells. The urinary excretion ratio of ⁵¹Cr-EDTA/L-rhamnose was used as an index of intestinal permeability.

There is now a huge amount of data suggesting a pathogenetic role of bacteria in Crohn's disease [249, 250]. As a consequence, antibiotics, including rifaximin [80], proved to be effective in both inducing and maintaining remission from the disease [251, 252]. All these data, taken together, are consistent with the hypothesis that rifaximin, by correcting SIBO, would prevent or lessen NSAID enteropathy. Conversely to metronidazole [253], this rifamycin derivative is virtually unabsorbed by the GI tract and is devoid of any carcinogenic potential, being therefore more suitable for long-term use. One could actually speculate that cyclic antibiotic administration, getting rid of enteric pathogenic bacteria, would protect in this way the intestine from the damaging effect of anti-inflammatory compounds. This could be particularly useful in elderly patients, in whom a coexistence of osteoarthritis and colonic diverticular disease [254] would make rifaximin administration very cost-effective. Studies to explore this appealing clinical use are on the way in our laboratory.

Safety and Tolerability

Some toxicological investigations of rifaximin were performed at the beginning of the 1980s and some additional studies have recently been completed. The old tests did not comply with good laboratory practice principles since they were carried out before 1986. However, overall, the experiments were sufficiently accurate to permit adequate assessment of the drug toxicity [59], particularly in view of its very limited oral absorption.

The studies were performed by the route proposed for human therapy, that is oral, and also by the intravenous

route as required by the guidelines; the animal species and the protocols used are considered as the best fitted for the different kinds of tests [59].

Acute and Chronic Toxicity in Animals

The single-dose toxicity studies were performed in two mammalian species, rat and mouse, by the route used in clinical practice, that is oral, as well as that ensuring adequate systemic exposure to the drug, that is intravenous. The subacute (3 months) toxicity studies were correctly carried out in the two animal species (rat, dog) in which also the pharmacokinetics was studied. Since in accordance with the International Conference on Harmonization (CPMP/ICH/286/95), 3-month toxicity studies support clinical trials for up to 1 month's duration (the longest duration of drug administration in clinical use), chronic toxicity studies have not been performed.

The acute oral treatment of rats and mice did not induce any important behavioral change. Only at the highest dose (2,000 mg/kg) was some excitement observed in rats, but no animal died. The LD₅₀ was, therefore, considered >2,000 mg/kg. Since only one mouse given that dose died, the lethal oral dose in that species was established to be 2,000 mg/kg [59, 255].

In subacute toxicity studies only the highest rifaximin dose (i.e. 100 mg/kg, corresponding to 25 times the therapeutic dose in humans) induced mild toxic effects (like, for instance, acute gastroenteritis) connected to the topical GI action of the drug [59, 255]. A dose-dependent increase of the total cholesterol value was recorded in female animals [255], most likely due to an alteration of biliary acid metabolism consequent to the antibiotic effect on gut flora [256].

Table 14. Adverse events with an incidence $\geq 2\%$ among patients receiving rifaximin (600 mg daily) or placebo in placebo-controlled studies (from rifaximin FDA label [34])

MedDRA preferred term	Patients, n	
	rifaximin tablets (600 mg/day; n = 320)	placebo (n = 228)
Flatulence	36 (11.3)	45 (19.7)
Headache	31 (9.7)	21 (9.2)
Abdominal pain NOS	23 (7.2)	23 (10.1)
Rectal tenesmus	23 (7.2)	20 (8.8)
Defecation urgency	19 (5.9)	21 (9.2)
Nausea	17 (5.3)	19 (8.3)
Constipation	12 (3.8)	8 (3.5)
Pyrexia	10 (3.1)	10 (4.4)
Vomiting NOS	7 (2.2)	4 (1.8)

Adverse events include those that may be attributable to the underlying disease. Figures in parentheses represent percentage. MedDRA = Medical Dictionary for Regulatory Activities (http://www.meddramsso.com/NewWeb2003/medra_overview/index.htm); NOS = not otherwise specified.

All these data, taken together, do suggest that rifaximin is devoid of any toxic effect in experimental animals.

Genotoxicity

It is clear today that no single test is capable of detecting all genotoxic agents. Therefore, the usual approach is to perform a standard battery of *in vitro* and *in vivo* tests for genotoxicity [257]. The Ames *Salmonella*/microsome mutagenicity assay (Ames test) is a short-term bacterial reverse mutation assay specifically designed to detect a wide range of chemical substances that can produce genetic damage that leads to gene mutations [258]. The test employs several histidine-dependent *Salmonella* strains each carrying different mutations in various genes in the histidine operon. Indeed, rifaximin was assessed with the Ames test in five different strains of *Salmonella typhimurium* with and without metabolic activation [59, 255]. Being an antibiotic, the drug was considerably toxic to all the five bacterial strains but was not mutagenic since there was no increase in the number of revertants. Furthermore, no mutagenic activity was detected by using the gene conversion test on *Saccharomyces cerevisiae* or the gene mutation test on *Schizosaccharomyces pombe* [59, 255, 259].

Additional studies were performed with eukaryotic cell systems (i.e. Chinese hamster ovary cells, rat bone mar-

row cells and human lymphocytes) and none of them revealed a genotoxic potential [59]. No carcinogenicity tests have been done, but the short duration of patient treatment and the nature of the drug itself, which is virtually unabsorbed, do not raise any concern for carcinogenic potential.

Reproductive Toxicity

In the early investigations oral administration of rifaximin did not induce any toxic effect on the reproductive ability of rats, on the embryo-fetal development of rats and rabbits and on the peri- and postnatal development of rats [59, 260, 261]. However, more recent embryo-fetal toxicity studies, performed in rabbits [262], revealed that fetuses from treated animals display an increased incidence of linked skeletal anomalies and variants associated with vertebral and rib configuration. Because of these findings the US FDA put rifaximin – like many other antimicrobial agents [263] – in the pregnancy category C¹ [34]. In the above experiments, however, the food intake of the female rabbits was significantly decreased during the first half of the treatment period and remained lower, but not significantly so, throughout the experiment. As a consequence, body weight loss for the majority of treated females was evident in the first few days of the treatment period. Subsequent body weight gains were, however, similar to control values but, overall, body weight gain of the rifaximin-treated group for the gestation period remained below that seen in the control group rabbits [262]. A similar reduction in body weight gain through preparturition and gestation was observed in rats given rifaximin [264]. These apparent effects on body weight were considered a reflection of the antimicrobial action of the drug on the intestinal flora (with consequent impairment of nutrient absorption) and, therefore, connected to its primary pharmacological action [264]. On the other hand, it is well known that maternal nutrition does affect fetal development [265] and that malnutrition can actually potentiate the teratogenic effects of some drugs, like for instance aspirin [266]. As a matter of fact, a careful examination of fetuses made it possible to conclude that the types and incidences of the observed skeletal anomalies are unrelated to maternal treatment with rifaximin [267]. It is worthwhile mentioning that the rabbit is generally not considered a suitable species for the study of teratological effects of antibiotics because its susceptibility to them can

¹ Animal reproduction studies have shown an adverse effect on the fetus; however, there are no adequate and well-controlled studies in humans, but potential benefits may warrant the use of the drug in pregnant women despite potential risks.

make clinical signs difficult to interpret [268]. It is however reassuring that a population-based case-control study [269] showed that maternal exposure to antituberculosis drugs during pregnancy does not reveal a detectable teratogenic risk to the fetus. It is, therefore, very unlikely that the minute amounts of absorbed rifaximin, which will never be taken throughout the full gestation period, affect fetal development in humans.

No studies evaluating the excretion of rifaximin into breast milk and its bioavailability to the infant have been performed. However, due to its very limited, if any, absorption from the GI tract and its physicochemical characteristics any milk excretion of the drug is unlikely [270].

Tolerability Profile in Humans

An evaluation of the rifaximin tolerability profile observed in almost 1,000 patients from 30 clinical trials was unable to identify a definite pattern of intolerance [33]. Very few adverse events have been reported during short-term treatment with the drug, the most frequently reported being gastrointestinal in nature (e.g. flatulence, nausea, abdominal pain and vomiting). It is worthwhile to emphasize that the detection of GI adverse reactions could have been difficult in rifaximin trials since the symptoms of the underlying diseases were often similar to the GI complaints observed after drug treatment.

The safety of rifaximin, 200 mg 3 times daily, was evaluated more recently in 320 patients from two placebo-controlled clinical trials [34]. All the adverse events for either rifaximin or placebo that occurred at a frequency $\geq 2\%$ are shown in table 14. With the exception of flatulence, which occurred significantly ($p = 0.0071$) less frequently after drug treatment, the adverse event profile of rifaximin overlapped that of placebo.

Prolonged therapy with high doses of the antibiotic has been associated with infrequent urticarial skin reactions [33]. A significant increase in serum potassium and sodium concentrations, although within the physiological range, has been observed after the drug. Since rifaximin was employed mainly for the treatment of diarrheal diseases, this finding could likely be connected to the electrolyte disturbances of underlying conditions.

Postmarketing Surveillance

The excellent safety profile observed in clinical trials has been confirmed by the postmarketing surveillance program [117]. More than 8.5 million patients have been treated in Italy and abroad with rifaximin since its introduction to the market. During the overall postmarketing

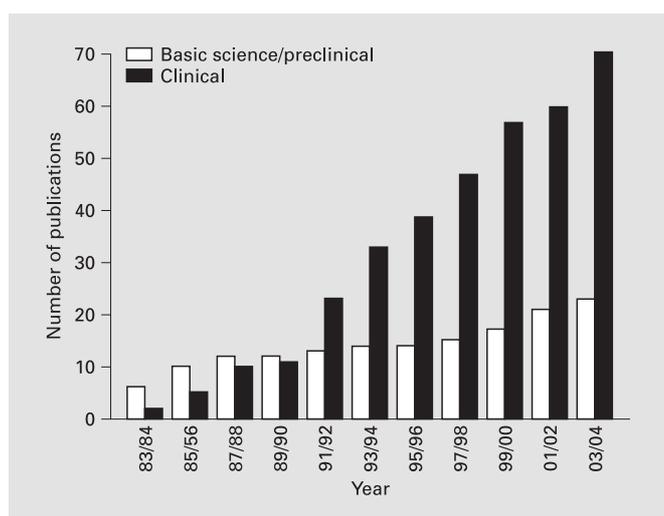


Fig. 10. Cumulative number of preclinical and clinical publications on rifaximin (courtesy of Lorin Johnson, PhD). Only papers quoted in Medline are presented.

period, 26 adverse reactions (17 cases of patients) were reported to the manufacturer, of which only 4 were judged to be serious. They consisted in 1 case of angioneurotic edema, 1 of cutaneous rash and 2 of urticaria.

In summary, rifaximin appears to be extremely safe with a very favorable risk-to-benefit ratio.

Conclusions and Perspectives for the Future

Rifaximin was first described in 1982 and was introduced into the Italian market 5 years later. Taking into account its excellent activity against a broad range of enteropathogens, the first 'logical' indication for this GI-targeted antibiotic was the treatment of infectious diarrhea in both adults and children. However, the appreciation of the pathogenic role of gut bacteria in several organic and functional GI diseases [12, 13] has increasingly broadened its clinical use.

Being virtually unabsorbed, this antimicrobial has little value outside the area of enteric infections, thus minimizing both antimicrobial resistance and systemic adverse events. It proved to be safe in all patient populations, including young children. Although pregnant women were purportedly excluded from controlled trials, clinical experience does suggest that the use of a nonabsorbable antibiotic, when strictly needed, represents the safest choice in this physiological condition. In this connection,

it was shown that treatment with oral neomycin, a poorly absorbed aminoglycoside, during pregnancy presents no detectable teratogenic risk to the human fetus [271]. Rifaximin, therefore, possesses almost all the characteristics of the 'ideal' antibiotic targeted at the GI tract [272].

As shown in table 7, there are established and potential clinical indications for this peculiar drug. In all these conditions, many of which share SIBO as a common feature, gut bacteria represent the specific target of rifaximin. The drug can be used alone (like, for instance, in the treatment of infectious diarrhea) or as add-on medication (as in the management of IBD) and given short-term (single course of treatment) or long-term (repeated courses of therapy, i.e. cyclically).

Although rifaximin has stood the test of time, it still attracts the attention of both basic scientists and clinicians as attested by the regular number of publications which appear every year in the medical literature (fig. 10). As a matter of fact, with the advancement of the knowledge on microbial-gut interactions in health and disease novel indications and new drug regimens are being explored.

Besides widening the clinical use, the research on rifaximin is also focused on the synthesis of new derivatives [173, 273] and on the development of original formulations, like soft capsules [274] and a gum-like device [275], designed for the controlled and continuous release of rifaximin. The former preparation, being bioadhesive, could be particularly useful in the *Helicobacter* eradication from the stomach whereas the latter should make the treatment of infections of the oral cavity possible, thus expanding the spectrum of clinical use.

All this ongoing research clearly indicates that the final chapter on this stimulating antibiotic has not yet been written.

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Note Added in Proof

The antibacterial activity of rifaximin has recently been confirmed in a clinical pharmacological study [1]. Two groups of healthy volunteers pretreated with either rifaximin (200 mg t.i.d. for 3 days) or placebo were challenged after the 4th dose of the antibiotic with *Shigella flexneri* 2a (1,000–1,500 CFU). The incidence of diarrhea was 0% in the rifaximin group compared with 60% in the placebo group ($p = 0.001$ at Fisher's exact test). The results of this study support the use of rifaximin in the prevention of infectious diarrhea in the traveler (TD). And indeed, DuPont et al. [2] have just reported that even once daily administration of rifaximin is able to prevent TD. The same team of investigators [3] has shown that up to 28% of the patients who developed TD still complain of chronic symptoms (namely loose stools, abdominal pain and fecal urgency) 6 months after the travel and that 11% of them met the Rome II criteria for irritable bowel syndrome. These findings underline the high incidence of postinfectious complications of TD and call for the use of antibiotic prophylaxis to prevent long-lasting complaints in the returning traveler.

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Rifaximin: In vitro and in vivo Antibacterial Activity – A Review

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Key Words

Rifaximin · Antibacterial activity · Gram-positive bacteria · Gram-negative bacteria

Abstract

In vitro inhibitory activity of rifaximin is directed against Gram-positive and Gram-negative, aerobic and anaerobic bacteria. It is effective in the treatment of gastrointestinal infections when given orally because of the high concentration of the drug remaining in the gut lumen. Laboratory investigations have been carried out to assess the in vitro activity of rifaximin on different bacterial strains isolated from both human and domestic animals. The objective of this project is to review the in vitro and in vivo activity of rifaximin against bacterial infection with Gram-negative rods, Gram-positive rods and Gram-positive cocci and their resistance to rifaximin. The available data suggest that rifaximin is active in vitro and in vivo in the treatment of bacterial infection of adults and children.

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Introduction

Rifaximin (4-deoxy-4'-methylpyrido-[1',2'-1'2'] imidazo[5,4-C] rifamycin SV) is a semisynthetic antibiotic molecule belonging to the rifamycin group [1] and synthesized by Alfa Wassermann (Bologna, Italy). Contrary to its structural analogue, rifaximin's chemical and physical properties do not allow its absorption following oral administration [2]. This feature prompted researchers to investigate the drug as a topical antimicrobial agent capable of acting against intestinal bacteria without causing systemic side effects.

Rifaximin is available in Europe for the treatment of acute intestinal bacterial infections, hepatic encephalopathy, bacterial overgrowth syndrome, diverticular disease of the colon, and for the prevention of infections after colorectal surgery [3, 4]. Rifaximin is also licensed in Mexico, Asia and Northern Africa and has recently been approved in USA for the treatment of traveler's diarrhea.

Rifaximin shows in vitro activity against a broad range of enteric pathogens. The minimal inhibitory concentration (MIC) at which 90% of isolates are inhibited (MIC₉₀) is 16–50 µg/ml for the various bacterial enteropathogens [5]. Ordinarily, isolates at this level of susceptibility would be considered 'resistant'. However, the drug is active against bacterial enteropathogens in vivo, which is

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Table 1. In vitro susceptibility to rifaximin (MIC) of enteric pathogens isolated from patients with bacterial diarrhea from multiple areas of the world [Mathewson et al., unpubl. data] [17, 18]

Species	Number of isolates	MIC ₅₀ µg/ml	MIC ₉₀ µg/ml	MIC range µg/ml
<i>Aeromonas</i> spp.	27	16	128	16 to >256
<i>Campylobacter jejuni</i>	54	12.5	>100	0.78 to >100
<i>Campylobacter</i> spp.	35	32	128	0.25 to >256
Enteroaggregative <i>E. coli</i>	50	64	128	16 to >256
Enterohemorrhagic <i>E. coli</i>	17	64	>256	32 to >256
Enteroinvasive <i>E. coli</i>	20	64	128	8 to >256
ETEC	153	64	128	8 to 256
ETEC with LT	50	64	256	8 to >256
ETEC with ST	76	64	128	8 to >256
ETEC with ST and LT	27	64	128	32 to >256
<i>Plesiomonas shigelloides</i>	25	64	256	16 to >256
<i>Salmonella</i> spp.	53	64	128	8 to >256
<i>Shigella</i> spp.	88	64	128	32 to >256
<i>Vibrio</i> spp. ¹	25	128	128	8 to 128
<i>Yersinia</i> spp.	91	12.5	25	0.2 to 25

LT = Heat-labile toxin; ST = heat-stable toxin.

¹ *Vibrio* spp. include non-cholera-causing vibrios.

explained by the extremely high luminal levels of the drug when it is administered orally. The fecal levels were in the range of 4,000–8,000 µg/g stool in a previous study [5]. The objective of this project is to review the in vitro and in vivo activity of rifaximin against bacterial infection.

Mechanism of Action

The mechanism of action of rifaximin depends on the inhibition of DNA-dependent RNA polymerase of the target microorganisms, leading to the suppression of initiation of chain formation in RNA synthesis.

Pharmacology

Animal and human studies have demonstrated that rifaximin has very poor intestinal absorption after oral administration, so that blood and urine concentrations of rifaximin are practically undetectable [6]. Rifaximin excretion is essentially exclusively by the fecal route [5]. Therefore, when rifaximin is administered by the oral route, it acts locally at the intestinal level and eliminates the bacterial organisms that are causing the infection. The important antibacterial activity of rifaximin appears to be directly related to the high intestinal concentration of the drug and inhibition of bacterial growth. The drug has

been used successfully in the treatment of infectious diarrhea in adults and children [7, 8], with low potential for inducing resistance by infecting bacterial strains [9].

Gram-Negative Rods Identified from Patients with Gastrointestinal Diseases

Diarrhea affects approximately 40% of travelers to tropical and semitropical areas of the developing world [10]. In up to 50–80% of patients with travelers' diarrhea, enteric bacterial pathogens are identified as the cause [11]. The important causes of travelers' diarrhea include enterotoxigenic *Escherichia coli* (ETEC) [11], enteroaggregative *E. coli* [12], *Salmonella* spp., *Shigella* spp., *Campylobacter* spp., *Plesiomonas shigelloides*, *Aeromonas* spp. and non-cholera-causing vibrios [11]. Since bacterial agents appear to be responsible for most causes of travelers' diarrhea and an important part of more severe diarrhea in children and adults in developing countries and industrialized regions, antimicrobial agents have become important for the treatment of this disease. Previous studies have shown that nonabsorbable antibacterial drugs were highly effective in treating travelers' diarrhea [13–15].

When bacterial isolates from patients with travelers' diarrhea from four geographic areas were tested [16], rifaximin was shown to have an MIC at which 50% of the

Table 2. Activity of rifaximin ($\mu\text{g/ml}$) against isolates of *H. pylori* according to the pH of the culture medium [20, 21]

Number of strains	pH 6.0			pH 7.2		
	MIC ₅₀	MIC ₉₀	MIC range	MIC ₅₀	MIC ₉₀	MIC range
30	4	4	0.5–8	1	2	0.25–4
43	2	8	0.5–8	NA	NA	NA

NA = Not available.

isolates tested were inhibited (MIC₅₀) of 12.5–128 $\mu\text{g/ml}$. The MIC₉₀ was 25 to >256 $\mu\text{g/ml}$. The results of in vitro susceptibility testing of specific enteric bacterial pathogens to rifaximin from three published studies [Mathewson et al., unpubl. data] [17, 18] are given in table 1. Differences were seen for individual enteric bacterial pathogens but no overall pattern of increased or decreased susceptibility was seen with the tested pathogens.

Helicobacter pylori has received much attention as a cause of gastritis, peptic ulcer and gastric cancer. In developing countries, the organism is acquired early in childhood, and up to 90% of children are infected by the age of 5 [19]. As the evidence linking *H. pylori* and gastric disease is impressive, eradication of *H. pylori* could reduce the rate of duodenal ulcer relapse and possibly gastric cancer. Megraud et al. [20] studied 43 *H. pylori* strains isolated from patients with gastritis for in vitro rifaximin activity (table 2). Rifaximin, amoxicillin and colloidal bismuth subcitrate all appeared to be active against *H. pylori* [20]. The MIC of rifaximin was influenced by the pH of the culture medium, with the agent being more active at basic pH. In another study, a total of 20 patients with *H. pylori* chronic gastritis were enrolled in a clinical trial of rifaximin. The trial demonstrated efficacy of rifaximin as treatment for *H. pylori* infection without important clinical side effects [21].

Gram-Positive Rods – *Mycobacterium* Identified from Patients with Tuberculosis

Tuberculosis remains a major global public health problem. The World Health Organization estimates that there are 8 million new cases of tuberculosis and 3 million deaths directly attributable to tuberculosis each year [22]. Drug resistance is a serious problem worldwide. In a study in New York City, 33% of patients with tuberculosis were infected by organisms resistant to at least one antituberculosis drug, and 19% were infected by organisms resistant

to both INH and rifampin [23], the first-line drugs in the treatment of tuberculosis.

Mycobacteria are also killed in vitro, as expected from an antibiotic sharing the properties of the rifamycin family [24]. In a study by Soro et al. [25], the MIC of rifaximin was determined for five *Mycobacterium tuberculosis* isolates from patients with tuberculosis. MIC concentrations were studied at 6, 20, 90 and 270 $\mu\text{g/ml}$, respectively. No resistant organisms were found. Growing *M. tuberculosis* in the presence of varying doses of rifaximin did not induce the occurrence of rifampicin-resistant strains [25]. In addition to this, experimental tubercular infection in the guinea pig was found not to be affected by an oral treatment course with rifaximin, therefore confirming the lack of absorption of the molecule after oral administration [26].

Anaerobic Bacteria – *Clostridium*

Rifaximin has been shown to possess good antibacterial activity against a variety of anaerobic bacteria (table 3) [24, 27, 28]. Anaerobes have been shown to be capable of producing ammonia (especially *Clostridia*), which has been incriminated in the pathogenesis of hepatic encephalopathy [29]. The authors suggested that since rifaximin is a nonabsorbable and effective antibiotic against anaerobic flora, it would be an ideal treatment for patients with compromised hepatic function. *Eubacterium* is inhibited by rifaximin with an MIC₉₀ $\leq 2 \mu\text{g/ml}$ [27].

Gram-Positive Cocci

Streptococcus, *Enterococcus* and *Staphylococcus* are clinically the most important of the Gram-positive cocci. Table 4 shows the range of MIC of rifaximin for 206 Gram-positive cocci where the MIC₅₀ was ≤ 0.015 –2 $\mu\text{g/ml}$ for the tested strains [27, 30, 31]. The MIC₉₀ was

Table 3. In vitro antibacterial activity of rifaximin against anaerobic bacteria [24, 27, 28]

	Number of isolates	Source of isolates	MICs, µg/ml		
			MIC range	MIC ₅₀	MIC ₉₀
<i>Bifidobacterium</i> spp.	6	cirrhotic patients, Italy	0.4–50	0.8	6.2
<i>Clostridium difficile</i>	4	cirrhotic patients, Italy	0.2–0.8	0.2	0.8
<i>Clostridium difficile</i>	93	human, Italy	0.004–128	0.004	128
<i>Clostridium perfringens</i>	15	various clinical material, unknown location	unknown	unknown	4
<i>Clostridium</i> spp.	26	cirrhotic patients, Italy	0.0125 to >100	0.4	50
<i>Propionibacterium</i> spp.	10	cirrhotic patients, Italy	0.025–12.5	0.2	12.5

Table 4. In vitro activity of rifaximin against Gram-positive cocci [27, 30, 31]

	Number tested	MIC ₅₀ , µg/ml	MIC ₉₀ , µg/ml	MIC range, µg/ml
<i>Staphylococcus aureus</i>	51	0.015	>8	≤0.01 to >8
<i>Staphylococcus epidermidis</i>	20	≤0.015	≤0.015	≤0.015
<i>Staphylococcus haemolyticus</i>	10	≤0.015	≤0.015	≤0.015 to >8
<i>Enterococcus faecalis</i>	21	2	8	0.5 to >8
<i>Enterococcus faecium</i>	11	2	>8	≤0.015 to >8
<i>Enterococcus</i> spp.	10	0.25	2	≤0.015 to >4
<i>Streptococcus</i> group A	19	0.12	0.25	≤0.03 to 0.25
<i>Streptococcus</i> group B	20	0.12	0.25	0.06 to 0.25
<i>Streptococcus</i> groups C, F and G	14	≤0.03	0.06	≤0.03 to 0.5
<i>Streptococcus pneumoniae</i>	30	≤0.03	0.06	≤0.03 to >4

≤0.015 to >8 µg/ml. Since Gram-positive cocci cause severe systemic infection, an orally administered drug without absorption would play no role in the therapy of these infections.

In vivo Rifaximin Activity in Intestinal Infections

Antimicrobial agents play an important role in the treatment of enteric infections caused by certain pathogens [32]. Several studies have been carried out to assess the effectiveness of rifaximin for eradication of causative bacterial enteropathogens [15, 33]. ETEC was the principal pathogen identified in these studies, since the target populations were travelers from industrialized countries to developing countries. The microbiologic eradication rates for rifaximin versus ciprofloxacin were not statistically significant. Furthermore, rifaximin has been shown

to shorten the duration of bacterial diarrhea in children and diarrhea due to ETEC, *Shigella* and *Salmonella* in nontravelers and to shorten the duration of excretion of the infecting organism [8].

Interestingly, in a small study on patients with AIDS, rifaximin was found to be effective against infectious diarrhea with stool cultures positive for protozoal pathogens, such as *Cryptosporidium parvum* and *Blastocystis hominis* [34]. The favorable effects of rifaximin on protozoal diarrhea have been also reported in a recent multicenter study on patients with travelers' diarrhea [33]. In fact, patients with pretreatment stools positive for *Cryptosporidium* infections obtained a clinical improvement with rifaximin significantly superior to the placebo-treated subjects.

Rifaximin Resistance

Two concerns with all antimicrobial agents intended for oral administration are the risk of depleting normal gut flora and the potential to induce antibacterial resistance. These aspects have been studied in several trials.

Minimal effects on intestinal flora were seen with rifaximin administration [9, 35]. In an early study, performed on healthy volunteers who received a short-term (5 days) rifaximin treatment, the observed changes in bowel flora returned to baseline levels within 1–2 weeks [9]. In a recent investigation fecal samples of patients with ulcerative colitis given three 10 day courses of the antibiotic were cultured and the different microbial species quantitated. Despite the high dose (i.e. 1800 mg daily) of rifaximin used there was only a minor change in bacterial counts which reverted back to pre-treatment values during the washout period [35]. It appears therefore that administration of this antibiotic does not disrupt intestinal microbial ecology.

Development of resistance to rifaximin is primarily due to a chromosomal one-step alteration in the drug target, DNA-dependent RNA polymerase. This differs from the plasmid-mediated resistance commonly acquired by bacteria to aminoglycoside antibiotics such as neomycin

[36]. In a recent study, we found no acquisition of rifaximin resistance in 27 rifaximin-treated subjects colonized by *Enterococcus* [37]. The MIC₅₀ and MIC₉₀ for the treatment group (rifaximin at a dose of either 400 or 200 mg twice daily for 3 days) were similar (16–64 µg/ml). In two published studies, rifaximin resistance was shown to occur in the bacterial flora of individuals who received treatment with rifaximin at a dose of 800 mg/day for 5 days [9, 27]. Within 1–2 weeks after the end of rifaximin treatment, resistance rates appeared to have decreased to less than 20% of the intestinal flora. The resistant strains detected during treatment appeared to be unstable and unable to persistently colonize the intestinal tract.

The available data suggest that bacterial resistance does not seem to be a major concern of the therapy with rifaximin. However, further monitoring of the occurrence of bacterial resistance to rifaximin would help to clearly define its clinical importance.

Conclusion

The available in vitro and in vivo studies suggest that rifaximin is active in the treatment of intestinal infections of adults and children.

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Rifaximin in the Treatment of Infectious Diarrhea

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Key Words

Rifaximin · Diarrhea · Treatment · Infectious diarrhea · Bacterial diarrhea · Travelers' diarrhea

Abstract

Rifaximin matches the criteria for an ideal agent for the treatment of infectious diarrhea. It has excellent activity against a broad range of enteropathogens. It is nonabsorbable, which may help explain its excellent side effect profile and lack of emergence of resistance because of high stool levels that are not likely to reach subinhibitory levels before the target Gram-negative bacilli are killed. It has shown excellent efficacy in numerous clinical trials of bacterial diarrhea. Because of the lack of systemic absorption and minimal adverse reactions, rifaximin should be useful in treating hosts such as pregnant women in whom the currently favored fluoroquinolones are contraindicated. Uses limited to enteric indications and its inherently low propensity to induce sustainable resistance among Gram-negative flora favor the sustained usefulness of rifaximin in the treatment of enteric infectious syndromes.

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Introduction

Antibiotics have a recognized role in the treatment of culture-proven bacterial causes of symptomatic enteric infection such as *Shigella* spp., *Campylobacter jejuni* and *Salmonella typhi*. The use of antibiotics in the treatment

of nontyphoidal salmonellosis is controversial, but the consensus is generally not to treat with antibiotics unless the patient is elderly or has another condition that predisposes to bacteremia. Also, when infection with enterohemorrhagic *Escherichia coli* is suspected in children, the consensus is not to use antibiotics because of the concern of increasing the risk of hemolytic uremic syndrome.

While fluid replacement remains the classic cornerstone of the treatment of diarrhea, especially in children, empiric antibiotic treatment is logical in certain situations. Children and adults with diarrhea and signs and symptoms suggestive of dysentery (fever, tenesmus, bloody, mucoid stools) can be given empiric antibiotics awaiting culture results. Severe diarrhea is more likely to be associated with bacterial causes. Some hosts and groups commonly have bacterial causes of their diarrhea even without clinical evidence of invasive pathogens, and the empiric use of antibiotics can be supported in such cases. Examples include diarrhea in patients with AIDS, in those with the so-called 'gay bowel syndrome' and in those traveling from developed to developing regions of the world.

As proposed in earlier publications, an ideal antimicrobial agent for the treatment of bacterial causes of infectious diarrhea would have the following features [1, 2]: (1) excellent activity against a broad range of bacterial enteropathogens; (2) nonabsorbable; (3) favorable side effect profile; (4) efficacious in the treatment of infectious diarrhea; (5) major indication is enteric disease, and (6) does not easily develop resistance or promote cross-resistance.

Is rifaximin, a rifampin-like antimicrobial agent that inhibits bacterial synthesis of RNA, an ideal agent for the

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Table 1. Rifaximin MIC₉₀ values for common enteropathogens causing travelers' diarrhea

Enteropathogen	Gomi et al. [4]		Sierra et al. [5]		DuPont et al. [6]		Peak stool concentration ³ /MIC ₉₀
	n	MIC ₉₀ , µg/g	n	MIC ₉₀ , µg/g	n	MIC ₉₀ , µg/g	
ETEC (= Enterotoxigenic <i>E. coli</i>)	97	32	38	16	72	32	250–500
EAEC	75	32	28	16	–	–	250–500
<i>Salmonella</i> spp.	46	64	14	4	9	16–32	125–2,000
<i>Shigella</i> spp.	36	64	64	16	11	32–64	125–500
<i>Campylobacter jejuni</i>	9	32	12	512	–	–	15–250
Others	21 ¹	4	11 ²	8	–	–	1,000–2,000

Gomi et al. [4] studied patients with diarrhea acquired in India, Mexico, Jamaica and Kenya, Sierra et al. [5] studied diarrhea in travelers returning to Spain from multiple locations, and DuPont et al. [6] studied patients with diarrhea acquired in Mexico. EAEC = Enterotoxigenic *E. coli*.

¹ Includes noncholera vibrios, *Plesiomonas shigelloides* and *Aeromonas* spp.

² *Aeromonas* spp.

³ 7,961 µg/g [3].

treatment of bacterial diarrhea? A critical examination of all the theoretic points suggests that it is the case.

Activity against Enteropathogens

First, rifaximin does have excellent activity against a broad range of enteropathogens. A problem is the definition of activity. Typically, a clinically acceptably low minimal inhibitory concentration (MIC) of an antibiotic is defined in terms of the pharmacokinetics of the antibiotic measured in serum. For some antibiotics (e.g. aminoglycosides), success is predicted by how much the concentration achieved in serum exceeds the MIC. For others (e.g. beta-lactam antibiotics), success is predicted by the percentage of the dosing interval that the serum level remains above the MIC. If an antibiotic like rifaximin is not absorbed, achieves concentrations in stool that far exceed the MIC of an organism and persists in stool at levels that exceed the MIC for days, clinical efficacy against the enteropathogen can be predicted.

When rifaximin is taken at clinical doses of 400 mg twice a day for 3 days, stool concentrations are high over the ensuing 5 days [3]. In that study, peak stool concentrations averaged nearly 8,000 µg/g, and remained in the range of 2–3,000 µg/g on days 4 and 5 after cessation of antibiotics [3]. In this same study, the authors measured the MIC at which 50% of the isolates were inhibited (MIC₅₀) and the MIC at which 90% of the isolates were inhibited (MIC₉₀) for 145 bacterial enteropathogens. The exact number of enterotoxigenic *E. coli* (ETEC), *Salmonella*, *Shigella* and other enteropathogens was not speci-

fied, but 50% of all bacterial isolates were inhibited at 12.5 µg/g and 90% were inhibited at 50 µg/g. The MIC ranges for ETEC and *Salmonella* were 0.098–200 µg/g. The MIC range for *Shigella* was 1.25–200 µg/g. Peak stool concentrations of rifaximin exceeded the MICs of these enteropathogens by 40- to over 80,000-fold. Defining susceptible as a ratio of stool concentration to MIC exceeding 1 during the anticipated dosing interval of 3–5 days, diarrheal illness caused by all of the 145 enteropathogens would be predicted to respond to treatment by rifaximin. As shown in table 1, Gomi et al. [4] determined the rifaximin MICs of 284 bacterial enteropathogens from four different locations around the world: India, Mexico, Jamaica and Kenya. Sierra et al. [5] determined the MICs of organisms causing diarrhea in travelers returning to Spain from various regions of the world. Table 1 also includes selected organisms from a comparative study of rifaximin versus ciprofloxacin in the treatment of travelers' diarrhea conducted by DuPont et al. [6] in Mexico. Taken in aggregate, these data show a consistent theme. Rifaximin MICs for common enteropathogens are similar from disparate regions of the world. Most MICs hover in the 32–64 µg/g range. Even relatively high MICs for *Campylobacter* noted by Sierra et al. [5] (but not by Gomi et al. [4]) are exceeded 15- to 250-fold by the concentrations of rifaximin achieved in stool.

Finally, one report of diarrhea in HIV-positive patients indicated that rifaximin might well prove useful in the treatment of protozoal causes of infectious diarrhea [4]. Thirteen patients had infections with *Cryptosporidium parvum*.

Lack of Absorption

The second feature of an ideal antibiotic for infectious diarrhea is that it should not be absorbed. Among 18 adult volunteers who received a 400-mg dose, no rifaximin was detected in serum 4 h later in 9 subjects, and the highest detected levels in the remaining 9 subjects was 5.3 ng/ml. The mean recovery in urine from a 400-mg dose was 0.007% in the first 24 h with a negligible amount detected in the second 24 h after the dose [7]. Furthermore, absorption of rifaximin is only minimally affected by colonic inflammation. Twelve subjects with mild to moderate ulcerative colitis took 400-mg doses. In only 4 subjects were levels ranging from 2 to 4.6 ng/ml noted irregularly, and 2 further subjects had single values of 13.4 and 22.4 ng/ml measured. The mean recovery in urine in the first 24 h was 0.009% [8]. That rifaximin is virtually non-absorbed is clear from these and other studies [9]. The reason that this is potentially important is that adverse reactions can be predicted to be higher for drugs with systemic absorption. A nonabsorbed drug should prove to be safe for special hosts like children and pregnant women.

Lack of absorption, however, was once considered to weigh against the predicted efficacy of an antibiotic in the treatment of bacterial diarrhea. Haltalin et al. [10] many years ago compared the efficacy of ampicillin with a non-absorbable aminoglycoside in the treatment of shigellosis. When the patients in the aminoglycoside arm failed to respond as those treated with ampicillin, the authors concluded that only an absorbable drug should be used in the treatment of bacterial diarrhea. It was later realized that poorly absorbed antimicrobials could be of value in the treatment of enteric infections. Indeed, oral bicozamycin and aztreonam proved to be efficacious in treating bacterial diarrhea, including cases caused by *Shigella* [11, 12].

Favorable Side Effect Profile

As might be expected from a nonabsorbed drug, the side effect profile of rifaximin has been shown to be excellent. Gillis and Brogden [9] summarized multiple studies in their excellent review of rifaximin. Although 963 patient experiences from 30 clinical studies were summa-

Table 2A. Summary of clinical studies of the efficacy of rifaximin in the treatment of acute infectious diarrhea: study design

Study	Design	Popula-tion	Definition of diarrhea	Enteropathogens	Medications				
					rifaximin		control		
					n	daily dose (duration)	drug	n	dose (duration)
Luttichau et al. [20]*	open label	adults	acute diarrhea	11 EPEC 4 <i>Shigella</i> 4 <i>Salmonella</i> 1 <i>Yersinia</i>	20	800–1,200 mg (5 days)	NA	NA	NA
Alvisi et al. [21]	open label	adults	acute diarrhea or bacterial super-infection of intestinal inflammatory diseases	1 <i>Yersinia</i> 2 patients with no pathogen various nonclassic enteric flora ¹	20	800 mg (5–10 days)	NA	NA	NA
Fiorentino et al. [22]*	open label	older children and adults	acute diarrhea	11 EPEC 7 <i>Salmonella</i> 1 <i>Shigella</i> 3 patients with no pathogen	22	600–1,200 mg (3–5 days)	NA	NA	NA
Lombardo and Santangelo [23]	open label	children	acute and chronic bacterial diarrhea	<i>E. coli</i> in almost all cases	31	>20 months: 200 mg load then 400 mg/day <20 months: 100 mg load then 200 mg/day (5 days)	NA	NA	NA
Sanfilippo et al. [24]	randomized, placebo-controlled, double-blind	children	acute diarrhea	rifaximin/control 10/8 <i>E. coli</i> 2/2 <i>Shigella</i> various nonclassic enteric flora	20	20–40 mg/kg/day (4 days)	placebo	17	matched placebo (4 days)

Table 2A (continued)

Study	Design	Popula- tion	Definition of diarrhea	Enteropathogens	Medications				
					rifaximin		control		
					n	daily dose (duration)	drug	n	dose (duration)
Vinci et al. [25]*	randomized, placebo-controlled, double-blind	adults	acute diarrhea	rifaximin/control 10/7 <i>E. coli</i> 8/5 ETEC 6/6 <i>Shigella</i> 1/2 <i>Salmonella</i> various nonclassic enteric flora	20	600 mg/day (5 days)	placebo	20	matched placebo (5 days)
Della Marchina et al. [26]*	randomized, placebo-controlled, double-blind	elderly adults (>73 years)	acute diarrhea	rifaximin/control 28/30 <i>E. coli</i> 6/4 <i>Shigella</i> 8/9 <i>Salmonella</i> 4/3 <i>Yersinia</i> various nonclassic enteric flora	63	600 mg/day (7 days)	placebo	58	matched placebo (7 days)
De Castro et al. [27]*	randomized 2:1 to rifaximin vs. oral rehydration, not blinded	children	acute episode of recurrent diarrhea occurring between cycles of antibiotic prophylaxis for UTIs	rifaximin/control 10/6 <i>Salmonella</i> 8/6 <i>Shigella</i> 6/1 <i>Yersinia</i> 4/2 <i>Campylobacter</i> 2/1 EPEC	30	400 mg/day (3–5 days)	oral re-hydration	16	NA
Alvisi et al. [28]	randomized, active drug-controlled, double-blind	adults	acute 'secretory' diarrhea	various nonclassic enteric flora	22	800 mg/day (5 days)	neomycin	21	1 g/day (5 days)
Stornello and Salanitri [29]*	randomized, active drug-controlled, not double-blind	children	acute diarrhea	rifaximin/control 8/7 EPEC 2/3 <i>Salmonella</i> various nonclassic enteric flora	20	200 mg load then 400 mg/day (3–6 days)	neomycin	20	<6 years: 125,000 IU/day 6–12 years: 375,000 IU/day (3–6 days)
Palermo et al. [30]	randomized, active drug-controlled, not double-blind	older children and adults	acute diarrhea	various nonclassic enteric flora	29	800 mg/day (4–7 days)	neomycin plus nalidixic acid capsules	20	1–2 capsules, 3–4 times/day (4–7 days)
Mazzitelli et al. [31]	randomized, active drug-controlled, not double-blind	adults	acute diarrhea	various nonclassic enteric flora	20	400–800 mg/day (7 days)	neomycin	20	1.5 g/day (7 days)
Beseghi and De Angelis [32]*	consecutive patient randomized, active drug-controlled, not double-blind	children	acute diarrhea	rifaximin/control 9/7 <i>Salmonella</i> 5/10 EPEC	14	400 mg/day (3–5 days)	neomycin plus bacitracin	17	5 ml, 4 times/day (3–5 days)
Frisari et al. [33]*	consecutive patient randomized, active drug-controlled, not double-blind	children	acute diarrhea	rifaximin/control 11/9 <i>Salmonella</i> 6/6 <i>Campylobacter</i> 3/3 <i>Shigella</i> 0/4 EPEC 1/0 <i>Salmonella</i> + <i>Campylobacter</i>	24	400 mg/day (3–5 days)	paromomycin	25	500 mg/day (4–5 days)
DuPont et al. [13]*	randomized, active drug-controlled, double-blind	adults	acute travelers' diarrhea	rifaximin/control 12/6 ETEC 7/1 <i>Shigella</i> , <i>Salmonella</i> , <i>Campylobacter</i>	55	18: 600 mg/day 18: 1,200 mg/day 19: 1,800 mg/day	trimetho-prim-sulfamethoxazole	17	320 mg trimetho-prim + 1,600 mg sulfamethoxazole/day
DuPont et al. [6]*	randomized, active drug-controlled, double-blind	adults	acute travelers' diarrhea	NA	93	800 mg/day	ciprofloxacin	94	1 g/day

Table 2B. Summary of clinical studies of the efficacy of rifaximin in the treatment of acute infectious diarrhea: study outcome

Study	Outcome of rifaximin treatment		
	clinical efficacy	adverse drug events	bacteriologic cure
Luttichau et al. [20]	50% reduction in stool frequency by end of day 1**	none	90%
Alvisi et al. [21]	50% reduction in stool frequency by day 4	none	89% (16/18) of original isolates eradicated
Fiorentino et al. [22]	50% reduction in stool frequency between day 1 and 2**	none	63% (7 <i>Salmonella</i> not eradicated despite clinical improvement)
Lombardo and Santangelo [23]	50% reduction in stool frequency between day 3 and 4	none	69% eradicated or markedly diminished
Sanfilippo et al. [24]	50% reduction in stool frequency between day 3 and 4; $p < 0.05$	none	90% of <i>Shigella</i> eradicated
Vinci et al. [25]	50% reduction in stool frequency by end of day 1; $p < 0.05$ **	none	86%
Della Marchina et al. [26]	50% reduction in stool frequency by end of day 2; $p < 0.05$ **	none	89%
De Castro et al. [27]	> 50% reduction in stool frequency by the end of day 1; 50% reduction in the control group at day 5; $p < 0.05$ **	none	93%
Alvisi et al. [28]	50% reduction in stool frequency by end of day 2 in both treatment groups	none	85%
Stornello and Stanitri [29]	50% reduction in stool frequency between day 3 and 4 in both treatment groups**	oral rifaximin therapy discontinued in 1 child due to vomiting	90%
Palermo et al. [30]	50% reduction in stool frequency by end of day 1 in both treatment groups	self-limiting urticarial rash	100%
Mazzitelli et al. [31]	baseline frequency lacking; patients passing 1–2 loose stools/day by day 2–3 of treatment	none	70%
Beseghi and De'Angelis [32]	> 50% reduction in stool frequency by end of day 1 in rifaximin group vs. 25% reduction in control group**	none	93%
Frisari et al. [33]	significant improvement in both groups by day 2*	none	90%
DuPont et al. [13]	50% well at 35 h**	minimal eosinophilia (5, 7 and 9%) in 3 subjects; 1 patient with posttreatment AST of 74; otherwise well tolerated	80%
DuPont et al. [6]	50% well at 27 h compared to 25 h for ciprofloxacin; $p = 0.3$ **	well tolerated in both groups	not available

EPEC = Enteropathogenic *E. coli*; NA = not applicable; UTI = urinary tract infections; AST = aspartate amino transferase.

* Indicates studies that documented a substantial number of cases of bacterial diarrhea; ** denotes studies with substantial isolation of bacterial enteropathogens.

¹ Nonclassic enteric flora includes *E. coli* with the designation of enteropathogenic *E. coli* or other recognized diarrheogenic *E. coli*, *Klebsiella*, *Proteus*, *Enterobacter*, *Pseudomonas* and *Enterococcus*.

rized, not all studies discussed drug tolerance specifically. Overall, only a handful of adverse events were recorded: 5 cases of flatulence, 2 of abdominal pain, 1 of nausea and 2 cases of vomiting in children. Overall, gastrointestinal side effects are estimated to occur in less than 1% of persons taking rifaximin. Other rare adverse events, like urticarial rash, headache, leg edema and weight loss, have been reported but often not considered drug-related by the investigators.

Efficacy in the Treatment of Infectious Diarrhea

Tables 2A and B summarize the results of clinical studies that assessed the efficacy of rifaximin in the treatment of bacterial diarrhea. Many of the studies suffer from small numbers and not uniform definitions of diarrhea. In the studies of DuPont et al. [6, 13], acute diarrhea was defined as three or more unformed stools passed in a 24-hour period accompanied by at least one symptom of

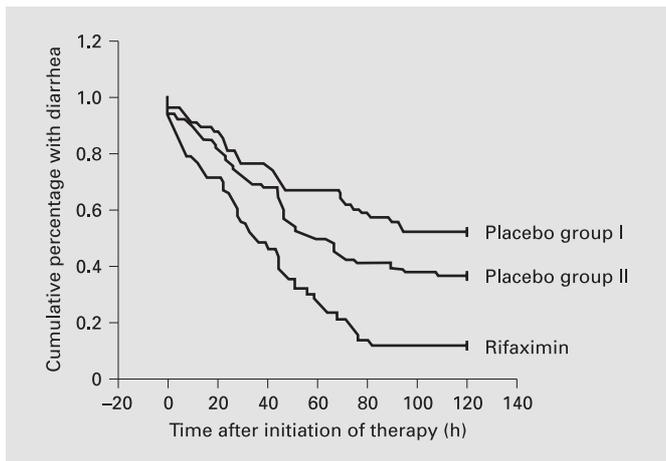


Fig. 1. Comparative effectiveness of treatment of patients with travelers' diarrhea. Percentage of subjects who continued to have diarrhea by hour after treatment in subjects receiving one of three doses of rifaximin compared with two similar placebo-treated groups studied earlier using identical study methods (from DuPont et al. [13]).

enteric disease such as abdominal cramps, nausea or vomiting. While most of the studies allude to the treatment of bacterial diarrhea, some are not accompanied by convincing stool bacteriology, with notable exceptions indicated by asterisks in the table. The studies, many of which were not rigorously assessed, included many different doses of rifaximin, ranging from as low as 400 mg to as high as 1,600 mg/day in divided doses. We determined the time to 50% reduction of the mean frequency of passage of unformed stools in an effort to compare the studies meaningfully. This median duration of diarrhea is the figure that best reflects the nonparametric statistics that would be performed when comparing therapeutic arms in a study and was chosen as a figure that could be derived from the published data and would permit a degree of comparison. Regarding adverse event data, most of the studies simply noted no adverse events, implying most likely that they passively recorded adverse events complained of by the subject rather than actively inquiring about possible adverse events and indicating when the incidence exceeded those reported in the placebo or control drug arm of the study.

Despite such limitations, the data are uniformly favorable and generally indicate that treatment with rifaximin successfully limits the course of bacterial and travelers' diarrhea to 1–2 days (fig. 1). In the case of travelers' diarrhea, for which bacterial causes are well known, when treatment with rifaximin was compared to treatment with either trimethoprim-sulfamethoxazole or ciprofloxacin

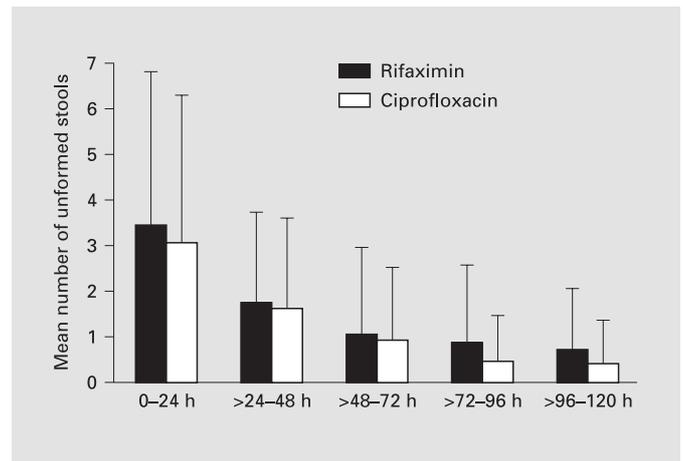


Fig. 2. Mean number of unformed stools passed per day of study by subjects with travelers' diarrhea taking rifaximin (400 mg twice a day) or ciprofloxacin (500 mg twice a day). The mean values for the two treatment groups were comparable for each day of the study (from DuPont et al. [6]).

(fig. 2), equivalent efficacy was noted. The average post-treatment duration of travelers' diarrhea was 35 h in one study [6] and 27 in another [13]. While data on the time required to clear bacteria from stool are lacking in most trials, the studies of DuPont et al. [6, 13] indicate that pathogens are removed from stool as effectively with rifaximin as with trimethoprim-sulfamethoxazole or ciprofloxacin and that the vast majority of pathogens were eradicated from stool when stool cultures were repeated at the conclusion of the therapeutic course.

While in patients with traveler's diarrhea due to *E. coli* the efficacy of the antibiotic was uniformly high, the cure rate was significantly lower when invasive pathogens (e.g. *Campylobacter jejuni*) causing fever and/or dysentery were present (DuPont, unpublished observations).

Rifaximin appeared to be effective and safe in both adults and children. Not only do these data support the efficacy of a nonabsorbable antibiotic in the treatment of diarrhea, the lack of absorption and degree of safety reported to date support the likelihood that rifaximin will be safe to use in pregnant women. Currently, the drugs of choice for the treatment of bacterial diarrhea, especially travelers' diarrhea, are the fluoroquinolones, which are contraindicated in pregnancy. While rifaximin will likely never be adequately studied in pregnancy, it should be safe.

One published study supports the use of rifaximin in the treatment of diarrhea with protozoal causes [14]. In this small study, Amenta et al. [14] showed that diarrhea

in patients with AIDS and from protozoal causes, including *Giardia lamblia* and *C. parvum*, responded favorably to treatment with rifaximin. In the Kenya arm of a recently reported multicenter trial of the treatment of diarrhea in travelers, rifaximin shortened the illness of European travelers with *Cryptosporidium* infection [15]. Further work should be done in these populations.

Finally, while a nonabsorbable antimicrobial agent has theoretic advantages as a choice for the treatment of infectious diarrhea, such an agent should not be used to treat hosts thought to have or be at risk for bacteremic disease (e.g. seriously immunocompromised patients).

Major Indication for Rifaximin Is Enteric Disease

As outlined in the excellent review by Gilles and Brogden [9], the current indications for rifaximin include surgical prophylaxis and the treatment of hepatic encephalopathy, infectious diarrhea and intestinal bacterial overgrowth syndromes. As such, rifaximin is aimed only at enteric flora. Owing to its lack of absorption, rifaximin will likely not be used for other conditions or indications. Such limited indications should help preserve the activity of the agent, since overuse for common conditions like urinary or respiratory tract infections will naturally not occur. Limited use should help retard the development of resistance among enteric flora.

Rifaximin Does Not Develop Resistance or Promote Cross-Resistance

While the emergence of rifaximin-resistant strains has been observed during the course of treatment, these strains disappear from the intestinal flora within 1–2

weeks of cessation of rifaximin [16]. It has been shown that in vitro, the emergence of resistant Gram-positive flora could be induced, but the emergence of aerobic Gram-negative flora was rare [Schito, personal communication]. Furthermore, emergence of resistance was much less common under anaerobic conditions that mimic the environment of the gut. Also, subinhibitory concentrations of an antibiotic encourage the emergence of resistance, a situation much less likely to occur in the gut because of the huge concentrations of rifaximin in stool.

Three studies have addressed the possibility of inducing cross-resistance to *Mycobacterium tuberculosis* during the use of rifaximin. In an experimental guinea pig model of *M. tuberculosis*, rifaximin was administered in an effort to induce resistance among *M. tuberculosis* strains of human origin. Not only did no resistance develop, cross-resistance to rifampin also did not occur [17, 18]. In another approach, *M. tuberculosis* strains were subjected to subinhibitory concentrations of rifaximin. No induction of resistance or cross-resistance to rifampin occurred [19].

Conclusion

Rifaximin appears to be an ideal agent for the treatment of infectious watery diarrhea. It has shown excellent efficacy in numerous clinical trials of bacterial diarrhea. Its excellent side effect profile and lack of systemic absorption predict that it should be useful in treating hosts for whom the currently favored fluoroquinolones are contraindicated. Uses limited to enteric indications and its inherently low propensity to induce sustainable resistance among Gram-negative flora favor the sustained usefulness of rifaximin in the treatment of enteric syndromes.

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Note Added in Proof

After the submission of this paper several publications have appeared in the literature that reinforce the role of rifaximin in the treatment of infectious diarrhea [1]. An entire issue of the *Journal of Travel Medicine* devoted to the use of this antibiotic in the treatment of traveler's diarrhea (TD) has been published [2-5]. In addition, a recent paper from our laboratory [6] confirmed the rifaximin efficacy also in enteroaggregative *Escherichia coli*-mediated TD. Furthermore, in a randomized, double-blind, placebo-controlled study [7] even once daily administration of the antibiotic proved to be capable of preventing TD. Finally, Lawler and Wallace [8] recently reviewed the treatment options for bacterial diarrhea and considered rifaximin a useful addition to our therapeutic armamentarium.

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Antibiotic-Associated Diarrhea and Pseudomembranous Colitis: Are They Less Common with Poorly Absorbed Antimicrobials?

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Key Words

Antibiotic-associated diarrhea · Pseudomembranous colitis · *Clostridium difficile* · Rifaximin · *Clostridium difficile*-associated diarrhea

Abstract

Diarrhea is a well-known complication of antibiotic therapy. Rates of antibiotic-associated diarrhea (AAD) vary from 5 to 25%. Some antibiotics are more likely to cause diarrhea than others, specifically, those that are broad spectrum and those that target anaerobic flora. This paper reviews the effects of antibiotics on the fecal flora as well as host factors which contribute to AAD. Clinical features and treatment of AAD are also described. Prevention of AAD rests on wise antibiotic policies, the use of probiotics and prevention of acquisition in the hospital setting. Data from clinical trials suggest that poorly absorbed antimicrobials might have a decreased risk of causing AAD and *Clostridium difficile*-associated disease, as concluded from studies of antibiotics used for preoperative bowel decontamination and poorly absorbed antibiotics used for traveler's diarrhea. Controlled trials would prove this but are not yet available. Probiotics may be a good adjunct to poorly absorbed antibiotics to minimize the risk of diarrhea associated with antibiotics.

Introduction

Diarrhea is a well-known complication of antibiotic therapy. Rates of antibiotic-associated diarrhea (AAD) vary from 5–25% [1]. Rates of AAD vary with antibiotic as well as with the population studied. AAD occurs in up to 29% of hospitalized patients, resulting in lengthened hospital stays, increased costs of medical care, a threefold increase in mortality and a fivefold increase in the acquisition of other nosocomial infections [2–5]. Some antibiotics are more likely to cause diarrhea than others – these are broad-spectrum compared to narrow-spectrum ones, and those that target anaerobic flora compared to those that do not. In addition, multiple antibiotics are associated with an increased risk of AAD. The most severe form of AAD is due to *Clostridium difficile*, which causes colitis and pseudomembranous colitis. The first case of pseudomembranous colitis, at the time described as ‘diphtheritic’ colitis, was reported in 1899 [6]. A young woman developed fatal colitis after gastrointestinal tract surgery. This was clearly in the preantibiotic era. Pseudomembranous colitis as a discrete entity was widely recognized in the 1970s in association with lincomycin and clindamycin therapy, even before the pathogen *C. difficile* was recognized.

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Table 1. Risk factors for AAD and CDAD

Risk factor	Degree of risk	Strength of evidence
<i>Antibiotics</i>		
Oral antibiotics		
Clindamycin	high	excellent
Cephalosporins	high	excellent
Ampicillin	high	excellent
Nonabsorbed antibiotic	low	minimal data to date
Parenteral antibiotics		
	moderate, varies with antibiotic	excellent
Multiple, broad-spectrum antibiotics	high	moderate
Hospitalization		
Gastrointestinal tract surgery	high	excellent
Severity of illness	high	moderate
HIV with immune suppression	high	cofactor
Gastrointestinal procedures, enemas, nasogastric tubes	moderate	unconfirmed
Renal failure	low	not uniform
	low	moderate

Epidemiology

Rates of AAD vary among different populations, and are probably higher for inpatients than outpatients. Two studies of adult patients in US community hospitals documented frequencies of 15 and 22%, respectively [7, 8]. In contrast, febrile neutropenic patients in another study had a frequency of AAD of 29% [9]. Rates of *C. difficile*-associated antibiotic diarrhea (CDAD) also vary – when an outbreak is documented, frequencies can be as high as 26–60% [10, 11]. Predisposing causes such as gastrointestinal tract surgery or the use of high-risk antibiotics such as clindamycin or third-generation cephalosporins can contribute to high rates as well. Other possible factors predisposing to AAD are comorbidity and altered immune status. AAD is common with HIV infection, but this may be a result of frequent antibiotic use rather than immunosuppression per se. In hospitalized patients, risk factors include prolonged hospitalization, gastrointestinal procedures such as endoscopy and enemas, and nasogastric tube feeding, though the latter is controversial [12, 13]. The lowest rates of AAD and CDAD generally occur in outpatients; perhaps they are generally more healthy and less susceptible to disruption of the normal flora (table 1) [14].

Pathophysiology and Antibiotic Risk

C. difficile overgrowth is present in only 15–25% of cases of AAD; thus, other explanations are needed to understand the pathophysiology of diarrhea. There are multiple mechanisms which can account for the diarrhea associated with antibiotics. These include altered fermentation of the flora, changes in dietary fiber and overgrowth of other potential pathogens.

Overgrowth of C. difficile

Antibiotics alter the normal colonic flora, leading to loss of colonization resistance, which is the ability of the normal flora to protect against overgrowth of pathogens, especially when the anaerobic flora are depleted [15]. In CDAD, the altered colonization resistance can allow for the overgrowth of *C. difficile* in the colon. The bacteria produces two toxins which cause disease (toxin A, an enterotoxin, and toxin B, a cytotoxin). The toxins of *C. difficile* inactivate Rho proteins, which results in the loss of cytoskeletal integrity in enterocytes. Cellular damage results in fluid loss, exudation and diarrhea. The most severe form of *C. difficile* diarrhea is pseudomembranous colitis, which can cause severe colitis, toxic colon and rarely colon perforation and death.

Altered Fermentation of Colonic Carbohydrates

The fecal flora ferment unabsorbed carbohydrates with production of short-chain fatty acids (SCFAs). One likely explanation for diarrhea due to antibiotics is altered fecal

flora, which results in changes in SCFA metabolism. Changes can result in acidic pH as well as osmotic changes in the lumen; either or both can contribute to the diarrhea. One study of seven patients with AAD showed reduced fecal concentration of SCFAs as well as changes in rates of production [16]. Since SCFAs stimulate salt and water absorption in the colon, a decrease in their concentration could also contribute to diarrhea.

Dietary Fiber

Dietary fiber may also play a role in AAD. Diarrhea with enteral feeding is well documented and is increased with antibiotics. Antibiotics can impair colonic fermentation of carbohydrates; studies of ampicillin documented effects on lactulose using breath hydrogen analysis and stool output measurements [17]. Lack of dietary fiber in enteral feeding can contribute to diarrhea. A study of pectin, a water-soluble fiber that stimulates epithelial growth in the colon, given to patients on enteral nutrition and antibiotics showed less diarrhea in the fiber-pectin group compared to the fiber-placebo group [18].

Overgrowth of Other Potential Pathogens

Rarely, other pathogenic bacteria have been reported to cause pseudomembranous colitis.

Klebsiella oxytoca. Two French studies of patients with acute AAD used culture of colorectal biopsy to detect *K. oxytoca*, despite negative stool culture [19, 20]. Colonoscopic findings included diffuse and focal colitis and right-sided hemorrhage colitis. The pathogenicity of these strains is still under debate.

Candida. The possible role of *Candida* as a cause of diarrhea has become more apparent lately. Early in the 1990s, two separate papers by Gupta and Ehrinpreis [21] and Danna et al. [22] documented elderly hospitalized patients with diarrhea and significant *Candida* in their stools whose diarrhea responded to antifungal therapy. Two studies of infants and children from India also suggested *Candida* as a pathogen causing diarrhea [23, 24]. However, subsequent studies in children indicated that the role of *Candida* may not be as a pathogen but as overgrowth due to the disruption of flora by antibiotic [25]. A critical literature review of *Candida*-associated diarrhea by Levine et al. [26] cast doubt on its pathogenicity, as many patients with *Candida* in their stools had had prior antibiotic therapy, but conceded that *Candida* species could cause diarrhea in selective clinical settings. Most recently, Krause et al. [27] published a large study analyzing stools from patients with AAD and controls and also concluded that elevated *Candida* counts were a result of

Table 2. Normal fecal flora

a Bacteriological analysis of the normal fecal flora

Bacteria	Log 10 bacteria counts
Enterobacteria	7.4
Enterococci	5.6
Lactobacilli	6.5
Clostridia	5.4
Bacteroides	9.8
Gram-positive anaerobes (eubacteria, bifidobacteria)	10

b Most common species

Anaerobes	Aerobes
Bacteroides	<i>E. coli</i>
Eubacteria	Enterococci
Bifidobacteria	Lactobacilli
Anaerobic cocci	

antibiotic therapy rather than a cause of AAD, an argument bolstered by their finding of no increase in a virulence factor (SAP) in diarrhea stools.

Antibiotics and the Fecal Flora

To understand the role of antibiotics, it is important to understand their effects on the fecal flora. The normal flora consists of a complex bacterial population with 400–500 distinct species of bacteria (table 2a). More than 99% of the total organisms are accounted for by non-spore-forming anaerobic rods [28]; the four major species are *Bacteroides*, bifidobacteria, eubacteria and peptostreptococci [29]. Other common species are *Escherichia coli*, *Streptococcus viridans*, *Streptococcus salivarius* and lactobacilli. Mette et al. [30] clarified the prevalence of species in fecal flora by listing the four most common anaerobes (*Bacteroides* spp., *Eubacterium* spp., *Bifidobacterium* spp. and anaerobic cocci) and three common aerobes (*E. coli* spp., *Enterococcus* spp. and *Lactobacillus* spp.) (table 2b).

The effects of antibiotics on anaerobic and aerobic flora are shown in table 3. These data are from a variety of studies of human volunteers or patients given preoperative antibiotics but receiving various antibiotics by various routes [31–44]. Variations in data can be explained by methodological differences between studies as well as

Table 3. Antibiotic effect on suppression of normal bacterial flora of humans

	Anaerobic flora	Aerobic flora	Comment	Reference
<i>Oral antibiotics</i>				
Tetracycline, doxycycline	minor (log 1–2)	no change to minor ↓	low risk of CDAD	30, 35
Neomycin + erythromycin	major (↓ log 5)	major (↓ log 5)		32
Kanamycin + metronidazole	moderate (↓ log 2)	minimal		33
Neomycin + erythromycin	moderate (log 5)	moderate (↓ log 5)		32
Penicillin	minor ↓			
Erythromycin	major ↓	marked ↓		35
Cephalosporins	major ↓	?		42
Cefoperazone				
Clindamycin	major ↓ (log 7) (log 4–5)	minimal to marked ↓	flora change noted especially in PMC patients	5, 7, 30, 31, 35, 84
Ampicillin + sulbactam	major ↓	minimal ↓	<i>C. difficile</i> correlated with abnormalities of flora	38
Ampicillin prodrugs (bacampicillin, pivampicillin)	moderate ↓	minimal ↓	<i>C. difficile</i> correlated with abnormalities of flora	38
<i>Parenteral antibiotics</i>				
Imipenem/cilastatin (i.v.)	minimal to moderate ↓	minimal to marked ↓	low rates of <i>C. difficile</i>	36, 41
Azlocillin	minimal to moderate	minimal to moderate	variability among volunteers	39
Ampicillin + sulbactam (i.v.)	moderate ↓ (log 3)	normal to minimal		34, 35
Temocillin (i.v.)	marked ↓	unknown	↓ <i>Enterobacter</i> likely low rates of <i>C. difficile</i>	37
Cephalosporins				
Cefoxitin (i.v.)	moderate ↓	moderate ↓		35
Cefoperazone (i.v.)	minimal	unknown	asymptomatic <i>C. difficile</i> in 1 volunteer	42
Ceftriaxone (i.m.)	minor to marked ↓	marked ↓	2 volunteers had <i>C. difficile</i> , 1 with diarrhea	43 44
Clindamycin	marked ↓	minor ↓		41
Aztreonam	minor ↓	minor ↓		41

PMC = Pseudomembranous colitis.

Data were summarized from the references noted in the table. Some articles present disparate data. In general, a minor decrease in flora was a decrease by 1–2 logs of bacteria; a moderate decrease was a decrease by 2–4 logs, and a major decrease was a decrease by 4–5 logs.

variations among studied populations. Nonetheless, some trends can be observed. Antibiotics with profound effects on suppression of anaerobic flora are the same ones which have an increased incidence of *C. difficile*-associated disease: clindamycin (p.o. or i.v.), ampicillin, sulbactam and broad-spectrum cephalosporins. Broad-spectrum parenteral antibiotics (e.g. ceftriaxone) have a lower but still significant risk. Antibiotics with less anaerobic suppression (tetracycline, penicillin) appear to have a lower risk of CDAD. In one study, a cephalosporin given intravenously showed minimal anaerobic suppression; this increased when given orally [43].

Clearly, the route of administration of antibiotics can affect the rates of AAD. Rates of diarrhea are lower with intravenous than oral antibiotics. Specific rates of diarrhea for all individual antibiotics are not available. Some

reports suggest that antibiotics with biliary secretion may have higher rates of AAD. Clindamycin remains a high-risk antibiotic [31]. A recent study of an epidemic in a Veterans Affairs hospital demonstrated dramatic declines in CDAD rates when the use of clindamycin was restricted [44].

In general, these high-risk antibiotics cause less suppression of aerobic bacteria. Decreases in aerobes appear to have less effect on colonization resistance (table 3). Interestingly, colonic lavage without antibiotics decreases colonic bacterial counts by log 2 [40].

A recent prospective study of AAD and CDAD in five Swedish hospitals showed an increased risk of AAD with cephalosporins, clindamycin and broad-spectrum penicillins [42]. While CDAD is well recognized as the most common nosocomial gastrointestinal pathogen, some

cases occur in the ambulatory setting. A retrospective study by Levy et al. [14] of enrollees in four community based HMOs (health maintenance organizations) showed a prevalence of 12 per 100,000 person-years. This is less than in the hospital setting, but the rate of testing for *C. difficile* is also lower – only 5% of patients with AAD were tested for *C. difficile*. In the study of Levy et al. [14], first- and third-generation cephalosporins (cephaloxin and cefixime) had a higher risk than other antibiotics. High rates with second- and third-generation cephalosporins have been shown by others, as well as with the use of multiple antibiotics [45–47]. It is likely that the key factor in antibiotic risk is the effect of the individual antibiotic on the colonic flora, specifically the anaerobic flora. Even metronidazole, used to treat *C. difficile*, can cause AAD and CDAD. Only vancomycin seems to have a low risk of CDAD – a single case report in a renal dialysis patient is less than convincing [48].

Nonabsorbable antibiotics are appealing because they have fewer systemic side effects and may be safer for children and pregnant women as well as in patients with renal and hepatic dysfunction. One such antibiotic, aztreonam, showed little effect on anaerobic flora in human volunteers, producing most of its effect on the aerobic flora [49, 50]. A trial showed efficacy of aztreonam for traveler's diarrhea, where most pathogens are aerobes [51]. While there are no data on rates of AAD for nonabsorbable antibiotics and *C. difficile*, these would likely be decreased. Given the preservation of the anaerobic flora, another poorly absorbed antibiotic, bicozamycin, has efficacy in traveler's diarrhea with its major effect being on fecal aerobes [52].

Rifaximin, a nonabsorbable antibiotic and a derivative of rifamycin which has been widely available in Europe since the 1980s, was shown early on to be active against anaerobic bacteria such as *Bacteroides* spp. Stool levels are high [53]. The drug has efficacy against both aerobes and anaerobes in fecal flora. It has been used in traveler's diarrhea [54], where pathogens are primarily aerobic. Its efficacy against anaerobic flora led to an in vitro study of its efficacy against 93 different strains of *C. difficile*. Seventy-four percent of the strains were susceptible to rifaximin, compared to 100% for vancomycin and metronidazole [55], suggesting a real potential use for this drug for the treatment of gastrointestinal diseases and possibly *C. difficile*. When used as an adjunct to cefotaxime in patients undergoing colonic surgery, fewer postoperative complications occurred and intestinal function recovered more quickly [56]. However, clinical trials would be necessary before concluding that the risks of AAD and

CDAD are substantially decreased with this antibiotic compared to others [57].

Rates of diarrhea also vary with host factors. Some people are more likely to develop diarrhea than others; predisposing factors are extremes of age (under 6 or over 65), severe underlying disease, chronic intestinal disorders, prior history of AAD, gastrointestinal tract surgery and nasogastric tube feeding [58]. Surgical patients are at increased risk for CDAD. In a 2-year prospective study, *C. difficile* accounted for 3% of all postoperative infections. The most commonly implicated individual antibiotics were ciprofloxacin (19%) and cefoxitin (16%). Of note, 16% of patients developed CDAD only after administration of perioperative antibiotics [59]. Factors not significantly associated with AAD are gender, antibiotic dose and inflammatory bowel disease. *C. difficile* can also be caused by nonantibiotic factors such as cancer chemotherapy, and rarely, sporadic cases do occur.

Clinical Features

Diarrhea or loose stools can start after even a few doses of antibiotics, or as long as 6 weeks after the cessation of antibiotics. Most patients develop AAD while still on antibiotics. Symptoms of AAD are watery diarrhea and cramping. Severe diarrhea with bleeding, fever and abdominal pain suggests colitis, the most severe form of AAD, which is almost always related to *C. difficile*.

Diagnosis

AAD is suspected in anyone who develops diarrhea while on antibiotics or following recent (previous 6–8 weeks) antibiotic therapy. Diarrhea that develops in hospital is almost never due to other enteric pathogens or ova and parasites, though causes in addition to antibiotics and *C. difficile* are other medications and tube feeding. Diagnosis of *C. difficile* relies on documenting *C. difficile* toxin A or B in the stool and exclusion of other causes of diarrhea. Culture positivity does not indicate disease since a carrier state can exist. Toxin B tissue culture assay is the gold standard, but it is expensive and time consuming. Thus, many hospitals are now using enzyme immunoassay tests for toxin A or toxin B or both. Fecal leukocytes are nonspecific as they indicate inflammation but not a specific cause. Colonoscopy or flexible sigmoidoscopy can give an immediate diagnosis of colitis (fig. 1), but this is rarely necessary.

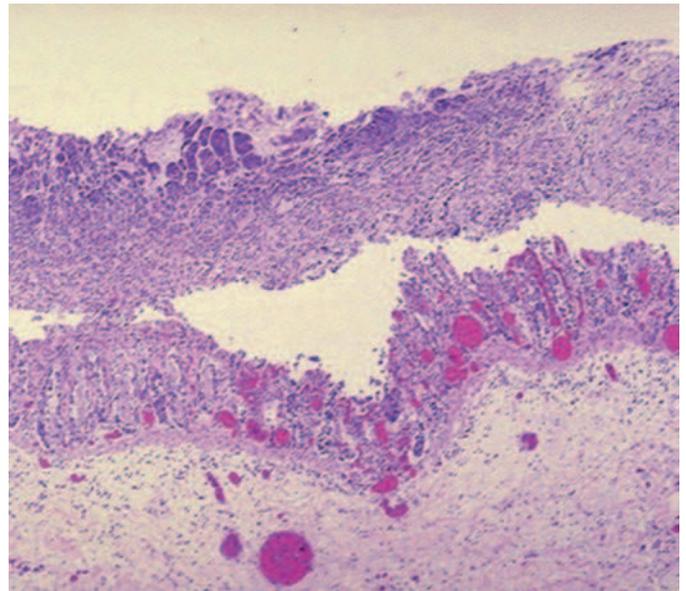


Fig. 1. Typical presentation of pseudomembranous colitis. At endoscopy (left), the mucosal surface of the colon appears hyperemic and almost completely covered by a yellow-green exudate. The mucosa itself is somewhat eroded. Microscopically (right), the pseudomembrane is composed of inflammatory cells, necrotic epithelium, and mucus in which the overgrowth of microorganisms usually takes place.

Treatment of AAD

When AAD is mild, the best first step is to discontinue the inciting antibiotics. Most cases will resolve spontaneously. Some advocate restriction of dietary carbohydrate as well [60]. More serious cases require specific therapy against the pathogen. Antidiarrheals and opiates should be avoided.

Treatment of CDAD

The two most frequently used antibiotics are metronidazole and vancomycin. Metronidazole is recommended as first-line therapy because of the risk of development of vancomycin-resistant enterococci with vancomycin use, as well as its much higher cost. Metronidazole and vancomycin have similar efficacy, though in one study, symptoms resolved sooner with vancomycin [61]. Metronidazole is given orally for 10 days, at a dose of 1 g per day. Vancomycin is given orally for 10 days; doses vary from 500 mg/day to 2 g/day. For mild to moderately severe CDAD, low-dose vancomycin is as effective as high-dose vancomycin. Vancomycin use is generally restricted to

patients who cannot take metronidazole (allergy, intolerant to side effects, pregnancy) or in those in whom metronidazole has failed. Other antibiotics have been used. One study showed similar cure rates for vancomycin (94%), metronidazole (94%), teicoplanin (96%) and fusidic acid (93%) [62].

Rifaximin is a derivative of rifamycin SV. The drug is not absorbed in the gastrointestinal tract and has activity against both Gram-positive and Gram-negative aerobic and anaerobic bacteria by inhibiting bacterial RNA synthesis. Active drug reaches the intestinal lumen. Rifaximin is widely used in Europe, and clinical trials have shown its efficacy in infectious diarrhea in children and adults, colonic diverticular disease, *H. pylori* eradication and treatment of hepatic encephalopathy [57]. It has also been shown to be effective in the treatment of traveler's diarrhea [63]. The antibiotic is active against a variety of intestinal pathogens in vitro, including *Campylobacter jejuni*, *Yersinia* spp. and *C. difficile*. In one study, 56 *C. difficile* strains were tested, and 34 strains (60.7%) were sensitive to rifaximin, compared to 27 strains (48.2%) with the same concentrations of rifampicin [64]. In the previously cited study, the efficacy of rifaximin against 74% of 93 *C. difficile* strains isolated from patients in Ital-

ian hospitals [55] led to a clinical trial that compared treatment of 20 patients with *C. difficile* pseudomembranous colitis using rifaximin, 200 mg thrice daily orally for 10 days, with vancomycin, 500 mg twice daily orally for 10 days. All 10 patients treated with vancomycin were cured, but there was 1 treatment failure in the rifaximin group. Thus, overall efficacy was 100% for vancomycin and 90% for rifaximin in this small trial. However, the time to disappearance of *Clostridium* toxins was more rapid in the vancomycin group (4.8 ± 1.8 days) compared to the rifaximin group (8.1 ± 1.8 days) [65]. The authors conclude that rifaximin can be used as a replacement for vancomycin when the use of the latter is not possible.

Recurrent CDAD is a particularly difficult problem which occurs in up to 20% of patients. Most patients will need repeated antibiotic treatment with metronidazole or vancomycin. Treatment strategies include pulsing or tapering antibiotics and adding probiotic agents, such as *Lactobacillus GG* or the nonpathogenic yeast *Saccharomyces boulardii* [66].

Prevention of AAD and CDAD

The cornerstone of prevention of AAD and *C. difficile* disease is wise antibiotic use policies. Restricting the use of high-risk antibiotics such as clindamycin has been shown to control hospital epidemics of *C. difficile*. In hospitals, hand washing and enteric precautions can minimize person-to-person transmission of *C. difficile*.

As diarrhea is a known and common complication of antibiotic therapy, many approaches have been used for prevention. The use of probiotics administered with the antibiotic is appealing. A probiotic is a live microbial food supplement that beneficially affects the host animal, improving its microbial balance [67]. Certain probiotics can reduce the risk and duration of diarrhea [68]. Agents used in prevention of AAD include lactobacilli, such as *Lactobacillus GG*, *Streptococcus faecium*, bifidobacteria and the yeast *S. boulardii*.

Lactobacilli used have included a commercial mixture of *Lactobacillus acidophilus* and *Lactobacillus bulgaricus* (Lactinex®) and *Lactobacillus GG*. decreased AAD in adults receiving amoxicillin (3.3%) compared to placebo (21%), but the difference was not statistically significant [69]. A study in children receiving amoxicillin failed to demonstrate a significant decrease in AAD with this probiotic mixture [70]. Variability in lots may explain differences in efficacy in many trials [71]. *Lactobacillus GG* may have a better efficacy than this probiotic mixture in

the prevention of AAD. Adults taking erythromycin and children taking oral amoxicillin had fewer diarrhea stools with *Lactobacillus GG* compared to placebo [72, 73]. In a controlled trial, prophylactic *Lactobacillus GG* also decreased AAD when given with antibiotics to children with respiratory infections [74]. In a recent study of *Lactobacillus GG* added to a week of standard anti-*H. pylori* triple antibiotic therapy, diarrhea was decreased compared to the placebo group, as was the symptom of bloating [75].

Bifidobacteria are part of the normal flora. Recent interest in this organism resulted from studies of its use in acute diarrhea in hospitalized infants. *S. faecium* (SF68) is a nonpathogenic nontoxigenic *Streptococcus*, available in some European countries as Bioflorin®, which has been shown to decrease AAD compared to placebo in two separate studies in adults [76, 77]. Its efficacy appears to be limited based on data from these two studies.

S. boulardii is a nonpathogenic yeast with proven efficacy in the prevention of AAD in multiple trials in adults and children [7, 8, 78]. The yeast may compensate for changes in microbial fermentation in response to antibiotic therapy [78]. Its use to prevent AAD in at-risk patients should be a good adjunct to other methods such as limiting *C. difficile* overgrowth [1].

Prevention of acquisition of *C. difficile* in a hospital setting includes careful attention to hand washing, disposable gloves, wise antibiotic policies and enteric precautions.

Conclusion

Data from a variety of clinical trials does suggest that poorly absorbed antimicrobials have a decreased risk of causing AAD and CDAD. These trials include studies of such antibiotics for preoperative bowel decontamination for gastrointestinal tract surgery as well as studies of antibiotics used for traveler's diarrhea (aztreonam, bicozamycin and rifaximin). Such antibiotics are appealing because systemic side effects are decreased. Their efficacy in traveler's diarrhea is a result of efficacy against aerobic pathogens. Rifaximin shows significant suppression of anaerobes, but its efficacy against many strains of *C. difficile* may offset its risk of *C. difficile* diarrhea and in fact decrease it. The use of probiotics as an adjunct to poorly absorbed antibiotics would be an interesting area for future study of minimizing the risk of diarrhea.

One of the leaders in the field, Carl Erik Nord, stated that 'ecological effects are difficult to predict and clinical studies of new antibiotics should include investigations of

their impact on the normal human intestinal flora' [41]. Nonetheless, the risk of CDAD may be difficult to predict accurately as other factors besides antibiotics contribute to this risk.

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Management of Hepatic Encephalopathy: Role of Rifaximin

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Key Words

Ammonia · Bacterial flora · Benzodiazepines, endogenous · Encephalopathy, hepatic · Liver cirrhosis · Rifaximin

Abstract

Hepatic encephalopathy (HE) is a neuropsychiatric syndrome, which develops in patients with acute or chronic liver failure. It is widely accepted to be due to impairment of hepatic clearance of toxic products from the gut such as ammonia. Accumulation of ammonia induces a glutamate neurotoxicity leading to an increased tone of the γ -aminobutyric acid A (GABA-A) receptor system in the brain which results in HE. Factors either increasing the ammonia levels (protein load, constipation, sepsis, or gastrointestinal bleeding) or potentiating the functional activity of the GABAergic system [natural benzodiazepine-like compounds (NBZDs) or exogenous benzodiazepines] may act as precipitating factors of HE. NBZDs are present in trace amounts in the blood of normal subjects and have been found to be increased in the blood of patients with liver cirrhosis, with or without HE. These compounds may derive either from the diet since they have been found in plants, vegetables and animals or from gut bacteria. The observation that intestinal bacterial flora is involved in the production of both primary

agent of HE (ammonia) and precipitating factors (NBZDs) suggests that the use of nonabsorbable antibiotics such as rifaximin may be useful in preventing episodes of HE in patients with liver cirrhosis.

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Introduction

Hepatic encephalopathy (HE) is a neuropsychiatric syndrome characterized by a general depression of the central nervous system, which occurs in fulminant hepatic failure or as a complication of liver cirrhosis. The neurological symptoms of HE range from minimal abnormalities of intellectual functions to deep coma [1]. From a pathogenetic point of view, HE is considered secondary to the accumulation of toxic products such as ammonia in extracellular fluids. These toxins, which increase in blood during severe hepatocellular disease and portal-systemic shunting because of the decreased clearance capacity by the liver, have been suggested to play a key role in HE. There are, however, other compounds, such as benzodiazepines [either natural benzodiazepine-like compounds (NBZDs) and the exogenously administered drugs] that are involved in HE and are today considered as precipitating factors.

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Ammonia

There have been debates and controversies for decades about ammonia, the first toxin claimed to be the primary agent of HE [1–4], but it is now recognized as the major factor of this syndrome [5, 6]. Ammonia toxicity does not only seem to be related to an altered brain metabolism, but more specifically to its ability to damage astrocytic cells by a glutamate-related neurotoxicity [7–9], which in turn may alter the entire γ -aminobutyric acid A (GABA-A) receptor system [10]. Moreover, ammonia has been shown to induce an increased production of neurosteroids that exert a positive modulatory effect on the GABA-A receptors [11] and this phenomenon has been ascribed to the upregulation of peripheral benzodiazepine receptors present in HE [12, 13]. These complex metabolic cerebral events, which affect both astrocytes and neurons, result in an imbalance in the functional activity of excitatory and inhibitory receptor systems leading to a prevalence of the latter thanks to an increased tone of the GABA-A receptor system [14, 15]. This is represented by a large supramolecular entity [16], which includes GABA receptors, benzodiazepine recognition sites, chloride ionophores and endogenous modulators. The plasticity of GABA-A receptors, that is up- and downregulation in the number of receptors, is regulated by the levels of neurotransmitters at the synaptic level. In the case of HE, the upregulation of GABA-A receptors, interpreted as an expression of a denervation supersensitivity phenomenon [14] with a consequent increase in GABAergic tone [14–17], is explained by a decreased presence of GABA at the neuronal level [18] rather than by an increased presence of GABA in blood and brain (gut-derived hypothesis) [17]. Indeed, studies performed in animal models of encephalopathy have shown that the mild stage of encephalopathy is characterized by a 45–50% increase in the number of both low and high affinity GABA receptors without changes in the affinity constants [14, 19] and also by an increased number of benzodiazepine receptors of the central type [20]. These findings have recently been confirmed in humans by *in vivo* studies using positron emission tomography [21] and ¹H magnetic resonance spectroscopy [22] in which an increased presence of central-type benzodiazepine receptors in the brain of patients with recurrent HE and a decreased cortical GABA level were demonstrated. These alterations seem to explain the supersensitivity to the administration of tranquillizers and in particular of benzodiazepines in patients with acute or chronic liver diseases. This phenomenon, originally attributed to an altered metabolism of benzodiazepines in such patients, is

now recognized to be mainly due to an increased central nervous system sensitivity to this class of drugs [23, 24].

Ammonia must be regarded as the main pathogenetic factor of HE but a variety of other events may precipitate the syndrome in patients with severe liver disease. Indeed, there are precipitating events which act by inducing an increased concentration of ammonia in brain tissues (diuretic therapy, hypokalemia, hyperazotemia, constipation, protein overload, sepsis) [1] and factors which enhance the latent but already present increased tone of the GABAergic system in the brain (NBZDs or exogenous benzodiazepines) [2].

Natural Benzodiazepines

Commercial benzodiazepines are widely used drugs for the treatment of anxiety and sleep disturbances. NBZDs such as diazepam and nordiazepam and other unknown benzodiazepine-like compounds are naturally present in several plants and vegetables [25–28], in different animal species and in humans [29–31]. Moreover, both plasma and brain contain other compounds with benzodiazepine-like activity called ‘endozepines’ apparently produced in mammalian cells [32, 33]. The observation that NBZDs are present in human brain samples stored since 1940 [30], when benzodiazepines were not yet synthesized, clearly indicates that NBZDs do not derive from environmental pollution by synthetic benzodiazepines, which entered the market only in 1959. Several attempts to find out the endogenous biosynthetic pathways of NBZD production have been made [31–34]. However, an endogenous biochemical pathway of NBZDs has not yet been identified in mammalian cells [33] while it has been found that microorganisms like *Streptomyces* and *Penicillium* can synthesize molecules like anthramycin and cyclopeptine, respectively, both containing the basic 1,4-benzodiazepine structure [35]. Moreover, evidence has been provided that some gut bacteria, such as *Acinetobacter lwoffii*, can produce precursors of benzodiazepine receptor ligands [36]. Since NBZDs have been found in food [25–28, 37], it could be speculated that at least part of these compounds found in the human body could be of alimentary source.

NBZDs are present in trace amounts in the blood of normal subjects but they may rise severalfold in the blood of patients with liver cirrhosis, with or without encephalopathy [2, 38]. Increased levels of these compounds are inconstantly present in cirrhotic patients. A significant albeit weak correlation [38] between circulating NBZDs

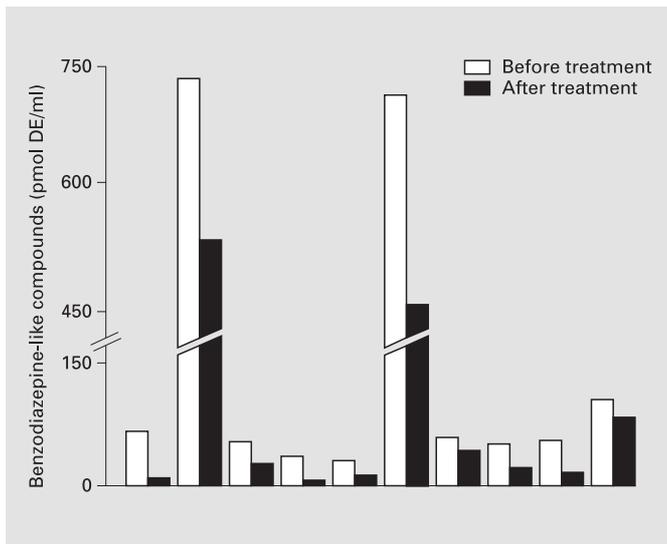


Fig. 1. Serum concentrations of benzodiazepine-like compounds in patients with cirrhosis of the liver before and after treatment with rifaximin (800 mg/day) lasting 7 days [normal values: 9.5 ± 2.9 pmol diazepam equivalents (DE)/ml, mean \pm SEM]. Each couple of columns refers to a single patient. One-way ANOVA showed a significant difference in the values before and after treatment ($p < 0.05$) (from Zeneroli et al. [39]).

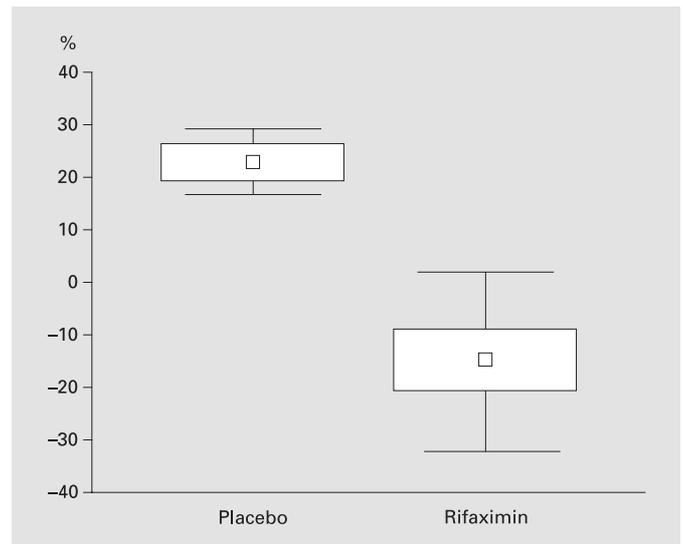


Fig. 2. Percent variation of beta rhythm in cirrhotic patients after a 1-week period of treatment with rifaximin (200–400 mg t.i.d.) or placebo. The relative beta power of the EEG decreased in the rifaximin-treated group (Wilcoxon paired test: $Z = 2.1$, $p = 0.03$), but not in the placebo-treated group ($Z = 1.07$, $p =$ nonsignificant) (from Del Piccolo et al. [40]).

in patients and the degree of HE does exist, suggesting a pathogenetic role of NBZDs in HE. In a large number of patients we [2] found, however, that NBZD levels correlate with the severity of the liver dysfunction but not with the degrees of HE. It is, therefore, likely that NBZDs represent an occasional precipitating agent of the syndrome rather than a 'true' pathogenetic factor.

Although food does contain NBZDs, it is unlikely that the minute amounts of dietary pharmacologically active substances (in the range of nanograms of diazepam equivalent/gram) cause any appreciable pharmacological effect [37]. However, it should be taken into account that in normal subjects benzodiazepines have a low clearance rate so that a chronic intake of even small amounts of NBZDs could lead to their accumulation in blood and brain, thus exerting a dietary influence on brain function and behavior. This phenomenon could be magnified in cirrhotic patients whose liver displays reduced clearance capabilities. Since food can only partially contribute to the increased NBZD levels in blood and brain of cirrhotic patients, it has been suggested that bacterial flora could also contribute to NBDZ formation [31, 36].

In order to ascertain whether bacterial flora is implicated in the synthesis of these compounds, we measured serum NBZD levels in patients with liver cirrhosis before

and after the reduction of bacterial flora with rifaximin, a poorly absorbed antibiotic and found – after treatment – a 40% decrease (fig. 1) [39]. Our results are in line with the finding of Del Piccolo et al. [40], who observed that the EEG beta activity, which is well known to be increased by benzodiazepines, is significantly reduced in patients with cirrhosis and HE treated with rifaximin (fig. 2). The results of both studies are consistent with the hypothesis that the intestinal bacterial flora is at least partially involved in the production of NBZDs.

Rifaximin in the Management of HE

Both dietary and endogenous ammoniagenic substrates are removed from the intestinal lumen by the osmotic cathartic action of nonabsorbable disaccharides such as lactulose and lactitol. These compounds are currently the main therapeutic agents for chronic HE. The efficacy of oral lactulose for the treatment of HE has been established in controlled trials [41–43]. Besides having a cathartic effect, lactulose lowers the colonic pH as a result of the production of organic acids by bacterial fermentation. The decrease in pH creates an environment that is hostile to the survival of urease-producing intestinal bac-

Table 1. Controlled clinical trials with rifaximin in the treatment of HE

Year	Authors	Ref.	Comparative agent (study design)	Duration of treatment	Evaluation criteria	Overall assessment
1984	De Marco et al.	49	Paromomycin (open)	6–12 days	NH ₃ , state of consciousness, intellectual functions, behavior, neurological symptoms	E: Rif \cong E Par T: Rif \geq Par
1985	Testa et al.	50	Paromomycin (open)	5 days	Antibacterial assays in vitro, NH ₃ , number connection test	E: Rif \geq Par T: Rif \cong Par
1991	Di Piazza et al.	51	Neomycin (double-blind)	7 days	Flapping tremor, bradylalia, patient self-evaluation, visual evoked potential reaction, trial making test	E: Rif \cong Neo T: Rif \cong Neo
1991	Pedretti et al.	52	Neomycin (double-blind)	21 days	NH ₃ , mental status, Reitan test, asterixis, EEG, PSE sum, PSE index	E: Rif \geq Neo T: Rif \geq Neo
1992	Parini et al.	44	Paromomycin (open)	10 days	NH ₃ , state of consciousness, intellectual functions, behavior	E: Rif \geq Par T: Rif \cong Par
1992	Festi et al.	53	Neomycin (open)	21 days	NH ₃ , asterixis, Reitan test, EEG	E: Rif \cong Neo T: Rif \cong Neo
			Lactulose (open)	21 days	NH ₃ , asterixis, Reitan test, EEG	E: Rif. \cong Lac T: Rif \geq Lac
1993	Massa et al.	54	Lactulose (double-blind)	15 days	Mental status, 'A' cancellation test, Reitan test, EEG, HE severity	E: Rif \geq Lac T: Rif > Lac
1993	Fera et al.	55	Lactulose (double-blind)	14 consecutive days each month for 3 months	Mental status, asterixis, cancellation test, Reitan test, EEG, NH ₃ , PSE severity	E: Rif > Lac T: Rif > Lac
1993	Bucci and Palmieri	56	Lactulose (double-blind)	15 days	Mental status, asterixis, cancellation test, Reitan test, EEG, NH ₃	E: Rif > Lac T: Rif > Lac
1997	Miglio et al.	58	Neomycin (double-blind)	14 consecutive days each month for 6 months	Disturbances in speech, memory, behavior and mood, gait, writing, asterixis, NH ₃ , Reitan test	E: Rif \cong Neo T: Rif \geq Neo
2000	Williams et al.	59	3 different rifaximin dose regimens: 200 mg \times 3 400 mg \times 3 800 mg \times 3 (double-blind)	7 days	HE index, mental status, asterixis, Reitan test, EEG, NH ₃	E: Rif 800 mg \times 3 \geq other doses No clear dose response explained by the short treatment period T: good for all doses
2003	Loguercio et al.	61	Rif + sorbitol Rif + lactitol Lactitol (double-blind)	14 consecutive days each month for 3 months	Mental status, asterixis, blood NH ₃ , number connection test	E: Rif + Lat = Rif + Sor > Lat T: Rif + Lat = Rif + Sor = Lat
2003	Mas et al.	62	Lactitol (double-blind)	5–10 days	HE index, mental status, asterixis, NH ₃ , number connection test, EEG	E: Rif \geq Lat T: Rif = Lat

E = Efficacy; T = tolerability; Rif = rifaximin; PSE = portal systemic encephalopathy; Par = paromomycin; Neo = neomycin; Lac = lactulose; Lat = lactitol; Sor = sorbitol.

teria and may promote the growth of non-urease-producing lactobacilli, resulting in a reduced production of ammonia in the colonic lumen.

Antibiotics with activity against urease-producing bacteria, such as neomycin [42], paromomycin [44] or metronidazole [45], also reduce the production of intestinal ammonia and have proved to be of value. Vancomycin has also been used in patients with lactulose-resistant chronic encephalopathy [46]. The efficacy of neomycin is similar to that of lactulose [42]. However, a small percentage of this drug is absorbed from the gastrointestinal tract and may cause ototoxic and nephrotoxic effects, especially with continuous use over several months [47]. This drug should be used with particular caution by patients with renal insufficiency. The efficacy of metronidazole for

1 week is similar to that of neomycin [45], although the occurrence of gastrointestinal disturbance and other systemic side effects limit the use of this agent for longer periods.

Rifaximin is a synthetic rifamycin derivative, which acts by inhibiting bacterial ribonucleic acid (RNA) synthesis [48]. It is virtually unabsorbed after oral administration and is, therefore, used primarily to treat gastrointestinal infections. Rifaximin possesses a broad spectrum of antimicrobial activity, covering Gram-positive and Gram-negative bacteria, both aerobic and anaerobic [49]. Several studies [44, 49–62] have shown that in patients with HE rifaximin displays an efficacy similar to that of lactulose and neomycin (table 1). A recently published study [62] compared the efficacy and safety of

rifaximin to that of lactitol. This large prospective, randomized, double-blind, double-dummy, controlled trial confirmed that this rifamycin derivative is as effective as lactitol in the treatment of grade I–III HE. Rifaximin, however, was significantly more effective in reducing plasma ammonia levels.

Altogether, these data clearly show that rifaximin represents a good alternative to nonabsorbable disaccharides

and may actually be effective in the management of patients resistant to their administration. Taking into account that circulating NBDZs are significantly reduced during administration of this antibiotic [39], cyclic administration of rifaximin – by preventing blood accumulation of both pathogenetic (e.g. ammonia) and precipitating (e.g. NBDZs) factors – might be capable of reducing the number of acute episodes of HE.

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Note Added in Proof

After the submission of the manuscript three interesting papers [1–3] dealing with the management of hepatic encephalopathy have been published. A Cochrane systematic review [1] evaluating 30 randomized controlled trials did conclude that antibiotics appear to be superior to nonabsorbable disaccharides in improving symptoms of portal systemic encephalopathy. The authors also emphasized that there is insufficient high-quality evidence to support the use of lactulose or lactitol. A combination of a disaccharide and an antibiotic has been suggested, but not consistently demonstrated to be beneficial [2]. Finally, the use of probiotics has been proposed [3], whose administration could actually follow that of antibiotics.

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Management of Inflammatory Bowel Disease: Does Rifaximin Offer Any Promise?

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Key Words

Rifaximin · Ulcerative colitis · Crohn's disease · Intestinal microflora

Abstract

An increasing number of both clinical and laboratory-derived observations support the importance of luminal components in driving the inflammatory response in ulcerative colitis and Crohn's disease. Although its role is unclear, antibiotic therapy is commonly used in clinical practice for the treatment of moderately to severely active ulcerative colitis. Metronidazole and/or ciprofloxacin are currently employed in active Crohn's disease, particularly in patients with colonic involvement and with perianal disease. Rifaximin, a rifamycin-derived antibiotic, is characterized by a wide range of antibacterial activity and a very low systemic absorption. Some preliminary data show its efficacy in severe active ulcerative colitis, pouchitis and prevention of postoperative recurrence in Crohn's disease.

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Role of Intestinal Microflora in Inflammatory Bowel Disease

Although the pathogenesis of inflammatory bowel disease (IBD) remains unclear, increasing evidence suggests that the enteric microflora plays a central role in this process. The distal ileum and the colon are the areas with the highest bacterial concentrations and represent the sites of inflammation in IBD; similarly pouchitis appears to be associated with bacterial overgrowth and dysbiosis.

Recent experimental data, coming particularly from animal models of IBD, are consistent with the hypothesis that gut flora and bacterial products are implicated in the initiation and/or perpetuation of chronic intestinal inflammation. Purified bacterial products can initiate and perpetuate experimental colitis [1, 2].

The leading hypothesis for the development of chronic intestinal inflammation is that an abnormal immune response to normal flora might be crucial. This loss of tolerance might be due to a lack of regulatory mediators or cells, or a breakdown in barrier function which makes possible the access of inflammatory bacterial products to the local immune system, thereby overwhelming the normal regulation [3]. These possibilities were supported by

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data obtained from several studies in IBD patients, reporting an important role for T cells in the proliferative response to intestinal flora [3], T cell-mediated immune responses to different autologous and heterologous species of bacteria from intestinal flora regulated by a complex network of T cell specificity [4], and enhanced IgG levels against cytoplasmic proteins from commensal bacteria in active IBD. In patients with Crohn's disease (CD) diversion of the fecal stream determines a decrease in disease activity, with disease recurrence occurring after restoration of the fecal stream [5]. Moreover, studies have demonstrated the ability of luminal contents, presumably dominated by bacteria or their products, to trigger postoperative recurrence in the terminal ileum within a few days, providing further evidence of the role of enteric microflora. In fact, D'Haens et al. [6] showed that the exposure of the mucosa of the excluded normal ileal loop to autologous fecal material in CD quickly (8 days) activates the mucosal immune system: biopsy specimens showed morphological evidence of focal inflammation with recruitment of inflammatory cells and differentiation of mononuclear cells into functional macrophages and epithelioid cells. Another study has tested the effect of the ultrafiltrated ileal effluent when infused into defunctioned colon; no signs of mucosal lesions were found, suggesting that the intact bacteria and large dietary particles may be responsible for the recurrence of inflammation. The most compelling evidence, however, comes from studies conducted in animal models of colitis; spontaneous colitis that consistently develops in many transgenic and knockout mutant murine models of colitis may not occur when these lines are maintained in a germ-free environment [7].

Taken together these experimental and clinical observations support the hypothesis that there is no gut inflammation without bacteria, and that the manipulation of enteric microflora may represent a possible therapeutic approach in IBD.

For many years, investigators have tried to find out whether a specific pathogen could determine IBD. For instance, much attention has been paid to the role of *Mycobacteria* at the onset of CD [8] and more recently it has been suggested that a particular subtype of *Escherichia coli* could play a pathogenic role in CD [9]. The presence of *Shigella* or *Shigella*-like toxin, *Salmonella* and *Yersinia* has been investigated as a possible cause of ulcerative colitis (UC), whereas *Clostridium difficile* toxin has been associated with disease exacerbation [10]; a similar role has been suggested for *Salmonella* infection perhaps associated with a diminished protective activity of the

mucus [11]. More recently, high serum antibody titers to the outer membrane protein of *Bacteroides vulgatus* were found in patients with UC [12], but all these results have been rather inconclusive. Since *E. coli* is the predominant aerobic Gram-negative species of the normal intestinal flora, much more attention has been paid to a possible role of its subtypes. Besides commensal bacteria, certain specific strains possess virulent properties and cause disease in humans; the diarrheagenic subtypes of *E. coli* belong to this latter group, showing properties such as adherence to the gut mucosa, production of enterotoxins and cytotoxins and tissue invasion [13].

The presence of *E. coli* in patients with UC has been investigated, and it has been reported that *E. coli* could be detected only in a small proportion of tissue samples [14, 15]. Studies on mucosal adhesion of pathogenic bacteria in UC are controversial. A significantly enhanced adhesion of isolates of *E. coli* from stool specimens and rectal biopsies from UC patients to buccal epithelial cells was found in comparison with patients with infectious diarrhea or normal controls. The adhesive properties were similar to those of pathogenic intestinal *E. coli*, suggesting that virulent *E. coli* strains might participate in the pathogenesis of UC [16, 17]. Another study reported adherence of only the DAEC and EAaggEC *E. coli* subtypes to rectal mucosa; however, no differences in adhesion could be found between UC patients and controls [18]. Adherence of a different species (EHEC) has also been described [19]. Using a hybridization in situ technique, a significantly higher number of bacteria was found within the mucus layer and not adherent to the surface of the epithelium in UC patients compared to controls, independently of the degree of inflammation. The bacteria belong, most likely, to a variety of species regarding the broad specificity of the probe used in this study [20]. In summary, incomplete information and controversies exist about the role, the adherent properties, and the subtypes of *E. coli* which might be important in the pathogenesis of IBD.

An alternative hypothesis suggests that alteration in normal intestinal ecology can cause inflammation through impairment of epithelial cell metabolism. Colonic anaerobic bacteria are able to break down the ingested carbohydrates and proteins through the process of fermentation into short-chain fatty acids, which are the main source of energy for colonocytes [21]. It has been postulated that a deficiency of this energetic support might lead to the onset of colitis [22]. Furthermore, in patients with active UC there is an overproduction of hydrogen sulfide, a metabolite very toxic for the intestinal mucosa, which seems to be related to an excess of sulfate-

reducing bacteria (*Desulfibrio desulfuricans*) in fecal samples [23, 24]. This theory is supported by the evidence that administration of sulfated polysaccharides (carrageenan) in guinea pigs induces a chronic colonic inflammation whose features are similar to human UC [25] and that treatment with 5-ASA is able to reduce fecal concentration of sulfide [26]. It is, therefore, evident that some bacteria do locate in the mucus and might possibly act by degrading its protective structure, leading as a consequence to mucosal invasion.

Therefore, the unresolved question remains of whether chronic, recurring inflammation is the result of a persistent infection with a specific pathogen, an exaggerated exposure to resident normal luminal bacteria products because of increased intestinal permeability or alteration of mucus composition, or an abnormally aggressive immune response to luminal components.

Antibiotic Treatment in IBD

Leaving from these data, few clinical trials have been performed using antibiotic therapy in UC and CD with contradictory results.

Antibiotics in UC

Only few trials with antibacterial agents have been carried out in IBD and the results are controversial.

Dickinson et al. [27], in 1985, published a double-blind controlled trial on the use of oral vancomycin as an adjunctive therapy in acute exacerbations of idiopathic colitis. No significant difference was found between the two treatment groups with only a trend in favor of a superior efficacy of vancomycin. It is important to underline that 7 of the 40 patients enrolled had colonic CD and that none of them had *C. difficile* infection that could explain the action of vancomycin. Subsequently, intravenous metronidazole, in addition to steroids, was effective similar to placebo in inducing remission [28].

In 1990, Burke et al. [29] published a double-blind, placebo-controlled trial on the use of oral tobramycin in acute UC. Eighty-four patients were randomized to receive steroid plus tobramycin or placebo. After 1 week of treatment, 74% of patients in the tobramycin treatment group versus 43% in the placebo group ($p < 0.003$) achieved a complete remission. Subsequently, tobramycin and metronidazole were associated with a standard steroid treatment in severely acute UC. At the end, no difference was found between the two groups [30].

Ciprofloxacin has been tested in a randomized, placebo-controlled study; 70 patients with mildly to moderately active UC were randomized to receive ciprofloxacin 250 mg b.i.d. or placebo for 14 days. At the end of the study, 70.5% of patients in the ciprofloxacin group versus 72% in the placebo group showed an improvement [31]. Nevertheless, in a more recent randomized, placebo-controlled trial, ciprofloxacin was administered for 6 months to patients with active UC poorly responding to conventional therapy with steroids and mesalazine. At the end of the study, the treatment failure rate was 21% in the ciprofloxacin-treated group and 44% in the placebo group ($p < 0.002$). Also endoscopic and histological evaluation showed a better improvement in the ciprofloxacin group [32].

Based on the experimental observation of a beneficial effect of intracolonic amoxicillin-clavulanic acid in a rat model of colitis, Casellas et al. [33] used an enteric-coated amoxicillin-clavulanic acid (1 g amoxicillin plus 250 mg clavulanic acid, t.i.d.) in active UC. They also evaluated the release of inflammatory mediators (IL-8, TXB₂, PGE₂) in rectal dialysates. After short-term treatment, this formulation decreased intraluminal release of IL-8 and other inflammatory mediators and led to an improvement of patients with active UC.

Antibiotics in CD

Metronidazole has been the mostly investigated agent. In 1978, Blichfeldt et al. [34] performed the first controlled study of metronidazole in CD. They did not find a difference between metronidazole and placebo-treated patients but a positive trend in favor of metronidazole was observed when the colon was involved. In the National Cooperative Swedish study, metronidazole was compared to sulfasalazine; no significant difference was found between the two groups; however, in the crossover section of the study, metronidazole was effective in patients not responding to sulfasalazine [35].

Metronidazole was used as single therapy or associated with cotrimoxazole compared to placebo in patients with a symptomatic relapse of CD. At the end of the 4 weeks of treatment there was no difference in response among the three groups [36]. In a Canadian study, two different dosages of metronidazole (10 and 20 mg/kg/daily) were used versus placebo in patients with active CD; no difference was found between the two groups, but in the metronidazole-treated group there was a high rate of dropout because of side effects or intolerance [37].

An antibiotic association was used in an Italian randomized controlled study in which metronidazole 250 mg

4 times daily plus ciprofloxacin 500 mg twice daily were compared to a standard steroid treatment. No differences were found and it was concluded that the antibiotic association could be an alternative to steroid treatment in acute phases of CD [38].

Ciprofloxacin has an excellent activity against enteric pathogens and Gram-negative Enterobacteriaceae associated with immunosuppressive properties. Ciprofloxacin 1 g/daily was compared to mesalazine 4 g/daily in a controlled study in mildly to moderately active CD. After 6 weeks an equivalence in efficacy was registered, offering an alternative treatment in active CD [39].

In a small study ciprofloxacin was shown to be effective in association with standard treatment in patients with resistant disease [40]. The results of this study were challenged by a controlled study in which ciprofloxacin (1 g/day) was associated with budesonide (9 mg/day) in ileocolic active CD. No difference was found compared to placebo, but surprisingly the overall response in both groups was lower than that reported in previous studies with budesonide [41].

A lot of studies have tried to evaluate the efficacy of antimycobacterial drugs in patients with CD, investigating the possibility that a strain of *Mycobacterium* might be an etiological agent in CD. Borgaonkar et al. [42] evaluated all randomized controlled trials in which antimycobacterial therapy was compared with placebo, suggesting an efficacy of antimycobacterial therapy only in few patients. However, the investigator emphasized that because of the small number of studies included in the meta-analysis, the data were not conclusive and a high rate of side effects was registered.

Metronidazole at the dose of 20 mg/kg/day was also tested by Rutgeerts et al. [43] in the prevention of postoperative recurrence. Sixty patients were randomized to

receive metronidazole or placebo for 12 weeks. At the end of the treatment, endoscopic relapse was evaluated by the Rutgeerts score. Metronidazole significantly reduced the incidence of severe endoscopic relapse (grade 3 or 4) but was complicated by a high incidence of side effects. Recently, ornidazole, another nitroimidazole derivative, was proposed in the prevention of postoperative recurrence in order to reduce the incidence of side effects. After 12 months, ornidazole was significantly more effective than placebo in the prevention of recurrence [44].

Rifaximin in the Treatment of IBD

Rifaximin is a rifamycin-derived antibiotic [45] with (1) a large antimicrobial spectrum covering most Gram-positive and Gram-negative bacteria, including aerobes and anaerobes and (2) poor absorption after oral administration and complete fecal excretion as unchanged drug.

Rifaximin has revealed as excellent safety profile: in clinical trials adverse reactions were observed in less than 2% of patients; most of the side effects were of gastrointestinal type or origin (such as nausea, vomiting, flatulence/meteorism, abdominal pain/cramp) while a mild to moderate urticarioid rash was infrequent [45].

A first open, uncontrolled study [46], performed in 12 patients with active IBD refractory to standard treatment who all had positive stool culture, suggested that adding rifaximin (800 mg daily) could be beneficial. A further small but controlled investigation performed in our unit [47] evaluated the efficacy and systemic absorption of rifaximin in patients with moderately to severely active UC refractory to steroid treatment. Patients were eligible if they had no response to intravenous corticosteroid therapy (methylprednisolone 1 mg/kg/day) after 7–10 days. Twenty-eight patients were randomized to receive rifaximin 400 mg b.i.d. or placebo for 10 days as an add-on

Table 1. Rifaximin vs. placebo in severe UC: outcome after 10 days of therapy (from Gionchetti et al. [47])

	Rifaximin			Placebo		
	before	after	p	before	after	p
Stool frequency	6.3±2.5	4.2±2.5	<0.02	6.3±2.8	5.2±2.5	NS
Rectal bleeding	1.41±0.51	1.0±0.0	<0.05	1.43±0.51	1.29±0.49	NS
Fever	36.6±0.3	36.5±0.18	0.07	36.7±0.4	36.9±1.0	NS
Sigmoidoscopic score	2.58±0.51	1.83±0.83	<0.01	2.14±0.36	1.57±0.85	0.06
Clinical activity	2.21±0.43	1.57±0.65	<0.05	2.42±0.51	1.83±0.83	<0.05

medication to standard steroid treatment. Clinical and endoscopic evaluation were performed before and after the treatment, and stool frequency, consistency and presence of blood were also recorded. Plasma and urine samples were collected before and after the treatment to determine the systemic absorption of rifaximin. Although there was no significant difference in clinical efficacy between the two treatments, only rifaximin determined a significant improvement of stool frequency, rectal bleeding and sigmoidoscopic score (table 1) [47]. The cumulative excretion of rifaximin in 24-hour urine after 10 days was 64,617 ng, confirming the poor systemic absorption also in presence of colonic inflammation [47]. The efficacy of this rifamycin derivative as add-on medication in patients with mild to moderate UC was recently confirmed in another open-label study [48], where the clinical activity index decreased by 30% after 4 weeks of treatment.

In patients, who experienced a clinical exacerbation of UC and who had a past history of serious adverse reactions to steroids, the antibiotic (400 mg b.i.d. for 4 weeks) was added to mesalazine (2.4 g daily) treatment [49]. In 7 out of 10 patients (i.e. 70%) a clinical remission was achieved without corticoid use, thus showing that rifaximin displays a steroid-sparing effect.

Rifaximin, at a very high dosage (2 g/day), was also used in association with ciprofloxacin 1 g/day in the therapy of chronic, treatment-resistant pouchitis [50]. This 15-day regimen induced a clinical, endoscopic and histological remission in 89% of treated patients with no side effects (fig. 1). Microbiological results have shown a significant decrease in the fecal concentration of most of the colonic bacterial species (table 2), suggesting the need for an antibiotic association in the case of refractory pouchitis to determine a wide antibacterial activity against either Gram-positive or Gram-negative bacteria, both anaerobes and aerobes. Pharmacokinetic data confirmed again the very limited excretion of unchanged rifaximin in the 24-hour urine samples (24.563 ng/ml, range 0–134.181) [50].

More recently, Campieri et al. [51] performed a randomized trial to evaluate the efficacy in the prevention of postoperative recurrence with rifaximin 1.8 g daily for 3 months followed by a probiotic preparation (VSL#3) 6 g daily for 9 months versus mesalazine 4 g daily for 12 months in 40 patients after curative resection for CD. After 3 months of treatment, patients on rifaximin had a significantly lower incidence of severe endoscopic recurrence compared to those on mesalazine [2/20 (10%) vs. 8/20 (40%)]. This difference was maintained since the end of the study using probiotics [4/20 (20%) vs. 8/20 (40%)].

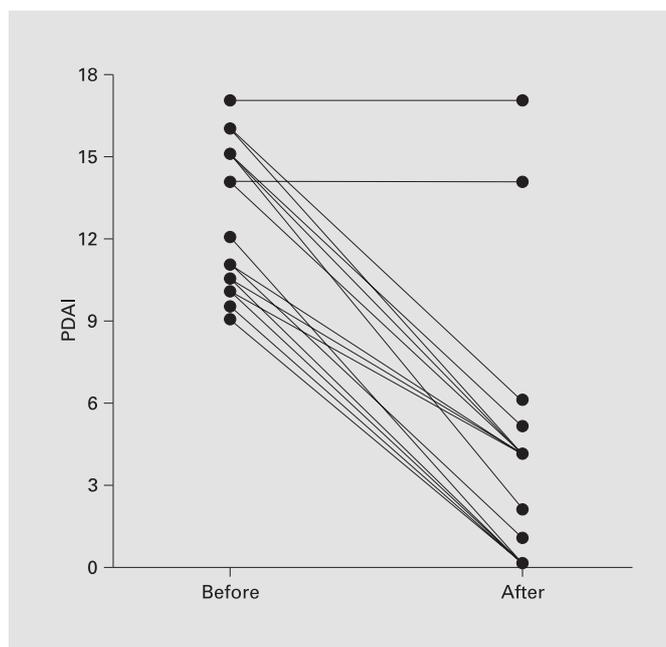


Fig. 1. Pouchitis Disease Activity Index (PDAI) score changes before and after antibiotic therapy in patients with chronic, treatment-resistant pouchitis (from Gionchetti et al. [50]).

Table 2. Bacterial counts in fecal samples of patients with active chronic pouchitis before and after combined antibiotic treatment (from Gionchetti et al. [50])

Bacterial species	Bacterial count (log ₁₀ CFU/g fecal dry weight)	
	basal values	values after antibiotic treatment
Total anaerobes	7.12 ± 2.03	5.17 ± 2.33**
Total aerobes	7.34 ± 1.49	5.37 ± 2.55**
Enterococci	6.14 ± 1.66	3.99 ± 2.39**
Coliforms	4.03 ± 2.56	3.57 ± 2.36
Bifidobacteria	3.82 ± 1.89	2.78 ± 1.38*
Lactobacilli	4.03 ± 2.03	2.89 ± 1.56**
<i>Clostridium perfringens</i>	3.67 ± 1.71	2.97 ± 1.69
Bacteroides	2.77 ± 1.70	2.17 ± 0.68*

Data are expressed as mean ± SEM. * p < 0.05; ** p < 0.01.

In conclusion, the results of this pilot study suggest the efficacy of the sequential combination of rifaximin and the highly concentrated probiotic preparation VSL#3 in the prevention of severe endoscopic recurrence of CD after surgical resection [51].

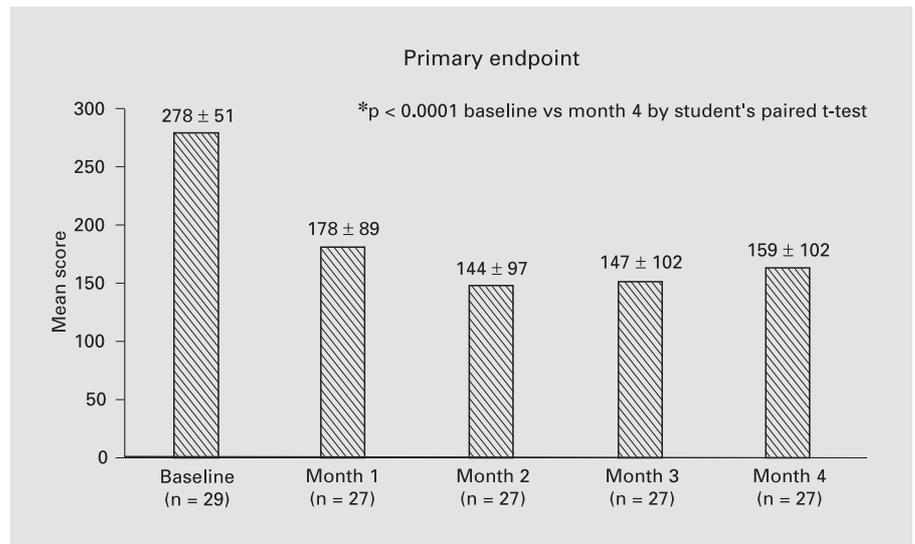


Fig. 2. Effect of rifaximin treatment on CDAI in patients with mildly to moderately active CD (from Shafran et al. [52]).

Finally, Shafran et al. [52] presented recently an open-label study on the efficacy and safety of rifaximin 600 mg/day for 16 weeks in the treatment of mildly to moderately active CD. At the end of the study, 59% of patients were in remission (as defined by a Crohn's Disease Activity Index, CDAI, <150) with a significant reduction of the

mean CDAI score compared to baseline ($p < 0.0001$) (fig. 2). Only one nonserious drug-related adverse event was reported, confirming the safety of the drug.

All these data, taken together suggest that this antibiotic is clinically useful in most cases of active intestinal inflammation.

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Treatment of Small Intestine Bacterial Overgrowth and Related Symptoms by Rifaximin

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Key Words

Rifaximin · Small intestine bacterial overgrowth · Therapy · Nonabsorbable antibiotics

Abstract

The treatment of small intestine bacterial overgrowth should address different aims: the removal of the predisposing condition, guarantee of adequate nutritional support to reintegrate both caloric and vitamin requirements and, obviously, suppression of the contaminating bacterial flora, which represents the major goal. The polymicrobial nature of contaminating flora suggests the administration of wide-spectrum antibiotics, but until now there has been no conclusive information on the most effective therapeutic approach. In this paper, the efficacy of the different therapeutic approaches used is reviewed.

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Introduction

The treatment of small intestine bacterial overgrowth (SIBO) is a clinical challenge for physicians, as data contained in the peer review literature do not offer conclusive

information on the most effective therapeutic approach. In this paper, available data were reviewed in order to suggest some indications.

Clinical and Pathophysiological Aspects

SIBO is due to the presence of more than 10^6 colony-forming units per milliliter of intestinal aspirate and/or colonic-type species [1]. Although asymptomatic cases, mostly among the elderly [2], have been described, the condition is generally accompanied by malabsorption and the consequent clinical syndrome is characterized by major symptoms, such as diarrhea, steatorrhea and weight loss, together with abdominal pain, bloating and flatulence. Therefore, an impairment of the nutritional status is frequently present and several nutritional defects have been described [3–9].

Malabsorption in SIBO is considered the consequence of abnormalities occurring mainly in the intraluminal environment; in fact, the excessive number of intraluminal bacteria interfere with the absorption process. However, in some cases, the presence of bacterial species capable of more aggressive adhesion to small bowel epithelium is probably the cause of direct damage to the absorptive surface, in particular in the blind loop syndrome [10, 11].

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Table 1. Aims of the treatment of SIBO

<i>Nutritional support</i>
Minerals
Vitamins
Caloric requirements
<i>Removal of predisposing conditions</i>
Surgery
Prokinetics (?)
<i>Suppression of contaminating flora</i>
Antibiotics
Probiotics (?)

Table 2. Conditions predisposing to SIBO

<i>Anatomical defects</i>
Blind loops
Strictures
Fistulae
Diverticula
Gastric resections
Ileocolonic resections
<i>Functional defects</i>
Impaired motility causing intestinal stasis
Ageing
Reduced gastric acid secretion
Reduced activity of intestinal immune system

Generally, it is not easy to determine how important the role of the predisposing condition and that of bacterial overgrowth are in the pathogenesis of malabsorption. In fact, conditions like gastrectomy, ileocolonic fistula and short bowel may be causes of malabsorption regardless of the concomitant presence of bacterial overgrowth. The pathophysiological role of bacteria lies in their ability to metabolize nutritional substances, such as carbohydrates, lipids and proteins, normally absorbed at the level of the small bowel. This concerns two different problems. The first is the nutritional defect, due to the lack of caloric substrates available for absorption. The second is a series of effects due to the products of bacterial metabolism, such as increased carbohydrate fermentation [12], which is responsible for the presence of symptoms like flatulence, bloating, abdominal colicky pain and abdominal distention; increased levels of short-chain fatty acids,

which trigger the irritation of the colonic wall, acidification of both the bowel lumen and feces and, finally, diarrhea accompanied by abdominal pain, and reduced mucosal disaccharidase activity, due to its inactivation by proteases secreted by anaerobic bacteria [13].

Aims of Treatment

The aims of the treatment of SIBO are listed in table 1. First, physicians should take into consideration the possibility of eliminating the predisposing condition. The importance of the role of these conditions (table 2) was shown by the demonstration of the presence of SIBO in around three-quarters of patients with malabsorption symptoms associated with a predisposing condition [14]. However, elimination of such conditions is not always possible. In patients who have undergone surgical reconstruction with loss of the gastric acid barrier or ileocecal valve, complete recovery from this syndrome is never possible and bacterial overgrowth will always represent a clinical problem to be taken into consideration. On the other hand, in patients with stenosing or fistulizing Crohn's disease, the timing of surgery is subject to complex evaluation and, therefore, relapsing symptoms of SIBO syndrome often have to be dealt with.

Moreover, adequate nutritional support is mandatory. The aim of this therapeutic measure should be the reintegration of both caloric and vitamin requirements, often defective in these patients. The nutritional defect is caused both by the predisposing condition and by the malabsorption syndrome.

Finally, suppression of the contaminating bacterial flora represents the major aim of treatment for SIBO.

Therapeutic Approaches

Prokinetics

In some patients SIBO is caused by intestinal stasis secondary to motility defects, and the restoration of normal intestinal motility may represent an effective approach. Prokinetic agents have been shown to improve intestinal motility [15, 16], and the use of this therapeutic approach has been shown to be effective: in patients with scleroderma, low-dose octreotide proved to be useful in the reduction of bacterial overgrowth [17]. Unfortunately, cisapride was recently removed from the market due to cardiotoxicity, and, apart from an erythromycin analog without antibiotic effects which showed no effects in rats [18],

Table 3. Antibiotic regimens used in SIBO

Drug	Dose	n	Predisposing conditions	Responders
<i>Tetracycline</i>				
Kahn et al., 1966 [21]	250 mg q.i.d.	4	scleroderma	75%
Goldstein et al., 1961 [22]	250 mg q.i.d.	1	Billroth II	+
Gorbach and Tabaqchali, 1969 [28]	250 mg q.i.d.	1	ileocolonic anastomosis in Crohn's disease	-
Bjorneklett et al., 1983 [23]	NA	3	small bowel diverticulosis	100%
Di Stefano et al., 2000 [42]	333 mg t.i.d.	11	GI surgery, intestinal stasis	27%
<i>Chloramphenicol</i>				
Goldstein et al., 1961 [22]	500 mg q.i.d.	1	Billroth II	+
<i>Lincomycin</i>				
Bjorneklett et al., 1983 [23]	NA	1	radiation fibrosis	-
Gorbach and Tabaqchali, 1969 [28]	500 mg t.i.d.	2	small bowel diverticulosis	50%
<i>Ampicillin</i>				
Goldstein et al., 1970 [26]	250 mg q.i.d.	1	diabetic autonomic neuropathy	+
<i>Metronidazole</i>				
Bjorneklett et al., 1983 [23]	NA	6	radiation fibrosis, small bowel diverticulosis	83%
<i>Cotrimoxazole</i>				
Elsborg, 1977 [30]	400 mg b.i.d.	1	small bowel diverticulosis	+
<i>Norfloxacin</i>				
Attar et al., 1999 [45]	400 mg b.i.d.	10	GI surgery or intestinal stasis	30%
<i>Amoxicillin-clavulanic acid</i>				
Attar et al., 1999 [45]	500 mg t.i.d.	10	GI surgery or intestinal stasis	50%
<i>Rifaximin</i>				
Trespi and Ferrieri, 1999 [43]	400 mg t.i.d.	8	chronic pancreatitis and Billroth II	100%
Di Stefano et al., 2000 [42]	400 mg t.i.d.	10	GI surgery or intestinal stasis	70%

NA = Not applicable; GI = gastrointestinal; + = positive effect of therapy; - = no effect of therapy.

no other drugs have been tested. Consequently, the role of these agents in the treatment of SIBO still needs to be explored.

Antibiotics

The polymicrobial nature of contaminating flora makes the use of wide-spectrum antibiotics mandatory [19, 20]. The choice of the drug is frequently empirical because small bowel aspiration and culture are impractical and, if performed, will show with certainty multiple organisms with different antibiotic sensitivity. However, the reason why we need to use wide-spectrum antibiotics lies on a still lacking information: we do not know which organisms should be eliminated to achieve the improvement of symptoms [19]. Several antibiotics have been shown to be effective (table 3). However, as can be clearly seen from this table, available data are based very frequently on the

description of single cases. Although anaerobes are responsible for the main metabolic alterations, tetracyclines [21–23] have been used for a long time and with satisfactory results, in spite of their poor activity against these bacteria [3, 20, 24]. A rapid improvement of symptoms was evident in most cases after a single therapeutic course of 10–14 days at a dose of 250 mg four times a day [3]. Aerobe suppression induced by tetracyclines probably render the intraluminal microclimate unfavorable to anaerobes, due to the increased bioavailability of oxygen [19]. However, it was recently reported that about 60% of patients do not respond to this treatment [25]. Metronidazole [3, 23], ampicillin [26] and erythromycin [27] have been used as an alternative to tetracycline, while other drugs active against anaerobes, such as lincomycin [23, 28, 29] and chloramphenicol [3, 22], are no longer employed due to the high risk of severe side effects. Neomy-

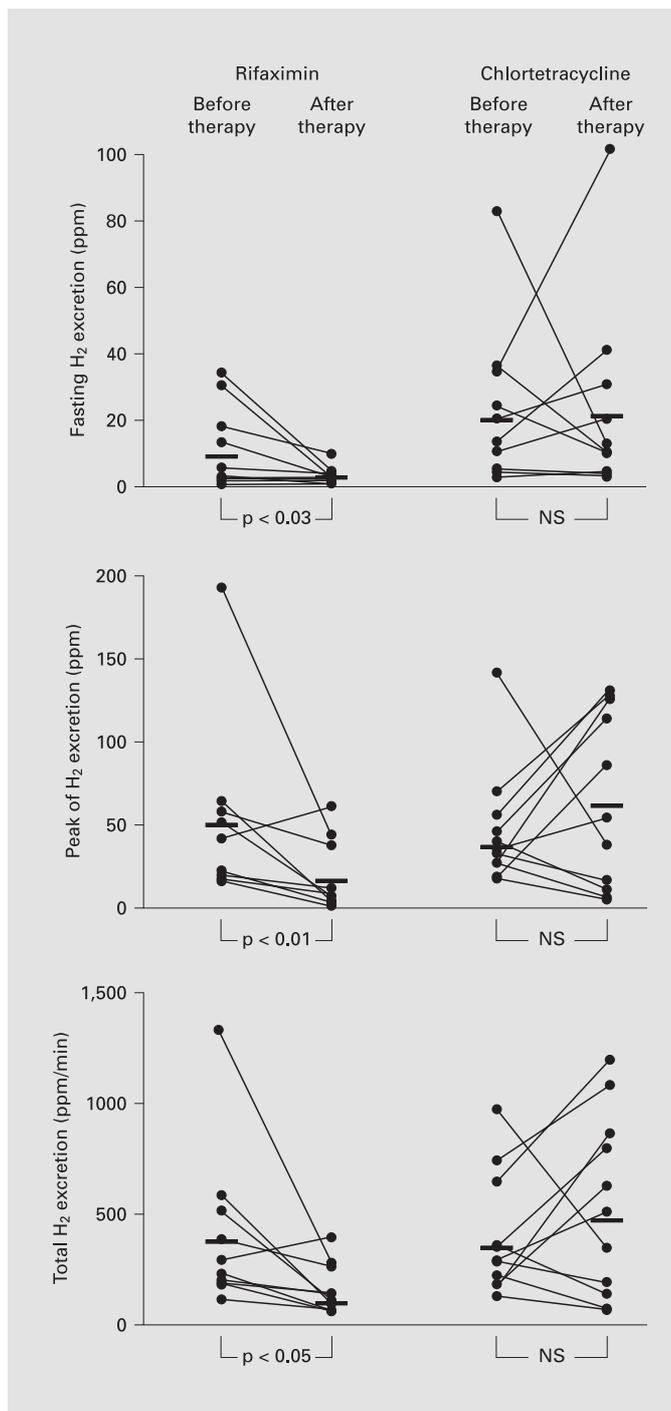


Fig. 1. Fasting, peak and total breath H₂ excretion before and after rifaximin or chlortetracycline therapy in two groups of patients with SIBO (from Di Stefano et al. [42]). NS = Not significant.

cin was shown to be of little efficacy when used alone in this condition [29]. Scanty information is available on cotrimoxazole, which was shown to be effective at a low dosage in a case report [30].

If contaminating flora is sensitive to the administered antibiotic, in most patients, a single course of 7–10 days of therapy is able to induce an improvement of symptoms [31]; however, a quick relapse of symptoms is often evident, but it can be treated with the same antibiotics [3, 27]. In these cases, good results can be achieved with intermittent antibiotic treatment [3].

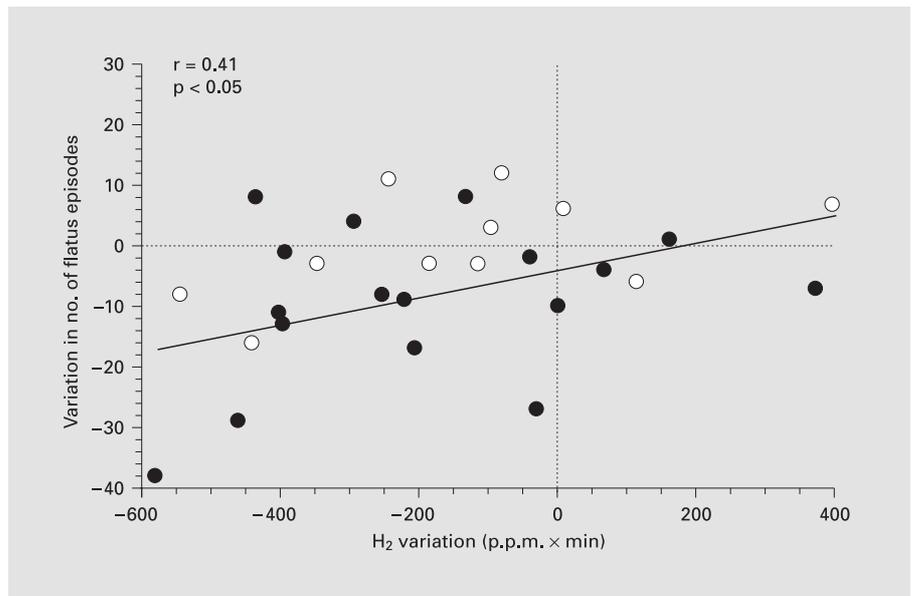
In a recent study, the bacterial populations contaminating the upper gut in SIBO patients and their antibiotic susceptibility were determined. Amoxicillin-clavulanic acid and cefoxitin were effective against >90% of anaerobic strains, while aminopenicillins, cephalosporins and cotrimoxazole were effective against the microaerophilic population. Erythromycin, clindamycin and rifampicin were ineffective. Data on metronidazole and fluoroquinolones are not available [32].

Rifaximin, a nonabsorbable derivative of rifamycin, has shown promising bactericidal action against both aerobes and anaerobes, such as bacterioides, lactobacilli and clostridia [33, 34]. The development of resistance to this antibiotic can occur, but resistant strains rapidly disappear from the intestine thus allowing cyclic administration of rifaximin. Controlled clinical trials showed efficacy of rifaximin in adult and pediatric patients with infectious diarrhea [36, 37], hepatic encephalopathy [38], post-surgical complications [39] and colonic diverticulosis [40]. Only recently was the efficacy of rifaximin in the treatment of SIBO demonstrated [41–43].

The most important evidence is probably offered by a recent double-blind, randomized trial which showed a better therapeutic effect of rifaximin in comparison to tetracycline in a cohort of SIBO syndrome patients [42]; in particular, rifaximin administration produced a significant reduction of breath hydrogen levels in fasting conditions, peak of hydrogen excretion and cumulative breath hydrogen excretion after an oral dose of 50 g of glucose (fig. 1). Normalization of the test results was evident in 70% of the sample studied.

A significant improvement of symptom severity and the absence of side effects was also evident after rifaximin administration but not after tetracycline, reinforcing, therefore, the validity of the therapeutic approach adopted. Rifaximin has proved to be effective in the treatment of gas-related symptoms; in fact, in a recent paper, it was reported that a 7-day course of therapy significantly improved the severity of symptoms in a cohort of patients

Fig. 2. Correlation between variation in breath H₂ excretion and variation in the number of flatus episodes recorded before and after therapy. Black circles indicate rifaximin-treated patients, white circles indicate charcoal-treated patients. Patients with score >0 were considered (from Di Stefano et al. [44]).



suffering from functional abdominal complaints (pain, bloating, flatulence) [44]. This effect should also be important in patients with SIBO syndrome; it would probably become more evident if courses of therapy longer than 1 week are prescribed, in view of the interference with the therapeutic efficacy due to the presence of predisposing conditions. A correlation between variation in breath H₂ excretion and variation in the number of flatus episodes was actually found in the above-mentioned study (fig. 2).

Another recent controlled trial showed a good therapeutic effect of both amoxicillin-clavulanic acid and norfloxacin in SIBO patients [45]. However, a rapid relapse of diarrhea just few days after the withdrawal of antibiotics was evident. In this paper, the efficacy of probiotics in SIBO patients was also evaluated, but no significant effect was described. While on the one hand these results confirm the frequent need of several courses of antibiotic therapy in SIBO patients, on the other they support the idea that rifaximin may represent a good choice on the basis of its excellent tolerability.

In our opinion, the pivotal topic of the treatment of SIBO syndrome is probably represented by the careful evaluation of clinical polymorphism in these patients; the presence of several predisposing conditions, very different with respect to both pathophysiological and clinical aspects, may modify the clinical response of individual patients, thus affecting the overall efficacy of the therapeutic programs. Preliminary results, in fact, have shown that patients with SIBO and blind loop syndrome display

a better clinical response after metronidazole therapy than after rifaximin, likely due to a lower drug concentration at the level of the blind loop [46]. It is, therefore, possible that in future the study of subgroups of patients will give us more informations and will clarify several still nonstandardized issues such as the optimal dosage and duration of therapy. Moreover, the availability of nonabsorbable antibiotics in clinical practice will represent an improvement of the current therapeutic strategies.

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Note Added in Proof

While this review was being typeset, some papers of interest were published. The efficacy of rifaximin in relieving functional symptoms such as bloating and flatulence was recently confirmed in a randomized, double-blind, placebo controlled trial [1]. In a cohort of 145 patients with Crohn's disease the presence of SIBO was found by hydrogen breath test after glucose administration in 20% of patients [2]. Both metronidazole and ciprofloxacin proved to be effective in the management of bacterial overgrowth and breath hydrogen excretion normalized in 86% and 100% of patients, respectively. In metronidazole group, 1 patient out of 15 withdrew after two days because of nausea and, together with the other 2 patients resistant to metronidazole, was successfully treated with ciprofloxacin. After a 1-year period of follow-up, only 1 patient presented a recurrence of bacterial overgrowth. This study suggests, therefore, that ciprofloxacin is more effective than norfloxacin in the treatment of SIBO and confirms that side

effects represent a major problem for the therapy with metronidazole. Moreover, in 50 consecutive patients with various malabsorption syndromes, 42% of jejunal aspirates showed a bacterial count greater than 10^5 CFU/ml. *Streptococcus* species and *Escherichia coli* were the commonest Gram positive and negative isolated bacteria, respectively, and proved to be more sensitive to quinolones than to tetracycline, ampicillin, erythromycin and cotrimoxazole [3]. Unfortunately, no data are available on rifaximin. Finally, the effect of probiotics on SIBO-related diarrhea was investigated in another paper. Two different *Lactobacillus* strains, namely *L. casei* and *L. acidophilus* strains Cerela, were administered and results were compared to placebo. A short-term improvement of the number of bowel movements and breath hydrogen excretion was achieved by probiotic treatment, suggesting that probiotics may represent an important therapeutic option in the treatment of SIBO provided prolonged courses are adopted [4].

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Management of Diverticular Disease: Is There Room for Rifaximin?

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Key Words

Diverticular disease · Diverticulitis · Antibiotics · Rifaximin

Abstract

Treatment of symptomatic diverticular disease of the colon is aimed at the relief of symptoms and the prevention of major complications. The efficacy of fiber supplementation and of anticholinergic and spasmolytic agents remains controversial. Antibiotics are commonly used in the treatment of inflammatory complications of diverticular disease. Data from open labelled and randomized controlled trials do suggest the efficacy of rifaximin in obtaining symptomatic relief in patients with diverticular disease. Approximately 30% therapeutic gain compared to fiber supplementation only can be expected after one year of intermittent treatment with rifaximin. Considering the safety and tolerability of rifaximin, this drug can be recommended for patients with symptomatic uncomplicated diverticular disease.

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Introduction

Diverticular disease of the colon is extremely common in developed countries and its prevalence is correlated with advancing age [1]. Estimates based on necroscopy or

radiologic findings indicate that it occurs in about 10% of people in the United States, the United Kingdom and Australia [2] and that it is currently found in one third to one half of all autopsies of people over 60 years of age [3]. In North America, diverticular disease is estimated to occur in one third of all individuals above 45 years of age and two thirds of individuals above 85 years of age [4].

The different prevalence rates in various geographic areas and ethnic groups have been confirmed by several reports; the high prevalence of diverticular disease in developed western countries is in contrast with the rarity with which the disease is observed in less industrialized countries and in Japan [4–6]. This led to the hypothesis that the high frequency of diverticular disease in western societies results from a low fiber consumption by an aging population [3, 4]. The dietary fiber hypothesis is largely supported by epidemiological observations [4, 5, 7–9] and case-control studies [10–12]. Recently, a prospective cohort study of 43,881 male health professionals 40–75 years of age demonstrated that dietary fiber consumption was inversely associated with the risk of developing diverticular disease and this association was particularly strong for cellulose [13]. In more recent years, with increasing globalization, factors previously uncommon in developing countries, such as the availability of highly processed foods, may be operating to cause a similar prevalence of diverticular disease in these populations as in European and American populations [14].

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The majority of patients harboring colonic diverticula remain asymptomatic throughout their life (asymptomatic diverticular disease); only 20% will develop symptoms and signs of illness [15]. Symptomatic diverticular disease is further subdivided into painful diverticular disease (symptomatic diverticular disease with no inflammation) and diverticulitis (symptomatic diverticular disease with inflammation). Diverticulitis is further subdivided into uncomplicated and complicated diverticulitis [16].

Treatment of Diverticular Disease of the Colon

The vast majority of patients with colonic diverticula will remain entirely asymptomatic. There are no data to support any therapeutic recommendations or routine follow-up in this large population, although it is reasonable to recommend a diet high in fruit and vegetable fiber.

Treatment of symptomatic diverticular disease is aimed at the relief of symptoms and the prevention of major complications. The efficacy of fiber supplementation in the treatment of painful diverticular disease remains controversial. Some controlled clinical trials have shown a benefit of high-fiber diets for symptomatic relief [17–20], but other studies failed to show positive results [21–23]. Despite these conflicting data, bran and bulking agents are commonly used in the treatment of symptomatic diverticular disease. The high intraluminal colonic pressure is of critical importance in the formation of diverticula, and this formed the basis for the high-fiber diet recommended to individuals with symptomatic diverticular disease [5]. Recent American College of Gastroenterology Guidelines [24] recommended a high-fiber diet for patients with symptomatic diverticular disease of the colon; some amelioration of symptoms can be expected, and other potential health benefits of fiber should be considered.

The observed hypermotility of the sigmoid colon in many symptomatic patients provides the rationale for using anticholinergic drugs and spasmolytic agents [25]. However, the efficacy of these agents has never been clearly documented in randomized controlled trials [3, 24].

Antibiotics are commonly used in the treatment of inflammatory complications of diverticular disease [3, 24]. In painful diverticular disease, when inflammation is excluded by definition, there is no rationale for using antibiotics. In a subset of patients with more severe symptoms, when an inflammatory component may be clinically suspected, a short course of antibiotics may be advisable [15].

However, some observations suggest a possible role of gut microflora in determining symptoms related to diverticular disease. Bacterial metabolism is the major source of intestinal gas such as H₂, CO₂ and CH₄ via carbohydrate fermentation [26]. Excessive production of bowel gas can play a role in determining abdominal symptoms such as bloating, pain and discomfort [27]; these symptoms, although nonspecific, are commonly observed in patients with diverticular disease of the colon. Antimicrobial drugs have been shown to reduce colonic H₂ production [28, 29] and gas-related symptoms [30]. Interactions between dietary fiber, bacterial metabolism and colonic functions are complex and not fully elucidated. It is well known that fiber increases stool bulk and affects colonic transit time in several ways: by water holding, by proliferation of bacteria and by the products of bacterial fermentation [31]. It has been assumed since the earliest times that fiber exerts its effect on bowel habits by virtue of retaining water in the gut and stimulating peristalsis through increased bulk [32]. However, water holding is an *in vitro* property of fiber which is inversely related to fecal bulking and colonic transit time *in vivo* [33]. This is not surprising, because virtually all fiber is broken down in the gut by intestinal bacteria [32]. Moreover, water holding can be equated with the solubility of fiber, and soluble fiber is more rapidly degraded by gut microflora [32]. This view is supported by a study in humans in which it was shown that antimicrobial therapy causes a rise in mean fecal weight in subjects with a constant fiber intake, probably due to the reduction of bacterial mass and bacterial degradation of fiber [34]. Therefore, a role of antibiotics in diverticular disease symptoms may be suggested, with respect to both bacterial gas production and fiber degradation.

Rifaximin in Diverticular Disease of the Colon

Rifaximin is a semisynthetic rifamycin derivative that acts by inhibiting bacterial ribonucleic acid synthesis [35]. *In vitro* data indicate a good antibacterial activity against Gram-positive organisms and less activity against Gram-negative organisms; conflicting data exist for activity against bacteroides [35]. Rifaximin is virtually unabsorbed after oral administration; thus, it is used primarily to treat local conditions within the gastrointestinal tract [35]. Rifaximin has been successfully used in the treatment of infectious diarrheas [36, 37], including travelers' diarrhea [38], and it has been shown to be at least as effective as neomycin in the treatment of portosystemic en-

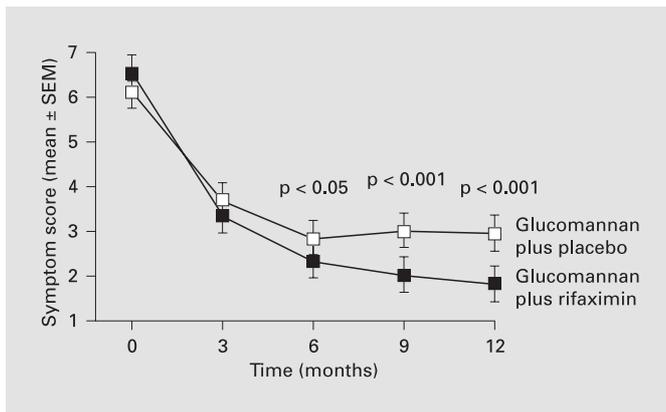


Fig. 1. Global symptom score in patients treated with glucomannan plus rifaximin and in patients treated with glucomannan plus placebo (from Papi et al. [45]).

cephalopathy [39, 40]. Furthermore, rifaximin has been shown to be effective in the treatment of intestinal bacterial overgrowth [41] and in the prophylaxis of postoperative complications following colorectal surgery [42].

Several clinical observations suggest a role of rifaximin in the management of symptomatic uncomplicated diverticular disease of the colon.

In a pilot multicenter open trial [43], 217 patients with symptomatic uncomplicated diverticular disease were treated with glucomannan 2 g/day or with glucomannan plus rifaximin 400 mg twice a day for 7 days each month. Clinical evaluation was performed at admission and at 2-month intervals for 12 months; a symptom score including 8 clinical variables was used as a measure of the therapeutic effect. After 12 months of treatment, a significant reduction of the symptom score was observed in patients treated with glucomannan plus rifaximin compared to patients treated with glucomannan only (63.9 vs. 47.6%; $p < 0.001$). At the end of the study period, 58% of patients treated with rifaximin and glucomannan were symptom free compared to 24% of patients treated with glucomannan only ($p < 0.001$).

Similar results were obtained in a large prospective open trial including 968 outpatients with symptomatic diverticular disease [44]. Patients were randomly assigned to receive fiber supplementation (glucomannan 4 g/day) or fiber supplementation plus rifaximin (400 mg twice a day for 7 days every month) for 12 months. After 12 months, 56.5% of patients in the rifaximin group were symptom free compared to 29.2% of patients in the fiber supplementation only group ($p < 0.001$).

In a multicenter double-blind placebo-controlled trial [45], 168 outpatients with symptomatic uncomplicated diverticular disease were randomly assigned to receive fiber supplementation (glucomannan 2 g/day) plus rifaximin (400 mg twice a day for 7 days every month for 12 months), or fiber supplementation plus placebo. A symptom score including 6 clinical indications (upper abdominal pain/discomfort, bloating, lower abdominal pain/discomfort, tenesmus, diarrhea and abdominal tenderness) was calculated to assess the clinical response. Both treatments were shown to be effective in reducing the score after the first 3 months of treatment; however, patients treated with rifaximin showed a significantly greater reduction in the score compared to patients treated with placebo at 6, 9 and 12 months (fig. 1). Symptoms such as abdominal pain, abdominal tenderness and bloating appeared to be particularly affected by rifaximin treatment. After 12 months of treatment, 68.9% of patients in the rifaximin group were symptom free compared to 39.5% in the placebo group [absolute benefit increase 29.7%, 95% confidence interval (CI) 15.3–44.1%]. Long-term intermittent rifaximin administration appears to be safe; no relevant side effects were reported in any of these trials [42–44].

Open-labeled and randomized controlled trial data support some evidence for the efficacy of intermittent long-term administration of rifaximin in patients with symptomatic diverticular disease. A therapeutic gain of approximately 30% compared to fiber supplementation only can be expected with respect to obtaining symptomatic relief after 1 year of treatment (table 1). Considering the safety and tolerability of rifaximin, this drug can be recommended for patients with symptomatic uncomplicated diverticular disease.

Although there is some evidence for the efficacy of long-term treatment with rifaximin for symptomatic relief in patients with uncomplicated diverticular disease, an unresolved issue is whether rifaximin can prevent major inflammatory complications of diverticular disease. In the two prospective open trials discussed above, the occurrence rate of complications in 12 months was lower in patients treated with glucomannan plus rifaximin compared to patients treated with glucomannan only: 2.7 versus 0.9% [43] and 3.2 versus 1.3% [44]. This observation was not confirmed in the double-blind placebo-controlled trial [45] in which no difference in the 1-year complication rate was observed between the rifaximin and placebo groups. However, in all the studies, the number of patients suffering complications in a 12-month period was too small to detect any statistically significant

Table 1. Studies addressing rifaximin for the treatment of symptomatic diverticular disease

Author	Patients	Study design	Treatment	Study period months	Asymptomatic patients, %	RD, % (95% CI)	Complications, %	RD, % (95% CI)
Papi et al. 1992 [43]	217	open	glucomannan 2 g + glucomannan 2 g + rifaximin ¹	12	24	34.3 (22.0–46.5)	2.7	–1.8 (–5.3 to 1.7)
					58		0.9	
Latella et al. 2003 [44]	968	open	glucomannan 4 g + glucomannan 4 g + rifaximin ¹	12	29	27.0 (20.9–33.1)	3.2	–1.8 (–3.8 to 0.1)
					56		1.3	
Papi et al. 1995 [45]	168	RCT	glucomannan 2 g + placebo glucomannan 2 g + rifaximin ¹	12	39	29.7 (15.3–44.1)	2.3	0 (–4.6 to 4.6)
					69		2.3	

RD = Rate difference; RCT = randomized controlled trials.

¹ Rifaximin 400 mg twice a day for 7 days each month for 12 months.

difference between the two treatment groups (table 1). In a retrospective study of 505 patients admitted to hospital for a major complication of diverticular disease of the colon (occlusion, perforation, fistula or bleeding) and discharged after conservative management, long-term treatment with poorly absorbable antibiotics (neomycin plus bacitracin, paromomycin or rifaximin) reduced by 50% the relative risk of readmission for complications compared to no antibiotic treatment [46].

Conclusions

There is some evidence that long-term cyclic administration of rifaximin combined with fiber supplementation is effective for inducing symptomatic relief in patients with uncomplicated diverticular disease of the colon. A therapeutic gain of approximately 30% compared to fiber supplementation only can be expected. The drug is well tolerated and no relevant side effects have been reported. Symptoms attributed to diverticula (abdominal pain or discomfort, bloating, disturbance of bowel habits) are nonspecific symptoms and are also features of irritable bowel syndrome. It has been suggested that irritable bowel syndrome and diverticular disease of the colon may coexist in many people and when bowel symptoms occur with diverticulosis coli, they may be due to a coexistent irritable bowel rather to the diverticula themselves [47, 48]. This hypothesis is supported by the fact that many patients with symptomatic diverticular disease show co-

lonic motility patterns similar to those with irritable bowel syndrome [25], and patients with asymptomatic disease have colonic myoelectric activity similar to that of normals [49]. A fundamental issue is that treatment with rifaximin may be related to colonic symptoms rather than to diverticula per se; this would suggest the inclusion, in future studies, of an additional arm of patients with symptoms of irritable bowel syndrome but not diverticula.

No definitive conclusions can be drawn concerning a possible role of rifaximin in preventing major complications of diverticular disease. Double-blind placebo-controlled trials with an adequate sample size are needed. However, such trials are difficult to perform considering the requirement of a large number of patients. Assuming a baseline risk of complications of diverticular disease of 5% per year [2], a randomized controlled trial able to detect a 50% risk reduction in complications should include 1,600 patients per treatment group considering a power of 80% ($1 - \beta$) and an α error of 5%.

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Mechanical and Antibacterial Bowel Preparation in Colon and Rectal Surgery

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Key Words

Preoperative colon preparation · Intestinal antisepsis ·
Oral antibiotics · Erythromycin-neomycin

Abstract

Colorectal surgery performed prior to 1970 was fraught with postoperative infectious complications which occurred in more than 30–50% of all operations. Diversion of the fecal stream appeared mandatory when operating on an urgent or emergent basis, thereby requiring the performance of multiple, staged operations instead of a single surgery encompassing resection and primary anastomosis as is performed commonly today. Multiple studies conducted in the early 1970s determined that anaerobic colonic microflora were causative agents in postoperative infections in colon and rectal surgery, and these studies initiated the development of effective oral preoperative antibiotic prophylaxis in combination with preoperative mechanical bowel preparation. This dual-tier regimen significantly reduced the incidence of postoperative infectious complications, thus allowing most uncomplicated colon and rectal surgeries to be performed in a single stage without the need for the diversion of the fecal stream and multiple operations. Therefore, a preoperative mechanical and antibacterial bowel regimen serves as the cornerstone of modern elective colorectal surgery, and these regimens now comprise three therapeutic directives. The first step is preoperative mechanical cleansing of the bowel, which is then fol-

lowed by preoperative oral antibiotic prophylaxis. Finally, perioperative parenteral antibiotics directed against aerobic and anaerobic colonic microflora are utilized.

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Introduction

The advent of this new century marked the 30-year anniversary of a revolution in colon and rectal surgery – preoperative bowel sterilization, resulting in a significant decrease in infectious complications and the commonplace practice of single-stage operations. The past three decades have produced a plethora of clinical trials that clearly show that preoperative bowel preparation is an advantageous therapeutic endeavor in a surgeon's armamentarium. For decades, medical science has recognized and understood the important symbiotic relationship of the enormous reservoir of aerobic and anaerobic bacteria that inhabit the distal ileum and colon from infancy [1]. Furthermore, the intestinal epithelium and mucous membrane, which serve as a vital shield against these microbes, have also been recognized as a vital component of the human body's defense against these microbes [2]. Should this defensive mechanism break down secondary to disease or traumatic processes – iatrogenic or otherwise – local or systemic spread of these microorganisms could result in a significant clinical infection. Accordingly, therapeutic directives targeted at diminishing this bacterial load for elective surgical intervention have been at-

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Table 1. Alternate approaches to bowel preparation in patients undergoing elective resection of the large bowel

Environment	Approach 1 home or outpatient	Approach 2 home or outpatient
Diet	clear liquids beginning at 6.00 a.m. the day prior to surgery	clear liquids beginning at 5.00 p.m. 2 days prior to surgery
Preoperative mechanical bowel preparation	polyethylene glycol-electrolyte solution lavage – 4 liters p.o. over 3–4 h starting at 6.00 a.m. on the day prior to surgery	sodium phosphate ¹ prep of 45 ml p.o. at 6.00 p.m. 2 days prior to surgery and 6.00 a.m. the day prior to surgery
Enemas	none	none
Preoperative oral antibiotic bowel preparation	1 g of erythromycin base and 1 g of neomycin SO ₄ p.o. at 1.00, 2.00 and 11.00 p.m. on the day prior to surgery (surgery at 8.00 a.m.)	1 g of erythromycin base and 1 g of neomycin SO ₄ p.o. at 1.00, 2.00 and 11.00 p.m. the day prior to surgery (surgery at 8.00 a.m.)
Perioperative parenteral antibiotics	yes: with appropriate aerobic and anaerobic antimicrobial coverage with a single dose within 30 min of incision	yes: with appropriate aerobic and anaerobic antimicrobial coverage with a single dose within 30 min of incision

¹ Sodium phosphate is contraindicated as an oral mechanical bowel preparation in patients with renal or cardiac disease [8].

tempted for over 100 years [3]. These efforts have culminated in a standard three-tier regimen practiced widely in the United States [4], which includes (1) preoperative mechanical cleansing to reduce the fecal load and facilitate the efficacy of the orally administered antibiotics, (2) preoperative oral antimicrobial bactericidal therapy targeting both aerobic and anaerobic species, and (3) perioperative parenteral antimicrobial therapy. This therapeutic attack has resulted in a dramatic fall in the incidence of infectious postoperative complications in colon and rectal surgery.

Mechanical Preparation

Mechanical cleansing of the colonic lumen before elective colon resection is a time-tested procedure that, when done appropriately, reduces total fecal mass, thus facilitating operative manipulation of the colon and enhancing the action of oral antibiotics. Vigorous mechanical cleansing alone, however, whether it includes lavage or follows the classic approach (dietary restriction, enemas and cathartics), does not significantly reduce the number of microorganisms in the residual colonic material [5] or mucosal-associated bacteria [6, 7].

Today, approaches to mechanical cleansing vary considerably. The historically touted 5-day preoperative preparation has been abandoned due to the risk of inducing severe metabolic abnormalities. Modern approaches fall into two general categories: (1) whole-gut lavage with

polyethylene glycol-electrolyte solution or sodium phosphate on the day before operation, and (2) dietary restriction and cathartics for a 2-day period [4]. These two procedures may also be combined (table 1). Though these techniques are considered safer, complications arising from their use have been reported [8].

Three separate reports in the literature some 12 years apart document the North American perspective of 100% compliance with preoperative mechanical bowel cleansing [4, 9, 10]. The first study is a 1990 survey of 372 clinically active, board-certified colon and rectal surgeons in the United States and Canada [10]. The authors reported that every surgeon surveyed employed some form of mechanical cleansing in addition to antibiotics in preparing for elective colon resection. Fifty-eight percent of those surgeons who responded to the survey used lavage with polyethylene glycol-electrolyte solution, while 36% of surgeons used a more traditional approach that included dietary restriction, cathartics and enemas. The remaining 6% of surgeons used either preoperative lavage with mannitol or whole-gut lavage. More recently, 808 board-certified North American colorectal surgeons were surveyed for their current bowel preparation practices before elective procedures [4]. Of the 471 surgeons who responded to the survey (58%), all used some form of mechanical preparation. Oral polyethylene glycol-electrolyte solution was the most frequently reported mechanical preparation employed (70.9% of the respondents), while oral sodium phosphate solution with or without bisacodyl was the next most prevalent mechanical bowel prepara-

tion method used (28.4%). The 'traditional' method of dietary restriction, cathartics and enemas was the regimen employed least often (28.4%). Finally, the extensive literature review of Zmora et al. [9] confirmed the continued need for preoperative mechanical bowel preparations, even in light of recent reports to the contrary [11, 12]. However, these commonplace North American practice guidelines are not uniform to all surgeons worldwide.

Multiple studies have been performed that investigated the necessity of performing mechanical cleansing before elective colon resection [11–17]. Two found no differences in the rates of anastomotic dehiscence or infectious complications between patients who received mechanical cleansing and those who did not [13, 14]. These investigators relied on intravenous antibiotics for preoperative intestinal antisepsis and did not employ oral regimens. Another similarly designed Brazilian study showed a higher wound infection rate in the group of patients receiving mechanical cleansing [15]. The authors suggested that 'mechanical bowel preparation is unnecessary and may be harmful in terms of preventing wound infection and anastomotic dehiscence in patients undergoing elective colorectal surgery' [15]. It must be stressed that these three studies did not employ oral antibiotics in their bowel preparation regimens. Furthermore, four recent prospective studies [11, 12, 16, 17] documented that mechanical bowel preparations are not needed, on the basis of low anastomotic leakage rates (0–5%), low postoperative infection rates (5–6%) and equivalent total hospital stays.

Antibiotic Preparation

All surgeons employ antibiotics in preparing patients for an elective colon resection. As noted earlier, it is agreed that the antibiotics used should be bactericidal to both aerobic and anaerobic species, but what constitutes the ideal antibiotic regimen is a topic of considerable debate. Advocates of the oral administration of antibiotics typically emphasize the importance of reducing the number of microorganisms in the colonic lumen and mucosa before opening the colon, whereas advocates of parenteral administration emphasize the importance of adequate tissue levels of antibiotics prior to the beginning of surgery.

Oral Antibiotics

Over 30 years ago, the three major requirements for an effective intestinal antiseptic were outlined: (1) rapid, highly bactericidal activity against gastrointestinal pathogenic organisms, (2) low local and systemic toxicity, and

(3) limited absorption from the intestine. Two oral regimens are now primarily used: (1) an aminoglycoside agent with erythromycin base, and (2) an aminoglycoside agent with metronidazole. The regimen most often chosen in the United States is neomycin and erythromycin base, which was introduced in 1972 [4, 10, 18, 19]. In Europe and Australia, however, physicians prefer kanamycin and metronidazole or neomycin and metronidazole when oral antibiotics are administered [20].

The timing of the administration of these oral agents is critical, with an elapsed time of 19 h from the first dose to the beginning of the operation being ideal. Therefore, it is recommended that 1 g each of neomycin and erythromycin base be given at 1.00 p.m., 2.00 p.m. and 11.00 p.m. on the day before operation (6 g in total) (table 1) [18, 19, 21]. The procedure should then be scheduled for about 8.00 a.m. the next morning. If the operation is scheduled for a different time, then the times at which the oral agents are administered should be changed accordingly to preserve the 19-hour interval. Giving more than three prophylactic doses of the oral antibiotic drugs is unwarranted, for it may induce the emergence of resistant flora. Authoritative reviews on antibiotic prophylaxis in colon surgery confirm the value of the oral neomycin and erythromycin base regimen in preventing infection after elective colon resection [22–24]. However, there appears to be no convincing evidence to suggest that erythromycin base is superior to metronidazole in this clinical setting. The pharmacokinetic profile of the oral neomycin and erythromycin base preparation has been studied in healthy volunteers [25] and in patients undergoing elective colon resection [26]. Results of these studies suggest that when adequate mechanical bowel preparation is combined with this preoperative antimicrobial bowel regimen, significant intraluminal (local) and serum (systemic) levels of erythromycin and significant local levels of neomycin are present. Therefore, combining both techniques serves to prevent infection after colon operation. Other nonabsorbable, orally administered antibiotics such as rifaximin have been proposed for use in preoperative colon prophylaxis, but larger, double-blind clinical trials are needed to definitely assess their efficacy [27]. In summary, the use of preoperative oral antibiotics is an efficacious and common therapeutic adjuvant as evidenced by the report of Solla and Rothenberger [10] on the clinical regimens utilized by 372 practicing surgeons. Oral antibiotics, most commonly neomycin and erythromycin base, were used in preoperative bowel regimens by almost 92% of the responding surgeons. Furthermore, surgeons who consider oral antibiotics important in the

Table 2. Most commonly used oral and parenteral antibiotics for preoperative bowel preparations before elective colon and rectal surgery in a North American survey [4]

Oral antibiotic(s)	Number using parenteral antibiotic(s)						total
	cephalosporin			β -lactamase inhibitor combinations	metro-nidazole	other	
	1st	2nd	3rd				
Neomycin	1	3	0	1	1	0	6
Plus clindamycin	0	2	0	0	0	0	2
Plus erythromycin	22	222	26	30	20	14	334
Plus metronidazole	18	153	8	21	16	3	219
Plus erythromycin and metronidazole	6	21	5	0	0	0	32
Metronidazole	4	7	2	0	0	0	13
Plus erythromycin	4	11	0	3	0	1	19
Total with oral antibiotics	55	419	41	55	37	18	625
Total without oral antibiotics	7	46	7	4	16	6	86
Total	62	465	48	59	53	24	711

Four hundred and seventy-one respondents listed all regimens commonly prescribed by them.

success of a preoperative bowel preparation also continue to realize the necessity of effective outpatient mechanical cleansing of short duration (i.e. 24 h or less) before operation [28, 29]. This combination of mechanical preparation and oral antibiotic preparation is time-tested, and it has yielded excellent clinical results during the past two decades [22, 30].

Parenteral Antibiotics

Parenteral agents that are effective in bowel preparation for elective colon resection, either alone or in combination with an aminoglycoside, include cefoxitin, cefotetan, metronidazole and doxycycline [31]. One study, reported by the Norwegian Study Group for Colorectal Surgery, advocated mechanical cleansing together with a single preoperative intravenous dose of doxycycline for aerobic coverage and tinidazole for anaerobic coverage [32]. Other studies have shown that a single dose of cefotetan is as effective as multiple doses of cefoxitin [33]. For elective operations, most investigators recommend restriction of parenteral antibiotic prophylaxis to no more than five doses during the first 24 h, beginning from the time of surgery. Use of parenteral antibiotic regimens for longer periods of time has been associated with the development of antibiotic-resistant strains of microbes in the colonic lumen [23, 33, 34]. North American surgeons were noted to employ parenteral antibiotics alone in preoperative colonic preparation in about 8% of their pa-

tients in a 1990 survey [10], and this percentage increased to roughly 11% of patients in a 1997 survey [4]. Finally, the use of parenteral antibiotic drugs alone is employed in emergency colonic resections.

Combination of Parenteral and Oral Antibiotics

Most North American surgeons now use both oral and parenteral antibiotics along with mechanical cleansing in preoperative preparation for elective colon resection [4, 10]. In a survey of more than 500 surgeons in 1979, only 8% used systemic antibiotics alone, 37% used oral antibiotics alone and 49% used both oral and systemic antibiotics before colon surgery [35]. In a later survey of 372 board-certified colon and rectal surgeons, 88% used both oral and systemic antibiotics in their preoperative preparation, 3% used oral agents alone and 8% used parenteral antibiotics alone [10]. In a more recent survey, 87% of surgeons surveyed used both oral and parenteral antimicrobial agents, while only 11% of respondents utilized parenteral antibiotics alone [4]. Almost 78% of the responding surgeons utilized oral neomycin and erythromycin base or metronidazole combined with a perioperative parenteral antibiotic. Most surgeons start the oral antibiotic preparation in an outpatient situation the day before surgery, and then parenteral agents are added within 2 h from the time that the procedure is to begin. The rate of outpatient antimicrobial bowel preparation is increasing, and patient selection and education is critical in order to

reduce the rate of complications. The most commonly used oral and parenteral antibiotics reported in the 1997 North American bowel preparation survey are listed in table 2.

The combined oral and parenteral antimicrobial regimen has obvious appeal in that it theoretically provides both intraluminal bacterial suppression and high serum and tissue antibiotic levels. These theoretical advantages were borne out in practical success as evidenced by the metaanalysis performed by Song and Glenny [36]. These authors demonstrated that oral antibiotics alone were not as effective as when they were combined with parenteral antibiotics as well. Furthermore, using this combined antimicrobial chemotherapeutic approach, Coppa and Eng [37] documented that the rates of wound infection, intraabdominal infection and anastomotic leakage were significantly higher in patients who received parenteral cefoxitin alone than in those who received parenteral cefoxitin in addition to oral neomycin-erythromycin base. These investigators also identified the two additional risk factors for an increased infection rate, and these were a surgical resection involving the rectum and an operation that lasted longer than 215 min [37]. When the duration of the procedure was less than 215 min and the resection did not involve the extraperitoneal rectum, infection rates were low: about 3% in both antibiotic groups. Although the evidence is conflicting, it now appears that supplementing mechanical and oral antibiotic preparation with a single parenteral dose of a cephalosporin with aerobic and anaerobic activity (given intravenously within 30 min of incision) may be beneficial, particularly in operations expected to last longer than 3 h. The parenteral antibiotic serves as a fail-safe mechanism when the oral agents have been administered at the wrong times or when the operation has been delayed. Finally, this combined antibacterial regimen is also frequently utilized in patients with a partial colonic obstruction who can tolerate oral intake (table 3).

Bowel Preparation for Emergency Colon Operation

The clinical conditions that most often necessitate emergency colonic operations are acute hemorrhage, perforation, ischemia, obstruction and trauma. In these circumstances, the operation must be performed without any bowel preparation because oral antibiotic prophylaxis and mechanical cleansing are either impossible or potentially harmful.

Table 3. Antimicrobial and mechanical bowel preparation in patients with partially obstructing intestinal lesions

Environment	Hospital
Diet	clear liquids as tolerated, supplemented with parenteral fluids
Preoperative mechanical bowel preparation	fractional doses of sodium phosphate on days 3 and 2 prior to surgery
Preoperative oral antibiotic bowel preparation	erythromycin base 1 g, neomycin SO ₄ 1 g p.o. at 1.00, 2.00 and 11.00 p.m. on the day prior to surgery
Perioperative parenteral antibiotics	yes: with appropriate aerobic and anaerobic antimicrobial coverage with dose given within 30 min of incision; may be continued 24 h postoperatively

Intraoperative Lavage

Several techniques for performing intraoperative lavage have been described [38]. The authors recommend that 8–10 liters of saline irrigation be instilled over 30 min directly into the colon through a balloon-tipped catheter in the distal ileum. The balloon is inflated to occlude the ileocecal valve and prevent reflux of the irrigant into the ileum and jejunum. This technique results in uniform cleansing of the colon, and it has enabled the performance of resections and primary anastomoses in many patients who otherwise would have undergone multiple, staged procedures. However, the most important variable in the success of this intraoperative regimen is a coordinated team approach to prevent spillage of fecal contents or other mishaps.

Parenteral Antibiotics

Prevention of infectious complications after emergency colon operation depends on a proper operative technique, clinical judgment and the appropriate choice and administration of parenteral antibiotics (table 4). As in elective operations, the antibiotics chosen should be active against both aerobic and anaerobic colonic microflora. They should be given intravenously at appropriate doses, starting shortly before the operation and continuing postoperatively for 1–7 days. The duration of administration is governed by the operative findings and by whether the antibiotic regimen is intended to be preventative (in which case the antibiotics are given for 1 day) or therapeutic in order to manage intraabdominal infections (in which case antibiotics are given for 2 days or more). Many single agents or combinations of agents appear to be

Table 4. Antimicrobial and mechanical bowel preparation in patients with obstructing intestinal lesions or undergoing emergency colonic surgery

Environment	Hospital
Diet	NPO with parenteral fluids for resuscitation and maintenance
Preoperative mechanical bowel preparation	none with the possible exception of an enema to cleanse the distal rectum
Preoperative oral antibiotic bowel preparation	none
Perioperative parenteral antibiotics	parenteral antibiotics with appropriate aerobic/anaerobic antimicrobial coverage should be administered preoperatively, intraoperatively and postoperatively depending upon the length of surgery, operative findings and clinical course

NPO = Nothing per os.

equally efficacious and are currently recommended (table 5). The choice of an agent or a combination of agents, therefore, depends on local hospital prices, toxicity profiles and the surgeon's familiarity with the agents.

Topical Antibiotics or Antiseptics

Some surgeons advocate direct application to the wound of or irrigation of the wound with either antibiotics or povidone-iodine during colon resection. Solutions containing povidone-iodine should almost never be placed in the peritoneal cavity, because they are likely to be absorbed and subsequently to cause toxic effects.

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Table 5. Parenterally administered antimicrobial agents which can be used alone or combined to provide effective coverage of the mixed aerobic-anaerobic infections arising from human colonic bacteria for patients requiring preventive therapy

Aerobic choices	Anaerobic choices	Aerobic-anaerobic choices
Amikacin	Clindamycin	Ampicillin-sulbactam
Aztreonam	Metronidazole	Ertapenem
Ciprofloxacin		Imipenem-cilastatin
Gentamicin		Meropenem
Levofloxacin		Piperacillin-tazobactam
Tobramycin		Ticarcillin-clavulanic acid

Conclusion

The clinical success of elective colon and rectal surgery as measured by decreased morbidity and mortality rates secondary to infectious complications has changed radically in the last 30 years. Due to the recognition of the importance of appropriate antimicrobial therapy directed at both aerobic and anaerobic species of bacteria, combined oral and parenteral antibacterial agents and mechanical cleansing, surgeons have been able to perform single-stage colonic resections with primary anastomoses routinely. Though recent reports from Europe have called into question the necessity of mechanical cleansing, no clear consensus has been determined, and North American surgeons are steadfast in its continued use. Ultimately, the use of a three-tiered regimen – preoperative mechanical bowel preparation, preoperative oral antibiotic therapy and perioperative parenteral antimicrobial therapy – is a clinical requisite as it has been shown to be efficacious by multiple reports from the last three decades with but a few, recent reports to the contrary.

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Note Added in Proof

Pursuant to editorial request, an additional literature search of relevant articles pertaining to the need for preoperative mechanical and antimicrobial bowel preparation in colorectal surgery since the time of

the initial submission of the article was conducted by the authors. Two studies were encountered, and both lend credence to our argument that the routine use of both antibacterial and mechanical bowel preparation is not only the common practice of American colorectal surgeons, but still has not been scientifically disproven as ineffective or deleterious. Zmora et al. surveyed 1,295 members of the American Society of Colon and Rectal Surgeons as to their preference and bias in regards to preoperative bowel preparation for elective colorectal surgery [1]. 515 responses (40%) were completed, and the tallied results showed that the three-tiered approach to preoperative bowel preparation espoused in our paper is practiced by the majority of the respondents. In fact, 75%, 99%, and 98% of the respondents made routine use of two-drug oral antibiotic prophylaxis, preoperative mechanical preparation, and perioperative intravenous antibiotics, respectively. These data further serve to corroborate our position as outlined in the main body of the manuscript.

The second study by Wille-Jorgensen et al. is a metaanalysis of 6 studies looking at the specific question of whether mechanical bowel preparation changed the rate of anastomotic leakage, mortality, peritonitis, or wound infections [2]. After analysis of the data, the authors found that the rate of anastomotic leakage was greater in the patients prescribed a mechanical bowel preparation over those that were not, and this result was statistically significant. However, there was no statistically significant difference in the rate of death, peritonitis, or wound infections between the two groups of patients. Though this study would refute our claim, an invited editorial concerning the statistical accuracy of this work [3] criticized the paper for using non-homogeneous data and publication bias. Specifically, the author writes that Wille-Jorgensen et al. are 'Comparing apples and oranges and the occasional lemon'.

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Rifaximin, a Peculiar Rifamycin Derivative: Established and Potential Clinical Use Outside the Gastrointestinal Tract

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Key Words

Rifaximin · Rifamycin · Antibiotic therapy · Skin infections · Vaginosis, bacterial · Periodontal disease

Abstract

Rifaximin is a poorly absorbed semisynthetic rifamycin derivative with a broad spectrum of antibacterial activity including Gram-positive and Gram-negative bacteria, both aerobes and anaerobes. Although originally developed for the treatment of infectious diarrhea, the appreciation of the pathogenic role of gut bacteria in several organic and functional gastrointestinal diseases has increasingly broadened its clinical use. The availability of a topical formulation (a cream containing 5% of the drug) and the lack of transcutaneous absorption pointed out in both animal and human studies has allowed its topical use in skin infections. Furthermore, since the spectrum of antibacterial action of rifaximin includes many organisms (e.g. *Bacteroides bivius-disiens*, *Gardnerella vaginalis*, *Haemophilus ducreyi*) causing genital infections, including *Trichomonas vaginalis* and *Chlamydia trachomatis*, its local application in the treatment of bacterial vaginosis (BV) has been attempted. Finally, since periodontal disease, caused by plaque (an aggregate of var-

ious bacteria), can be considered a 'local' infection, intra-pocket rifaximin was tried in the treatment of periodontal infections. While the efficacy in pyogenic infections of the skin has been confirmed by several investigations, which showed an improvement of both subjective and objective parameters significantly better than that of the reference drug (i.e. chlortetracycline or oxytetracycline), the usefulness of rifaximin in BV and periodontal disease needs to be further studied in well-designed clinical trials.

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Introduction

Rifaximin, a virtually nonabsorbed antibiotic, is a semisynthetic rifamycin derivative, with a broad antimicrobial spectrum that includes most Gram-positive and Gram-negative bacteria, both aerobes and anaerobes [1, 2]. Unlike systemically available antibiotics, this antimicrobial allows localized targeting (e.g. enteric or cutaneous) of pathogens and is associated with a minimal risk of systemic toxicity or side effects [3, 4]. Provided that nonabsorbed antibiotics are as effective as systemically absorbed drugs for the target illness, their safety and toler-

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ability profiles may render them more appropriate for certain patient groups, such as young children, pregnant or lactating women, and the elderly, among whom side effects are a particular concern.

Taking into account its excellent activity against a broad range of enteropathogens, the first 'logical' indication for this gastrointestinal (GI)-targeted antibiotic was the treatment of infectious diarrhea in both adults and children [5]. The appreciation of the pathogenic role of gut bacteria in several organic and functional GI diseases [6, 7] has increasingly broadened its clinical use [2]. However, the availability of a topical formulation (a cream containing 5% of the drug) and the lack of transcutaneous absorption pointed out in both animal [8] and human [9] studies have allowed its topical use in skin infections. Furthermore, since the spectrum of antibacterial action of rifaximin includes many organisms (e.g. *Bacteroides bivius-disiens*, *Gardnerella vaginalis*, *Haemophilus ducreyi*) causing genital infections [10], including *Trichomonas vaginalis* [10] and *Chlamydia trachomatis* [11], its local application in the treatment of bacterial vaginosis (BV) has been attempted. Finally, since periodontal disease, caused by plaque (an aggregate of various bacteria), can be considered a 'local' infection [12], intrapocket rifaximin was tried out in the treatment of periodontal infections.

All these extra-GI clinical uses of rifaximin will be reviewed below.

Skin Infections

Human skin provides a great living environment for the growth of microbes. The types and numbers of microorganisms vary according to the part of the body and the age and gender of the individual [13]. Despite this, normal healthy skin presents a natural physical barrier to bacterial invasion. An intact stratum corneum layer provides a barrier to a wide variety of pathogens. The natural resistance of the skin to bacterial penetration and multiplication is not completely understood, but elements involved include the following factors [14]: (1) the inability of organisms to penetrate the keratinized stratum corneum, (2) desquamation, which sheds bacteria as it sloughs keratinocytes, (3) natural acidity of the skin (pH 5.5), (4) presence of antibacterial substances in sebaceous secretions and intracellular lipids of the stratum corneum, (5) relatively low moisture content of the skin, and (6) normal cutaneous microflora.

Changes in any of these factors can greatly influence an individual's susceptibility to infection, as can changes in the overall ability of the host to mount an inflammatory response. Nonpathogenic microbes are capable of becoming disease-producing pathogens in individuals with reduced cellular or humoral defenses or defects (e.g., immunocompromised or nutritionally compromised individuals).

Staphylococcus aureus and group A streptococci are the two most dominant pathogens involved in primary and secondary infections of the skin and minor skin wounds in outpatient settings [15]. Dominant pathogens in the hospital setting are tracked through the SENTRY Antimicrobial Surveillance Program established in 1997, which follows skin and soft tissue infections reported through a worldwide network of more than 100 hospitals. There is a remarkable consistency over the years that *S. aureus* is the most common hospital-acquired pathogen followed by *Pseudomonas aeruginosa*.

In order for bacteria to be pathogenic, they must be able to adhere to, grow on, and invade the host [16]. Bacteria possess numerous virulence genes that allow for growth in these privileged niches. Epidermal infections caused by *S. aureus* and *Streptococcus pyogenes* include impetigo and ecthyma. Dermal infections consist of erysipelas, cellulitis, and necrotizing fasciitis. The pilosebaceous unit is involved in folliculitis, furunculosis, and carbunculosis. Moreover, *S. aureus* and *S. pyogenes* produce toxins that may elicit a superantigen response, causing massive release of cytokines. The staphylococcal scalded skin syndrome, toxic shock syndrome, and scarlet fever are all superantigen-mediated.

Systemic therapy with a variety of β -lactams, macrolides and lincosamides (clindamycin) has been the cornerstone of skin infection therapy for many years [17]. However, topical antibiotics can play an important role in both treatment and prevention of many primary cutaneous bacterial infections commonly seen in the dermatological practice [18]. Indeed, while systemic antimicrobials are needed in the complicated infections of skin and skin structure, the milder forms can be successfully treated with topical therapy alone [18]. The topical agents used most often in the treatment of superficial cutaneous bacterial infections are tetracyclines, mupirocin, bacitracin, polymyxin B, and neomycin.

Due to the frequent difficulty of doing cultures before giving an antibacterial drug, topical treatment often requires the use of broad-spectrum antibiotics. However, the selection of resistant strains, especially via the use of broad-spectrum antimicrobials with marked systemic dif-

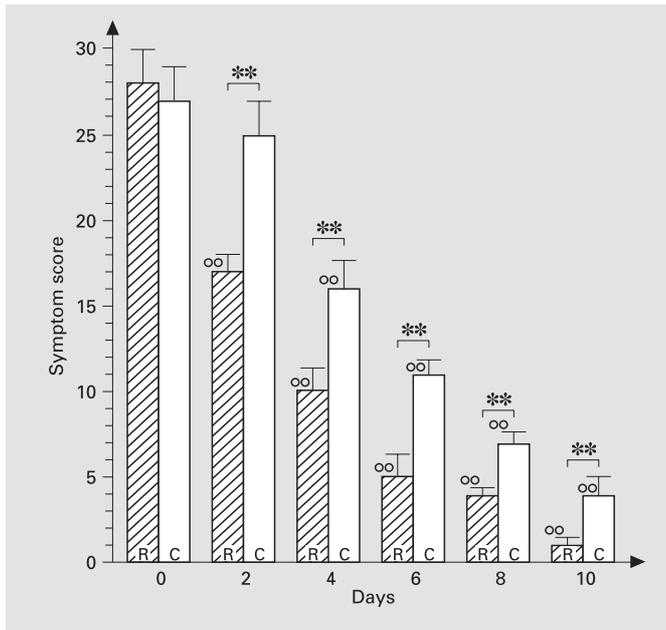


Fig. 1. Cumulative subjective plus objective symptom score after 10-day topical application of rifaximin (5% cream, n = 30, R) or chlortetracycline (3% cream, n = 30, C) of patients with pyogenic skin infections (from Della Marchina et al. [30]). ∞ p < 0.01 versus basal value (day 0); ** p < 0.01 between the two groups.

fusion even after topical application, represents a continuous challenge also in the field of skin infection [19]. *S. aureus* strains have developed worldwide a resistance to penicillin due to β -lactamase production in >90% of cases, and methicillin resistance is now a major problem with resistance levels of >50% in certain areas of the world. These resistant strains are often multiresistant, being not sensitive to erythromycin and tetracycline, with resistance to quinolone developing rapidly. Group A streptococci are still susceptible to penicillin, but increasing problems with erythromycin and tetracycline have been reported [19].

Rifaximin is broad-spectrum antibiotic, which covers many skin pathogens, whose lack of transcutaneous absorption has been well documented by both animal [8] and human [9] studies. On these grounds, a topical formulation (i.e. cream) containing 5% of the active compound was developed and tested in the treatment of pyogenic skin infections. Some open trials [20, 21] showed the efficacy and safety of the formulation and pointed out the lack of selection of resistant strains after topical application of rifaximin. In any event, drug delivery from the topical formulation is orders of magnitude higher than the

'breakpoint' level set for resistance, allowing to overcome even a resistant-defined strain.

Randomized, controlled studies (some of which were single blinded) compared rifaximin to chlortetracycline or oxytetracycline [22–30] in the therapy of skin infections caused by different pyogenic bacteria. In almost all these trials the clinical efficacy of rifaximin was significantly better than that of the reference drug. Indeed, the improvement of both subjective (i.e. itching, burning, pain) and objective (i.e. erythema, edema, papules, blebs, erosions, scabs) parameters was generally more rapid with the rifamycin derivative (fig. 1), which often led to a reduction in the duration of treatment [26]. In many studies (table 1) a bacteriological investigation was performed before and after therapy: on average, bacterial eradication at the infection site was achieved in 96.5% of the rifaximin-treated patients and in 79.2% of the tetracycline-treated ones, the overall odd ratio being 6.51 (95% CI 2.71–15.64, p < 0.0001).

The dogma of antibiotic-associated contact dermatitis has been perpetuated and generally accepted throughout the dermatological and general medical community over the past decades. The acceptance of this dogma has in part been fueled by surveys of identified allergens from contact allergy clinics [31, 32]. For example, the most recent study conducted by the North American Contact Dermatitis Group [32] showed a positive patch test response to neomycin (20% formulation) of 11.6% in a population of 3,104 patients referred for diagnostic patch testing. In this connection, being rifaximin unabsorbed, sensitization phenomena with this antibiotic are virtually unlikely. As a matter of fact, sensitization was never observed in clinical trials. One study [26] also examined whether photosensitization occurred after topical application of the drug. To this end, a skin area, where rifaximin cream had been applied, was exposed to medium and long UV rays and a photopatch test was performed. In no patient was photosensitization observed.

These data, taken together, demonstrate that topical application of rifaximin represents an effective and safe treatment of pyogenic skin infections. An additional application of this dermatological formulation would be infection prophylaxis in superficial skin wounds, particularly when used with a dressing that occludes the wound. Prophylactic topical antibiotic use makes particular sense for wounds in which the risk of infection is high, such as those that are likely to be contaminated (accidental wounds, lacerations, abrasions, and burns). Because all traumatic wounds should be considered contaminated, topical antibiotics are a logical measure to prevent wound

Table 1. Bacterial eradication at the infection site in patients with pyogenic skin infections after short-term treatment with topical rifaximin or either chlortetracycline (C) or oxytetracycline (O)

Authors, year	Ref. No.	Rifaximin	Reference drug	
Lazzaro et al., 1988	22	30/30 (100)	25/30 (83.4)	C
Marincola Cattaneo et al., 1988	23	31/32 (98.8)	21/31 (67.7)	C
Lazzaro et al., 1992	24	12/12 (100)	11/12 (91.7)	C
Palazzini and Palmerio, 1993	25	13/14 (91.7)	4/7 (57.1)	O
Moroni et al., 1993	26	16/16 (100)	11/15 (73.4)	O
Pasi and Palazzini, 1993	27	11/11 (100)	9/10 (90)	O
Cannata and Piccardo, 1993	28	7/7 (100)	11/11 (100)	C
Rafanelli et al., 1993	29	16/19 (84.2)	14/20 (70)	O
Total		136/141 (96.5)	106/136 (79.2)	

Figures in parentheses represent percentage.

infection [33]. Thanks to its pharmacokinetic and pharmacodynamic properties, rifaximin would be particularly suitable for preventing wound infection.

Bacterial Vaginosis

BV, previously known as nonspecific vaginitis or *Gardnerella* vaginitis, is the most common cause of vaginal discharge in women of childbearing age [34]. Estimates of its prevalence depend on the population studied, but include 17–19% in family planning clinics and 4–10% in student health clinics [35], increasing to 24–40% in sexually transmitted disease clinics. BV has been observed in 16–29% of pregnant women and is more frequently detected (30%) in women attending infertility clinics. It predominantly affects young, sexually active females but can occur in the absence of sexual intercourse.

This clinical syndrome is now recognized as polymicrobial superficial vaginal infection involving a loss of normal hydrogen peroxide (H₂O₂)-producing lactobacilli and an overgrowth of anaerobes [34]. While commonly found in increased numbers in women with BV, *G. vaginalis* is not invariably present [36]. The massive overgrowth of vaginal anaerobes is accompanied by an increased production of proteolytic carboxylase enzymes, which act to break down vaginal peptides to a variety of amines (especially trimethylamine) which, in high pH, become volatile and malodorous. The amines are associated with increased vaginal transudation and squamous epithelial cell exfoliation, creating the typical discharge [37]. Indeed, an increased pH (greater than 4.5 units) of the vaginal secretion, a fishy odor of the vaginal discharge

before or after the addition of 10% KOH (the so-called whiff test) and the presence of clue cells (epithelial cells with adherent coccobacilli) are, together with the symptoms, the main diagnostic criteria for BV [34, 37]. The term vaginosis is used because of the superficial nature of the infection. The wet mount usually does not show the increased number of leukocytes seen in other types of vaginitis. Although lactobacilli are essential for normal vaginal acidity, they are themselves acidophilic and hence attracted to an acid environment. The more anaerobic vaginal milieu of BV is not conducive to lactobacillary growth and dominance. What remains unknown, however, is whether the loss of lactobacilli precedes or follows this massive upheaval in flora [37].

BV is associated with an increased risk of several pathological conditions, including postoperative infection following hysterectomy and postabortion pelvic inflammatory disease [34]. In addition, 7 published studies (2 case-control and 5 cohort studies) have reported an increased risk of preterm birth in women with BV [37]. An appropriate diagnosis and treatment of BV may, therefore, lower a patient's risk of associated pelvic inflammatory disease, endometritis and adverse pregnancy outcomes. In this connection, a Cochrane systematic review [38] did show that eradicating BV significantly decreases the risk of preterm prelabor rupture of membranes without lowering that of preterm birth.

BV being an infectious disease, antibiotics represent the logical therapeutic approach [39]. However, a high spontaneous cure rate in some patients without symptoms has been found [40]. For this reason, placebo-controlled groups are recommended for antibiotic effectiveness trials. Besides systemic toxicity [4], antibiotic treatment

Table 2. Efficacy of two different rifaximin formulations in the treatment of BV in nonpregnant women (from Palazzini and Desai [52])

Rifaximin formulation	Positive vaginal cultures		Microbiological and clinical success rates, %	
	before treatment	after treatment	eradication rate	cure rate
Foam	15/15 (100)	2/15 (13.4)	86.7	86.7
Cream	16/16 (100)	7/16 (46.7)	56.2	56.2

Rifaximin (5% cream or foam) was applied vaginally from the delivery system at bedtime for 5 consecutive nights. Figures in parentheses represent percentage.

of BV can also lead to vulvovaginal yeast infections [41]. The benefit of antimicrobial therapy should, therefore, be weighed against the potential risks. Metronidazole and clindamycin are the most effective antimicrobial agents [34, 37, 39]. Erythromycin fails, because macrolide antibiotics are not effective at the acidic pH level of the vagina [42]. Ampicillin too is ineffective, probably because of poor activity against anaerobes, especially those producing of β -lactamases [43]. Other newer antimicrobials, such as ciprofloxacin [44] and amoxicillin/clavulanate [43, 45], show some effectiveness but appear to be less efficacious than metronidazole or clindamycin. Topical therapy with 2% clindamycin once daily for 7 days or 0.75% metronidazole gel once daily for 5 days has been shown to be as effective as oral metronidazole [46]. More recently, abbreviated 3-day courses of topical clindamycin and metronidazole have achieved comparable early cure rates, but long-term follow-up suggests higher rates of early recurrence [47]. Vaginal antimicrobials obviously result in minimal risk to the fetus in pregnant women.

Although metronidazole represents the most effective therapy of BV, no single treatment is completely successful in either cure or prevention. This is why several attempts have recently been made to use alternative and safer treatments [48]. The absence of lactobacilli in the vagina, a specific feature of BV, raised the question as to whether the restoration of lactobacilli by probiotic therapy can restore the normal flora and improve the chances of having a healthy term pregnancy. As a matter of fact, certain lactobacilli strains can safely colonize the vagina after oral and vaginal administration, displace and kill pathogens including *G. vaginalis* and *Escherichia coli*, and modulate the immune response to interfere with the inflammatory cascade that leads to preterm birth [49]. During pregnancy, local treatment restoring the normal acidity and vaginal flora, without systemic effect, is of course preferable to any other approach.

The very fact that several alternative therapies have been tried in BV is evidence that no single treatment provides the symptomatic and bacteriological cure rate the physician requires. Thanks to its antibacterial activity [10, 11], covering *G. vaginalis* and other pathogens responsible for urogenital infections, as well as its 'topical' antimicrobial action, rifaximin could be a suitable alternative for the local treatment of BV even in pregnant women. To this end, a 5-day and a 28-day toxicological study [50, 51] were performed in rabbits. Animals were given 1 ml of rifaximin 20% in K-Y[®] lubricating jelly applied deep into the vagina using a rubber catheter. All rabbits survived throughout the study and were free of signs of systemic toxicity. No signs of drug-related vaginal irritation were seen either by macroscopic observation or microscopic examination. A randomized, open-label pilot study [52] was performed in Italy to compare the effectiveness of two rifaximin formulations specifically developed for vaginal use (5% vaginal cream and 5% vaginal foam) in the treatment of BV. As shown in table 2, patients treated with the foam delivery system had a cure rate, which was higher than that achieved with the cream formulation. Only 1 patient in the cream arm experienced local irritation resulting in discontinuation of the treatment. The foam delivery system was, therefore, selected for further development.

Since a high spontaneous cure rate has often been observed [40], double-blind, placebo-controlled trials are needed to definitely assess the efficacy of rifaximin in the treatment of BV.

Periodontal Disease

Chronic inflammatory periodontitis occurs frequently in the adult population. The exact prevalence of the disease in the US and worldwide has been estimated to be

20% of the adult population, but has not been definitively determined because studies lack a consistent definition of the disease and a consistent methodology [53]. To this end, the American Academy of Periodontology [54] updated its classification system for periodontal diseases in 1999 to create a common terminology compatible with current scientific knowledge of periodontal diseases. Gingivitis and periodontitis are classified as separate diseases. Gingivitis is an inflammation of the marginal gingiva that does not produce attachment loss or loss of bone. Pockets that may occur with gingivitis are actually 'pseudopockets', and are due to gingival enlargement and do not involve apical migration of the gingival attachment or bone loss [55]. Periodontitis occurs when the junctional epithelium and periodontal attachment move apically along the tooth root. Alveolar bone also resorbs towards the apex of the tooth during the disease process (fig. 2). It is believed that pathogenic bacterial plaque induces an inflammatory immune response, which may compromise periodontal structures. The plaque would be comprised of pathogenic organisms, predominantly Gram-negative anaerobic bacteria [56], large enough in numbers to effect a response in a susceptible host. Genetic and systemic factors may affect these events [55]. Disease severity ranges along a continuum of slight to severe, and localized to generalized, depending upon the amount, location and rate of attachment loss. Chronic periodontitis often affects different areas of the mouth to different degrees. It is usually progressive, characterized by bursts of disease progression followed by periods when the disease is more quiescent. If left untreated, tooth loss may result due to the progressive nature of the disease [55, 57].

It is now well accepted that the goal of all periodontal therapies is to reduce, eliminate and/or repair the damage to the periodontal structures resulting from the deleterious effects of the host response to pathogenic organisms [55]. The nature and severity of chronic periodontitis are a result of the interactions between microbial factors and a susceptible host. Pathogenic bacteria in the gingival sulcus or periodontal pocket are necessary for periodontitis to occur, but bacteria alone are only part of the picture. It is well known that the body's local and systemic response to bacteria and their toxins determines the extent and severity of the disease. Controlling pathogenic bacterial colonization minimizes the host response. This is best accomplished by routine brushing, flossing and otherwise disrupting plaque and biofilm. The recommendations of dental professionals to the patient for plaque control procedures and products are essentially to achieve a healthy periodontal attachment for life. Likewise, nonsurgical ap-

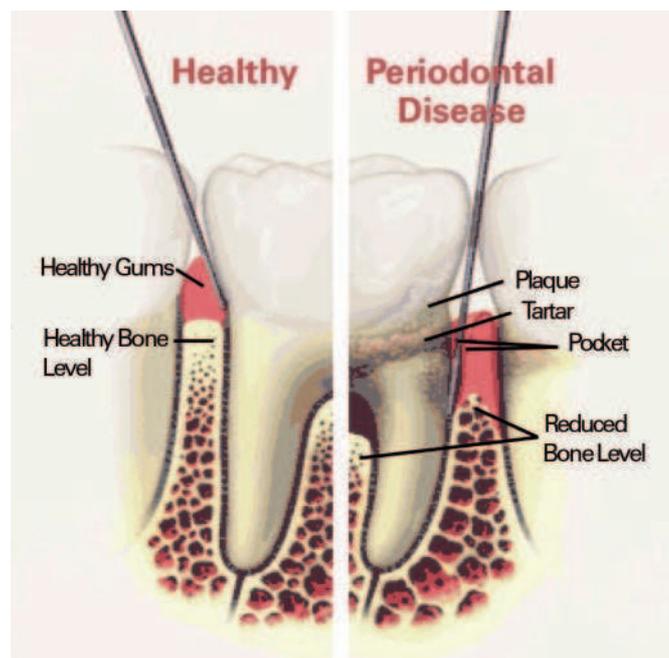


Fig. 2. Pathophysiological sequence of events in periodontal disease. Bacteria produce byproducts (e.g. toxins or enzymes) which, along with mucus, constantly form a sticky, colorless 'plaque' on teeth. If not removed, plaque can harden and form bacteria-harboring 'tartar' around teeth. Tissue that attaches the gums to the teeth can be destroyed by the irritants of plaque. If this is the case, gums pull away from the teeth and small pockets arise between the teeth and gums. The pockets then become filled with more plaque, deepen, and it becomes impossible to clean plaque out. At this stage the bone structure supporting teeth can actually be destroyed (courtesy of Drs J. Chavez and S.E. Zaragoza, El Paso Community College, Texas, USA).

proaches such as scaling and root planing (SRP) as well as the antimicrobial use have all been proven to improve periodontal health [58]. Several different surgical therapies have been evaluated as a last resort, but clinical outcomes are less predictable due to a variability in factors such as the dental surgeon's skill, experience, bias, procedures, and reproducibility of outcome measures [59].

Although the mechanical periodontal treatment alone is adequate to ameliorate or resolve the clinical condition in most cases, adjunctive antimicrobial agents, delivered either systemically or locally, can enhance the effect of therapy. In this connection, two recent meta-analyses [60, 61] have shown that systemic antimicrobials in conjunction with SRP can offer an additional benefit over SRP alone in the treatment of periodontitis both in terms of change in the clinical attachment level and probing pocket depth. When examining the effect of individual antibiotics, it was found that there were statistically significant improvements in clinical attachment level with tetracy-

Table 3. Inhibitory activity of rifaximin on microorganisms isolated from periodontal pockets in patients with periodontal disease (from Mangiante et al. [70])

	MIC ₅₀ µg/ml	MIC ₉₀ µg/ml	Range µg/ml
<i>Bacteroides fragilis</i>	0.1	12.5	0.025–100
<i>Bacteroides</i> spp.	0.1	50	0.025–100
<i>Clostridium</i> spp.	0.4	50	0.0125–100
<i>Fusobacterium</i> spp.	0.4	50	0.05–100
<i>Bifidobacterium</i> spp.	0.8	6.2	0.4–50
<i>Eubacterium</i> spp.	0.2	0.4	0.05–6.2
<i>Propionibacterium</i> spp.	0.2	12.5	0.025–12.5
<i>Peptococcus</i> spp.	0.1	3.1	0.025–100
<i>Peptostreptococcus</i> spp.	0.2	25	0.025–100
<i>Streptococcus</i> spp.	0.2	32	0.025–100
<i>Veillonella</i> spp.	0.1	6.0	0.025–100

cline and metronidazole [61]. The advantages of systemic antibiotics include the ability of the drug to control pathogens in all periodontal tissues and other sites where they may have an impact. However, systemic antibiotics may not reach the needed concentrations in the periodontal pocket, may have side effects and interactions, may be subject to compliance issues and may actually select resistant strains. Due to all these concerns, they are not routinely used in chronic periodontitis but reserved for selected patients, such as refractory cases, where often antimicrobial combinations are used [62].

Over the past 20 years, locally delivered, anti-infective pharmacological agents, most recently employing sustained-release vehicles, have been introduced. As outlined in a recent systematic review [63], controlled clinical trials have consistently shown that these formulations, along with SRP, have resulted in a clinically and statistically significant increase in the percentage of patients achieving predetermined periodontal benefits compared to scaling and SPR alone. Some trials have also shown that an antimicrobial agent alone can reduce probing depths as much as SRP alone [63]. Here again, tetracyclines [64] and metronidazole [65] are the most widely used antimicrobials.

The broad antibacterial activity of rifaximin as well as its ‘topical’ action make this antibiotic suitable for intrapocket administration in periodontal disease. As a matter of fact, local application of rifaximin compares well with tetracyclines and metronidazole in other extra-GI diseases, i.e. skin infections and BV, respectively (see above). On the other hand, rifampicin (rifampin), another rifamycin derivative, has been successfully used in the treatment

of some dental infections, including chronic periodontitis [66–69].

In a preliminary study [70], intrapocket administration of rifaximin in patients with periodontal disease was studied from both microbiological and clinical points of view. The activity of this antibiotic against the microorganisms isolated from periodontal pockets is shown in table 3. The high local concentration of rifaximin should likely exceed the observed MIC values. Indeed, its topical application was followed by a quick disappearance of anaerobic bacteria and a marked reduction of aerobic microorganisms. This was accompanied by a significant clinical improvement.

In the above trial [70] rifaximin was dissolved in chloroform and applied by repeated painting. After the solvent had dried a red sludge persisted over the dental structures allowing a continuous antimicrobial effect. Better delivery systems, such as subgingival controlled release preparations [12], are, however, needed to fully exploit the rifaximin potential in periodontal disease. In this connection, a gum-like device [71] has been developed that allows a controlled and continuous release of the antibiotic within the oral cavity. Large double-blind controlled trials using this and other formulations are now needed to establish the best therapeutic regimen for this indication.

Conclusions

Due to its excellent activity against a broad range of enteropathogens as well as its lack of absorption after oral administration the first ‘topical’ application of rifaximin was within the GI tract [1, 2]. The appreciation of the pathogenic role of gut bacteria in several organic and functional GI diseases [6, 7] has broadened its clinical use, which now goes beyond the original indication, i.e. GI infections.

It was later realized that rifaximin can be applied directly to the skin and the vaginal and gingival mucosae without absorption and without any risk of irritation. The spectrum of antimicrobial activity being large enough to cover many of the pathogens involved in skin infection, BV and periodontal disease, topic application of this antibiotic was attempted in these clinical conditions. While the efficacy in pyogenic infections of the skin has been confirmed by several investigations, the usefulness of rifaximin in BV and periodontal disease needs to be further studied in well-designed clinical trials. No doubt, new clinical applications of this peculiar antibiotic will appear in the future.

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