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The impact of proton pump inhibitors on the human gastrointestinal microbiome

Daniel E. Freedberg, MD, MS¹, Benjamin Lebwohl, MD, MS¹, and Julian A. Abrams, MD, MS¹

¹Division of Digestive and Liver Diseases, Columbia University Medical Center, New York, NY, USA.

Abstract

Potent gastric acid suppression using proton pump inhibitors (PPIs) is common in clinical practice yet may have important effects on human health that are mediated through changes in the gastrointestinal microbiome. Acting through pH-dependent or pH-independent mechanisms, PPIs have the potential to alter the normal microbiota throughout the human gastrointestinal lumen. In the esophagus, PPIs change the normal bacterial milieu to decrease distal esophageal exposure to inflammatory Gram-negative bacteria which may lower the risk of Barrett's esophagus. In the stomach, PPIs alter the abundance and location of gastric *Helicobacter pylori* and other bacteria, which has implications for peptic ulcer disease and gastric malignancy. In the small bowel, PPIs cause polymicrobial small bowel bacterial overgrowth and have been associated with the diagnosis of celiac disease. In the colon, PPIs associate with incident but not recurrent *Clostridium difficile* infection, putatively through alterations in commensal colonic anaerobes. Our understanding of the effect of gastric acid suppression on the human gastrointestinal microbiome is incomplete but is rapidly advancing.

Keywords

Proton pump inhibitors; Gastric acid suppression; Hypergastrinemia; Human microbiome; Barrett's esophagus; *Helicobacter pylori*; Small bowel bacterial overgrowth; *Clostridium difficile* infection

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Correspondence to: Daniel E. Freedberg, Columbia University Medical Center, Division of Digestive and Liver Diseases, 630 West 168th Street, PH Building, Floor 7, New York, NY 10032. Phone: 212-342-0238; Fax: 212-342-5759; def2004@cumc.columbia.edu. Benjamin Lebwohl, Celiac Disease Center at Columbia University, 180 Fort Washington Avenue, New York, NY 10032. bl119@cumc.columbia.edu

Julian Abrams, Columbia University Medical Center, 630 West 168th Street, PH Building, New York, NY 10032. ja660@cumc.columbia.edu

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INTRODUCTION

For centuries, it has been known that dietary factors influence gastrointestinal bacteria; Dorlencourt hypothesized that pH differences between breast milk and cow's milk explained the higher proportions of *Lactobacillus* observed in the stools of breastfed children.¹ Today, the role of gastric acidity in the human gastrointestinal microbiome is intertwined with the development and increasing use of proton pump inhibitors (PPIs). Other medications can alter the pH of the human gastrointestinal lumen. However, PPIs are the most potent, the most common, and have received the most attention. This review focuses on PPIs and covers the physiology of gastric acid production and suppression, and the evidence and clinical consequences of acid-related changes in the normal microbiome.

PROTON PUMP INHIBITORS AND GASTROINTESTINAL ACIDITY

Normal gastrointestinal acidity

Acidity within the human gastrointestinal tract varies by anatomic location and is part of essential physiologic processes including digestion and nutrient absorption.² In the stomach, luminal pH can approach 1.0; gastric acid plays a role in breakdown of food particles, and the pH-dependent separation of intrinsic factor from R-protein.³ Outside of the stomach, luminal pH is often discussed in the context of optimizing drug delivery. In general, pH tends to rise gradually from 6.5 in the small bowel to a high of 7.5, drop in the cecum (to as low as 5.5), and again rise gradually in the left colon to a high of 6.5 – 7.0.⁴ The invariant pattern of gastrointestinal pH seen between individuals suggests that pH plays crucial physiologic roles throughout the gastrointestinal tract. Local pH partially determines the absorption of biotin and folate in the small bowel,^{5,6} vitamin B12 in the distal ileum,⁷ and calcium and other electrolytes in the colon.⁸ Thus, in addition to the influence that pH exerts on the microbiome, gastrointestinal acidity is important and tightly regulated.

Physiology of gastric acid production

Food, stress, and other central and hormonal mechanisms stimulate gastric acid secretion acting via autonomic and paracrine signals. The primary signals are gastrin from pyloric and duodenal G-cells, acetylcholine from postganglionic neurons in the gastric submucosa, and histamine from enterochromaffin-like cells; the common target of these signals and the acid-producing cell of the stomach is the parietal cell.⁹ In response to stimuli, transmembrane H^+/K^+ -ATPase pumps are translocated from tubulovesicles into parietal cell canaliculi, increasing their concentration on the cell surface by 10-fold. These powerful pumps then acidify the stomach by utilizing ATP for energy to drive protons or hydronium ions against enormous concentration gradients.¹⁰

Proton pump inhibitors

Proton pump inhibitors were independently synthesized by two companies from 2-pyridylthioacetamide by screening modified compounds (Figure 1); the first PPIs were omeprazole (1988) and lansoprazole (1991).¹¹ There were initial safety concerns surrounding omeprazole, which was linked to increased risk for gastric carcinoids.¹² Subsequent studies suggested that PPIs did not confer increased risk for malignancy and

more PPIs were developed including enantiomers (esomeprazole and dexlansoprazole) of the original PPIs.¹³ There are currently 7 PPIs available in the United States by prescription, 2 PPIs (omeprazole and lansoprazole) that are available over-the-counter, and 3 PPIs (omeprazole, lansoprazole, and pantoprazole) which are available as generics. Because PPIs are metabolized through the hepatic cytochrome P450 system, drug levels can vary between formulations for individuals with certain pharmacogenetic characteristics.¹⁴ However, there is little evidence that the various PPI formulations differ significantly in clinical efficacy or in side effects.¹⁵

All PPIs are pro-drugs that are concentrated in a pH-dependent manner in the canaliculi of parietal cells. PPIs are concentrated within acidic parietal cell canaliculi, protonated, and covalently bound to cysteine residues of parietal cell H⁺/K⁺-ATPase antiporter pumps.¹⁶ Because stimulation at the prospect of food causes H⁺/K⁺-ATPases to be translocated into parietal cell canaliculi, PPIs are most effective if taken before meals when the maximal number of H⁺/K⁺-ATPases are available as targets. Once bound by PPIs, parietal cell H⁺/K⁺-ATPases are irreversibly fixed into an inactive configuration, which lasts approximately 24 hours until more H⁺/K⁺-ATPases can be inserted from resting intracellular vesicles into the apical membrane of the parietal cell. The key to the tremendous efficacy of PPIs is that they inhabit the end pathway of gastric acid production and thus, unlike other acid suppressive medications, cannot be overwhelmed by normal physiologic compensatory mechanisms.

In the stomach, PPIs induce profound hypochlorhydria. Serum concentration peaks after 2–5 hours; after 3–4 hours, a single oral PPI dose will raise gastric pH in most patients from 2.0 to over 6.0, a 10,000-fold change.¹⁷ The pH-raising effect of PPIs persist in the proximal duodenum, but are attenuated by the distal duodenum. In a study of healthy volunteers who underwent continuous pH monitoring, median pH in the distal duodenum was 5.85 after 1 week of PPIs compared to 5.95 after 1 week of placebo.¹⁸ Using wireless capsule pH measurement, there is similar overall small bowel pH between users and non-users of high-dose PPIs.¹⁹ The best available evidence thus suggests that, by the proximal jejunum, the direct pH effect of PPIs has been fully attenuated and is no longer significant.

Proton pump inhibitors have established clinical efficacy for many health conditions including peptic ulcer disease, gastroesophageal reflux, eosinophilic esophagitis, and acid hypersecretory conditions (e.g., Zollinger-Ellison syndrome). Because they are effective and are believed to be benign, PPIs have gained widespread use. They are perennially among the top three drug classes by sales in the world; one PPI, esomeprazole, was the fourth most prescribed drug by sales in the United States in 2012 and the top drug by sales through the first six months of 2013.²⁰ When used for appropriate indications, PPIs have great benefits. However, they are often prescribed in situations where they have no potential clinical benefit.²¹ Over half of all inpatients who receive PPIs do not have an appropriate indication for the drugs and, among these patients, over one third are discharged on PPIs.²² Among outpatients, 80% of PPI prescriptions are repeats and 40 to 50% are for non-specific abdominal pain.²³

Non-pH dependent effects of PPIs

The influence of PPIs on the gastrointestinal microbiome is presumed to depend upon their capacity to raise gastric pH. However, PPIs also have the potential to influence the microbiome through pH-independent mechanisms. First, proton pump inhibitors induce hormonal changes including hypergastrinemia and hyperparathyroidism that have the potential to alter the gastrointestinal bacterial milieu.²⁴ Second, PPIs can alter luminal contents to interfere with nutrient absorption and change the amount or location of bacterial food substrates. Case reports and cross-sectional studies document increased hypomagnesemia among patients on longterm PPIs, suggesting the possibility that PPIs interfere with small bowel magnesium transport.^{25,26} Finally, PPIs have been shown to bind non-gastric H⁺/K⁺-ATPases, both on human cells and on commensal bacteria and fungi.²⁷ The P-type family of ATPases, which includes H⁺/K⁺-ATPases, is present on fungi, *Helicobacter pylori*,²⁸ and *Streptococcus pneumoniae*,²⁹ but little is known about the effect of PPIs on specific bacteria aside from *H. pylori*.

EFFECTS OF PROTON PUMP INHIBITORS ON THE MICROBIOME

Esophagus

Proton pump inhibitors are first-line treatment for acid-related esophageal disorders including gastroesophageal reflux disease (GERD), erosive esophagitis, Barrett's esophagus (BE), suspected eosinophilic esophagitis, and non-erosive reflux disease.³⁰⁻³² Esophageal disorders are the most common reason for prescribing a PPI. Since the 1970s, there has been a 5 to 10-fold rise in BE and esophageal adenocarcinoma (EAC), with a parallel rise in GERD.^{33,34} In a large pharmacy database, over 60% of patients on long-term PPIs reported heartburn and 68% carried diagnoses of GERD, dyspepsia, or both.³⁵ But diagnostic testing was rare; only 27% of these patients underwent upper endoscopy and only 3% had testing for *Helicobacter pylori*.

The esophageal microbiome is altered in esophagitis and BE compared to normal controls.³⁶ A study of distal esophageal specimens from 34 subjects who had esophagitis, BE, or an endoscopically normal esophagus found that the microbiome could be separated into 2 types: a pattern dominated by *Streptococcus* that associated with a normal esophagus, and a pattern dominated by Gram-negative anaerobes or microaerophilic bacteria that associated with esophagitis or Barrett's. These Gram-negative bacteria may increase esophageal inflammation by activating Toll-like receptor 4 and the NF- κ B pathway through surface lipopolysaccharides (LPS).³⁷ Alternatively, these bacteria may increase distal esophageal acid exposure by lowering lower esophageal sphincter tone or by delaying gastric emptying.^{38,39}

Proton pump inhibitors are believed to protect against progression of Barrett's esophagus to EAC by decreasing distal esophageal mucosal acid exposure. PPIs simultaneously alter the distal esophageal microbiome in ways that may affect inflammation and carcinogenesis. The mucosal-associated microbiota of the distal esophagus, which is altered in patients with esophagitis or BE,^{36,40} is further modified by PPIs. A study of 34 patients with Barrett's, esophagitis, or a normal distal esophagus used 16S rRNA gene sequencing to assess the

microbiome from distal esophageal biopsies and gastric aspirates, comparing results before versus after PPIs.⁴¹ Before PPIs were administered, there were no major differences in distal esophageal mucosal bacteria comparing patients with esophagitis/BE to controls. After PPIs were administered, however, there were significant increases in distal esophageal *Lachnospiraceae*, *Comamonadaceae*, and unclassified Clostridial families. The family *Methylobacteriaceae*, which were increased in gastric aspirates among BE/esophagitis patients before PPIs, were dramatically depleted in these patients after PPI therapy. This bacterial family has also been associated with inflamed tissue in patients with inflammatory bowel disease and found in patients with irritable bowel syndrome, suggesting that these bacteria can only thrive on altered mucosa.⁴²

Although *Helicobacter* is not a dominant organism in the esophagus, *H. pylori* exerts control over the distal esophageal microbiome. There is a strong inverse correlation between *H. pylori* infection (especially *cagA* positive *H. pylori*) and Barrett's esophagus or EAC.^{43,44} A recent study by Fischbach *et al* investigated the role of acid suppression in the *H. pylori*-Barrett's relationship.⁴⁵ The authors found that the odds ratio for the association between *H. Pylori* and BE was 0.56 among those who used PPIs compared to 0.90 among those who did not, implying that PPIs augment the protective effects of *H. pylori* for BE. These results are surprising because PPIs have powerful anti-*H. pylori* activity and *H. pylori* appears to be protective for esophageal neoplasia. The most likely explanation is that the direct protective effects of PPIs in Barrett's (via decreased distal esophageal acid exposure) outweigh indirect and less potent anti-*H. pylori* effects. Future studies should further elaborate the influence of PPIs in the *H. pylori*-Barrett's relationship and determine the precise mechanisms by which PPIs alter the distal esophageal microbiome.

Stomach

Proton pump inhibitors are a mainstay of *H. pylori* eradication therapy and have direct bacteriostatic activity against *H. pylori*⁴⁶ as well as indirect *Helicobacter* activity via increases in gastric pH. Because *H. pylori* and an acidic environment are necessary for the formation of most gastric and duodenal ulcers, PPIs effectively prevent peptic ulcer disease and dramatically speed the healing of ulcers that have already formed.⁴⁷ PPIs are often used in non-ulcer dyspepsia and other functional gastric conditions, although their utility under these circumstances is less clear.

The acidity of the stomach distinguishes the gastric niche from the rest of the human gastrointestinal tract and determines the composition of the gastric flora. *H. pylori* is the dominant microorganism of the stomach, accounting for at least 70% of the gastric microbiome by 16S rRNA sequencing in positive individuals.⁴⁸ Gastric acidity both allows *Helicobacter pylori* to thrive and is influenced by the presence of *H. pylori*. Acid suppression with PPIs decreases *H. pylori* abundance and, in antrum-predominant infection, shifts *H. pylori*'s location to the corpus; meanwhile, corpus-predominant *H. pylori* infection can cause atrophic gastritis and achlorhydria.⁴⁹

PPIs cause gastric bacterial overgrowth, and PPI-induced gastric bacterial overgrowth is related to *H. pylori* infection. *H. pylori*-infected individuals have greater pH changes with PPIs than do uninfected individuals and are consequently more susceptible to overgrowth.⁵⁰

When *H. pylori* is absent, dominant gastric bacteria include oral flora such as *Streptococcus* (primarily in the mitis group)⁵¹ and common commensals such as *Lactobacillus* and *Clostridium* spp. that are seen elsewhere in the gastrointestinal tract.^{52,53} When gastric pH is raised above 4.0 by PPIs, *Lactobacillus* spp., *Streptococcus* spp., and other gastric bacteria proliferate and can cause nausea, bloating, and altered concentrations of upper GI anaerobes, which in turn affects conjugation of bile acids and can lead to diarrhea.^{54,55}

In susceptible individuals, chronic *H. pylori* infection leads to multifocal atrophic gastritis, gastric epithelial dysplasia, and gastric cancer.⁵⁶ This stepwise inflammatory process, termed the Correa cascade, has been demonstrated in animal models and corroborated by human studies; in 1994, *H. pylori* was recognized as a class I (definite) carcinogen by the World Health Organization.⁵⁷ Because of improved hygiene and increased use of antibiotics, *H. pylori* infection is declining in the developed world. However, in areas at high risk for gastric cancer, PPIs have been successfully used with antibiotics to eradicate *H. pylori* for the chemoprevention of gastric cancer. Two large, randomized and placebo-controlled trials have been conducted in areas in China with very high baseline rates of gastric cancer. The first study, conducted among 1,630 participants with *H. pylori* infection in the Fujian Province showed that antibiotics and PPIs decreased incident gastric cancer among those without precursor lesions after 7.5 years of follow-up.⁵⁸ A second, larger trial in the Shandong province showed a significant reduction in incident gastric cancer among all participants, comparing PPIs and amoxicillin versus placebo after 15 years of follow-up.⁵⁹ Because of this and similar data, short courses of PPIs are recommended as part of a chemopreventive strategy in high-risk individuals with *H. pylori* infection in guidelines from the United States, Europe, and Asia.^{47,60,61}

In *H. pylori* negative individuals, the effect of chronic PPI use on gastric dysplasia is less clear, and recent data suggest that *H. pylori* is not the only gastric microorganism that contributes to dysplasia and gastric cancer. Ironically, *H. pylori* does not thrive in the relatively high pH environment associated with gastric cancer. In patients with gastric cancer, *H. pylori* decreases in abundance and there is a shift towards *Streptococci* genera that are not often found in normal individuals.⁶² Recent data from mouse models have contributed to our understanding of the role of the non-*H. pylori* gastric microbiome in the pathogenesis of gastric cancer, although the high gastric pH of mice (baseline 3.0 to 4.0) may limit the ability to generalize murine microbiome findings to humans.⁶³ A well-established mouse model of gastric cancer is the transgenic INS-GAS mouse, which overexpresses gastrin and almost invariably develops gastric cancer.⁶⁴ When raised in a germfree environment, *H. pylori*-monoinfected INS-GAS mice had delayed progression of gastric dysplasia compared to mice with a complex gastric microbiome.⁶⁵ Introduction of complex microbiota or of defined species (altered Schaedler's flora) into the stomachs of INS-GAS mice was sufficient to accelerate dysplasia.⁶⁶ This interesting finding raises the possibility that PPIs, if continued after *H. pylori* eradication, could promote gastric cancer pathogenesis by causing non-*H. pylori* gastric dysbiosis that perpetuates the Correa cascade.

The preponderance of data does not support the idea that PPIs accelerate cancer in the stomach in humans through the microbiome or other mechanisms. Early animal studies of omeprazole showed increased rates of enterochromaffin (ECL) cell carcinoids, but

subsequent lifelong studies of rats failed to confirm this finding.¹² Further animal studies did not indicate risk,¹³ and longterm prospective cohort data in humans have not shown an association between PPIs and gastric carcinoids or gastric adenocarcinoma.⁶⁷

Small Bowel

The profound effect of PPIs on pH is limited to the stomach and proximal duodenum, with little-to-no effect on the pH of the majority of the small bowel.¹⁸ Nevertheless, gastric acid suppression by PPIs exerts a downstream effect on small intestinal bacterial composition. The increase in the quantity and diversity of the gastric microbiome in PPI users is paralleled by an increase in the quantity of bacteria in the proximal small bowel. A study of 450 consecutive patients undergoing glucose hydrogen breath test for suspected small intestinal bacterial overgrowth (SIBO) found that 50% of PPI users tested positively, compared to 6% of non-users.⁶⁸ Using duodenal aspirates and a diagnostic criteria of 10^3 colonic-type organisms per cc of fluid, a study of over 300 patients found that that 36% of PPI users had SIBO compared to 22% of non-users.⁶⁹ The most common organisms in SIBO patients were *Escherichia coli* (37%), *Enterococcus* spp. (32%), and *Klebsiella pneumoniae* (24%). Finally, a recent meta-analysis found a nearly three-fold increase in the risk of SIBO among adult users of PPIs compared to non-users (OR 2.28, 95% CI 1.24–4.21).⁷⁰ The association was particularly strong when the endpoint of SIBO was classified solely on the gold standard of duodenal or jejunal aspirates (OR 7.59). While individuals with SIBO are often asymptomatic, clinical sequelae can include gas and bloating sensation due to increased intraluminal carbohydrate fermentation, iron and vitamin B12 deficiency due to competitive microbial uptake, and fat malabsorption as a consequence of bacterial deconjugation of bile acids.^{71,72}

PPIs are often prescribed to provide gastric protection in patients who are co-ingesting nonsteroidal anti-inflammatory drugs (NSAIDs), but the combined use of these agents may exert a paradoxical cytotoxic effect on the small bowel. In a study of rats administered omeprazole or lansoprazole for 9 days plus celecoxib or naproxen for the final 4 days, PPI use was associated with reductions in jejunal *Actinobacteria* and *Bifidobacteria*, and exacerbated intestinal damage.⁷³ This injury appeared to be mediated by dysbiosis, because injury was ameliorated when PPI-treated rats were repleted with a *Bifidobacteria*-enriched microbiota. Also, when germ-free mice were given jejunal bacteria from PPI-treated rats, the germ-free mice had more severe NSAID-related injuries than did germ-free mice given bacteria from control rats. While this effect has not been demonstrated in humans, these results suggest that PPIs, when co-administered with NSAIDs, may potentiate cytotoxicity in the small bowel via a microbiome-mediated effect.

The rise of PPI use in recent decades has coincided temporally with an increased incidence of celiac disease, an immune-mediated enteropathy characterized by intraepithelial lymphocytosis and villous atrophy in response to the ingestion of gluten. Children with celiac disease appear to have distinct duodenal microbial characteristics including reduced *Lactobacillus* and *Bifidobacterium* and increased *Bacteroides* and *E. coli*.⁷⁴ A population-based case-control study found that a prescription of a PPI was far more likely in patients prior to being diagnosed with celiac disease compared to age and sex-matched controls (OR

4.79; 95% CI 4.17–5.51). Given the possibility that PPIs may have been prescribed in response to symptoms of undiagnosed celiac disease, a sensitivity analysis excluding all PPI prescriptions in the one year immediately preceding this diagnosis found that the effect, though diminished, remained significant (OR 2.28; 95% CI 1.67–3.10).⁷⁵ While these results do not prove causality, the potentially mediating effect of the microbiome on the PPI-celiac disease relationship warrants further investigation.

A second link between PPI use and the increased risk of celiac disease may relate to the effect of PPIs on *H. pylori*. In a large cross-sectional study of simultaneously submitted gastric and duodenal biopsy specimens to a national commercial pathology laboratory, there was a strong inverse association between *H. pylori* and celiac disease; this remained significant after adjusting for age, gender, and socioeconomic status (OR 0.48; 95% CI 0.40–0.58).⁷⁶ This apparently protective effect of *H. pylori* may be due to the local recruitment of regulatory T lymphocytes, damping the immune response to potentially antigenic dietary exposures.⁷⁷ Because PPIs exert a bacteriostatic effect on *H. pylori*, the potentially protective effect of this bacteria on celiac disease risk may be diminished by PPIs; in support of this hypothesis, the rise in diagnosis of celiac disease correlates with decreased rates of *Helicobacter* infection in western societies.

Colon

The large intestine contains most of the human gastrointestinal microbiome in part because the colonic pH of 5.5 to 7.0 is permissive for the growth of many microbial species.⁷⁸ Elegant interspecies experiments show that, when a mammalian microbiome is transplanted into a germ-free zebrafish, the microbial structure quickly changes to resemble a conventional zebrafish microbiome.⁷⁹ This suggests that very basic host characteristics – pH, temperature, and motility – are important determinants of overall microbiome structure. PPIs do not directly alter the pH of the colon, yet they may have clinically important effects on the distal gut, and interest has focused around the relationship between PPIs and *Clostridium difficile* infection (CDI).

C. difficile infection is a highly morbid form of infectious colitis that has been associated with exposure to PPIs in over thirty observational studies.^{80,81} During a period of declining use of antibiotics, the rise in CDI correlates with increased use of proton pump inhibitors (Figure 2).⁸² An association between PPIs and CDI has been found among outpatients,⁸³ inpatients,⁸⁴ and patients in intensive care units.⁸⁵ Multiple meta-analyses and population-based data support these findings.^{80,81} A program of active surveillance undertaken by the Centers for Disease Control that covers over 11 million people found that PPI exposure was 5% higher among those with CDI who did not report exposure to antibiotics, compared to those with CDI who did report exposure to antibiotics.⁸⁶

The mechanism linking PPIs and CDI is uncertain, but is believed to be via the microbiome. *C. difficile* spores are acid resistant, and acid suppression has little impact on their survival.^{87,88} Antibiotic use causes CDI by depleting commensal bacteria that normally block *C. difficile* proliferation and reducing the diversity of the colonic microbiome.^{89,90} Proton pump inhibitors cause small bowel bacterial overgrowth with predominantly colonic species; it follows that, with overgrowth in the proximal gut, an altered bacterial load is

delivered to the colon that may predispose to CDI. Distal gut bacteria interact with colonic epithelial cells and patients using PPIs have been shown to have increased colonic intraepithelial leukocytes⁹¹ and fecal calprotectin levels,⁹² suggesting colonic mucosal inflammation. Proton pump inhibitors may also directly bind colonic epithelial H⁺/K⁺-ATPases, or act on the colonic mucosa through NF-κB or other systemic immune pathways.⁹³ Further evidence supporting the hypothesis that the microbiome mediates the PPI-CDI relationship can be found in studies examining PPIs as a risk factor for recurrent CDI. Unlike studies of incident CDI, the relationship between PPIs and recurrent CDI is not clear.^{94,95} It is biologically plausible that PPIs cause incident CDI by altering the colonic microbiome and that this effect is blunted after the microbiome has already been perturbed by CDI.

Few studies have investigated the changes within the fecal microbiome that precede CDI. In a prospective cohort study of 599 patients during a *C. difficile* outbreak, decreased *Bacteroidetes* at the time of admission was associated with subsequent development of CDI.⁹⁶ When 16S rRNA sequencing was used to compare the fecal microbiome from 25 patients who developed CDI to 25 randomly selected controls who did not, the patients who developed CDI had significant depletions in Clostridiales Incertae Sedis XI, a family that belongs to the same order as *C. difficile*.⁹⁷ In humans, the combination of antibiotics and PPIs produced a pattern of reduced fecal bacterial diversity and reduced *Bacteroidetes* abundance although the effect of PPIs alone in humans is unknown.⁹⁸

Under controlled conditions, PPIs have effects on the fecal microbiome of animals. In dogs, administration of high-dose PPIs increased *Lactobacillus* and, in male dogs, reduced commensal fecal bacterial types including *Bacteroidetes*.⁹⁹ Another study used quantitative real-time PCR to assess the effect of achlorhydria on the fecal microbiome in Wistar rats treated with high-dose PPIs and humans with chronic atrophic gastritis.¹⁰⁰ They observed significant increases in the levels of *Lactobacillus* in acid-suppressed rats and achlorhydric humans compared to controls, without comparable increases in *Bacteroidetes*. These fecal microbiome changes resemble some of the alterations seen after administration of antibiotics, raising the possibility that PPIs may act like antibiotics to decrease microbiome diversity or otherwise alter normal microbiome structure and lower normal colonization resistance to *C. difficile*.⁹⁰

SUMMARY

Proton pump inhibitors irreversibly bind and inactivate gastric H⁺/K⁺-ATPases to induce profound gastric achlorhydria. PPIs are highly effective treatment for acid-related disorders but are widely overused. PPIs alter the microbiome throughout the human gastrointestinal tract with important potential consequences for human health (Figure 3). The ability of PPIs to heal erosive esophagitis and slow progression of Barrett's esophagus may be partly mediated by PPI-related decreases in Gram negative bacteria. In the stomach, PPIs have a chemopreventive effect when used for eradication of *Helicobacter pylori*, yet contribute to gastric carcinogenesis in animals by causing dysbiosis if given after *H. pylori* is eradicated. In the small bowel, PPIs may cause diarrhea through bacterial overgrowth and may be a risk factor for celiac disease. Finally, epidemiological studies show that PPIs are associated with

Clostridium difficile infection, although the mechanism linking PPIs and incident *C. difficile* remains unclear. Further research is needed to determine the effect of PPIs on the gastrointestinal microbiome and on human health.

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Key points

- Proton pump inhibitors (PPIs) have the potential to affect human health via interactions with the gastrointestinal microbiome.
- PPIs reduce esophageal Gram negative bacteria and may decrease risk for distal esophageal neoplasia.
- Given for *Helicobacter pylori* eradication, PPIs can prevent gastric cancer yet may cause gastric dysbiosis after *H. pylori* has been eradicated.
- PPIs may cause small intestinal bacterial overgrowth and are associated with the diagnosis of celiac disease.
- PPIs are associated with *Clostridium difficile* infection (CDI), although the mechanism linking PPIs and CDI is uncertain.

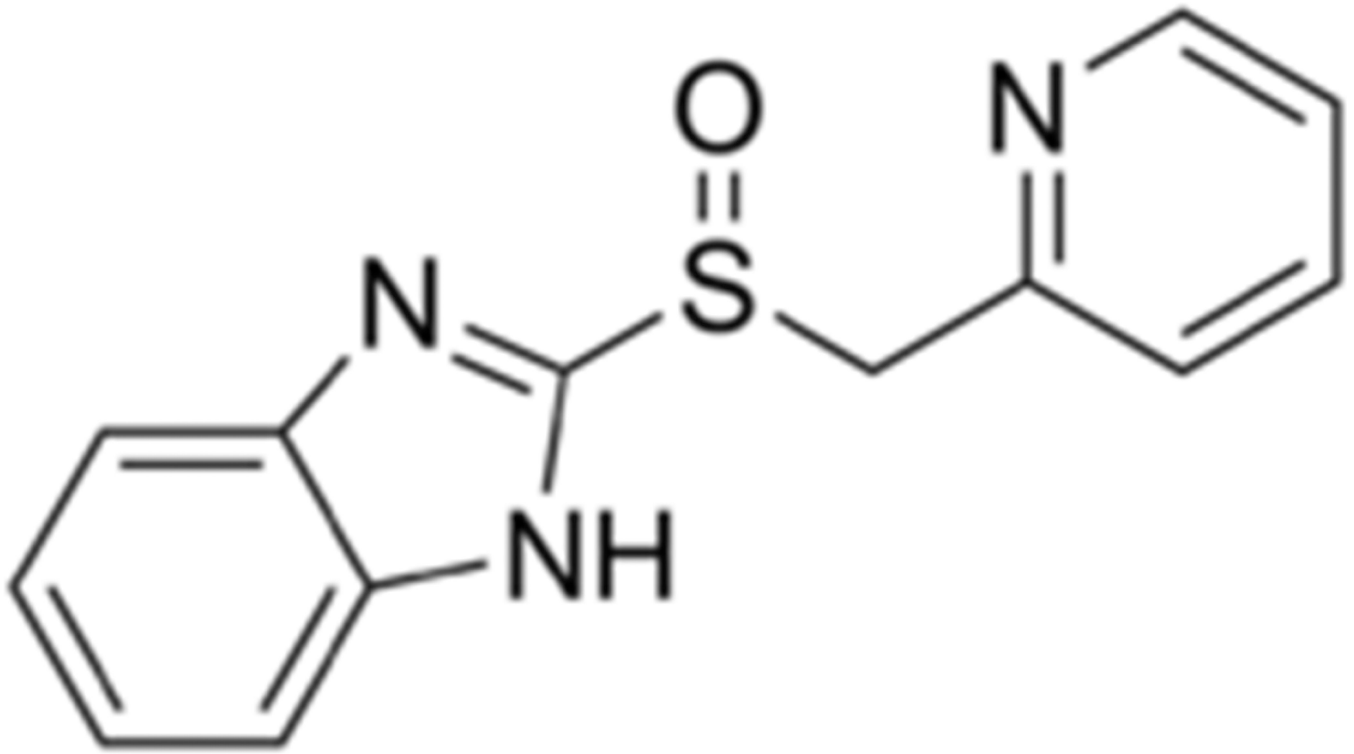


Figure 1. Common structure of proton pump inhibitors (PPIs). All PPIs share a common backbone, with a pyridine linked to a benzimidazole.

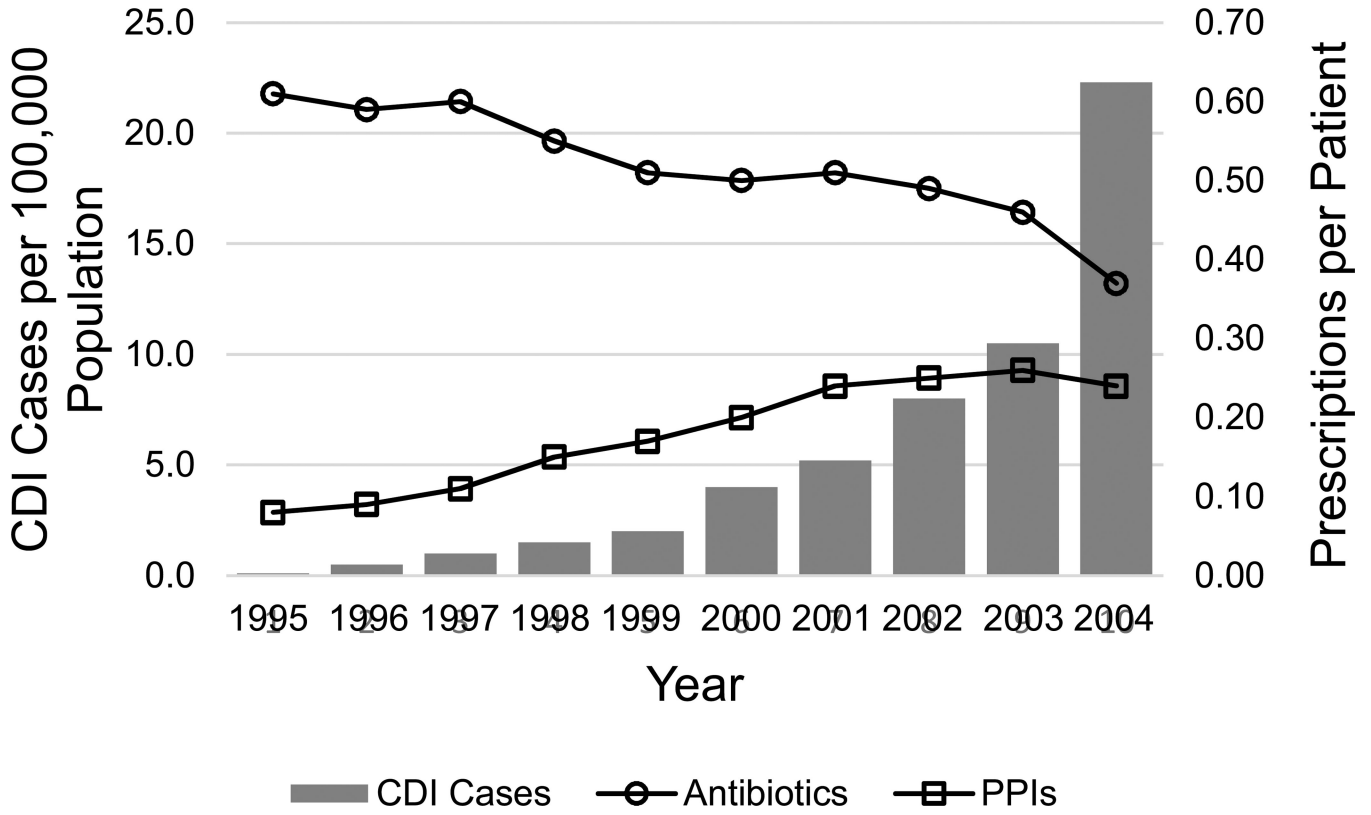


Figure 2. Corresponding rises in the incidence of *C. difficile* infection (CDI) and rate of proton pump inhibitor (PPI) use, during a time of decreasing antibiotic use. Adapted from: Dial S, Delaney JA, Barkun AN, Suissa S. Use of gastric acid-suppressive agents and the risk of community-acquired *Clostridium difficile*-associated disease. *JAMA*. Dec 21 2005;294(23): 2989–2995.

Normal pH by anatomic location

Microbiota changes with PPIs

Potential disorders impacted

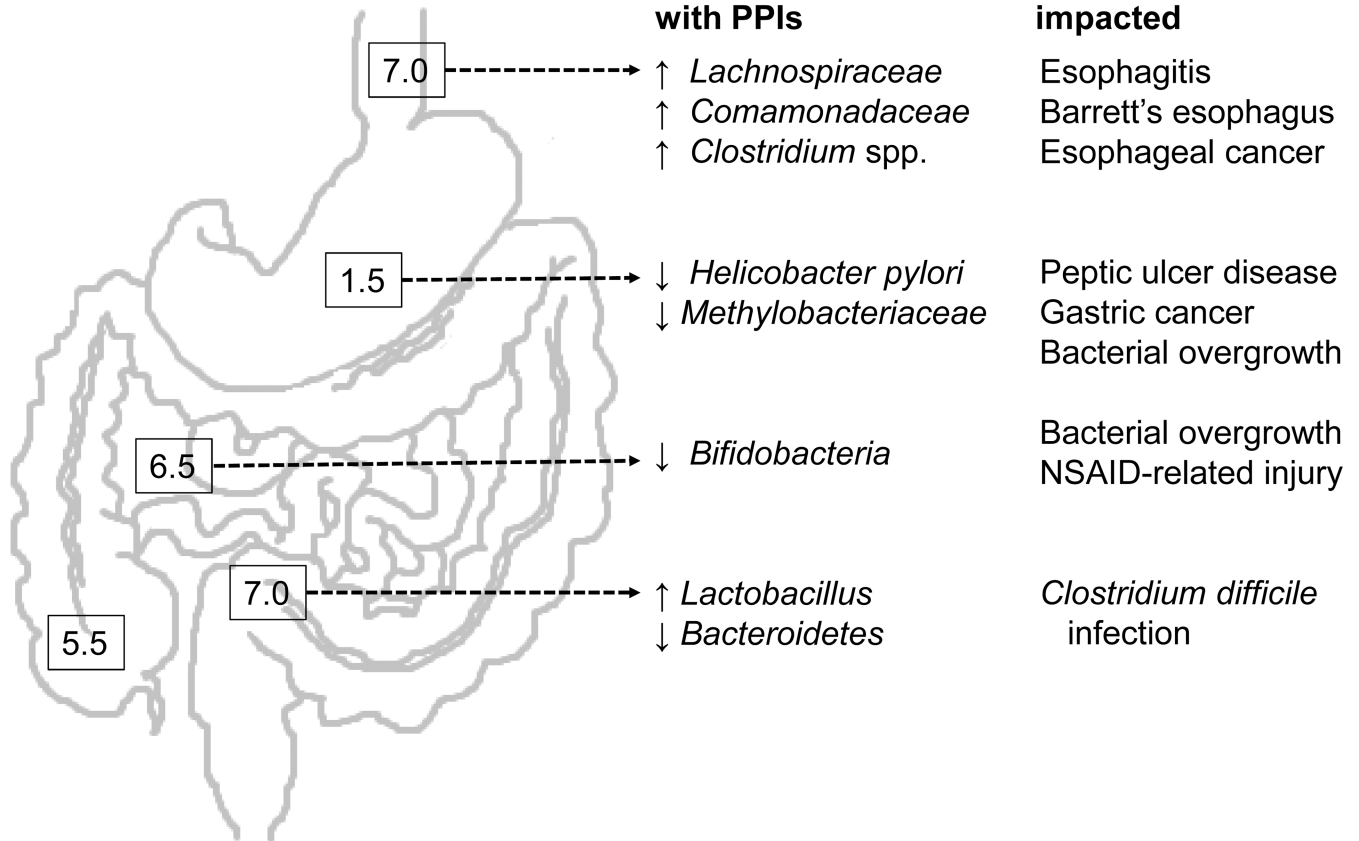


Figure 3. Bacteria that may be affected by proton pump inhibitors (PPIs) are shown by anatomical area; small arrows indicate directionality of changes with PPIs.