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The Leaky Gut: Mechanisms, Measurement and Clinical Implications in Humans

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Abstract

The objectives of this review on “leaky gut” for clinicians are to discuss the components of the intestinal barrier, the diverse measurements of intestinal permeability, their perturbation in non-inflammatory “stressed states”, and the impact of treatment with dietary factors. Information on “healthy” or “leaky” gut in the public domain requires confirmation before endorsing dietary exclusions, replacement with non-irritating foods (such as fermented foods), or use of supplements to repair the damage. The intestinal barrier includes surface mucus, epithelial layer, and immune defenses. Epithelial permeability results from increased paracellular transport, apoptosis, or transcellular permeability. Barrier function can be tested *in vivo* using orally administered probe molecules or *in vitro* using mucosal biopsies from humans, exposing the colonic mucosa from rats or mice or cell layers to extracts of colonic mucosa or stool from human patients. Assessment of intestinal barrier requires measurements beyond the epithelial layer. “Stress” disorders such as endurance exercise, nonsteroidal anti-inflammatory drugs administration, pregnancy, and surfactants (such as bile acids and dietary factors such as emulsifiers) increase permeability. Dietary factors can reverse intestinal leakiness and mucosal damage in the “stress” disorders. Whereas inflammatory or ulcerating intestinal diseases result in leaky gut, no such disease can be cured by simply normalizing intestinal barrier function. It is still unproven that restoring barrier function can ameliorate clinical manifestations in gastrointestinal or systemic diseases. Clinicians should be aware of the potential of barrier dysfunction in gastrointestinal diseases and of the barrier as a target for future therapy.

Keywords

permeability; mucus; tight junctions

INTRODUCTION

The objective of this review is to address three questions based on data almost exclusively acquired in humans: What does the “leaky gut” mean? Should clinicians diagnose leaky gut

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and, if so, how can it be diagnosed? Is the leaky gut treatable? There are vast numbers of papers that link “leaky gut” with altered microbiota in disease models in experimental animals, from allergy to non-alcoholic steatohepatitis to depression and amyotrophic lateral sclerosis. This review focuses almost exclusively on the evidence from human studies, since the clinicians need to address the three questions in the context of their patients. In addition, it focuses on the evidence of leaky gut in non-gastrointestinal diseases, which are the focus of many diseases or disorders in which leaky gut and microbiota are considered to be etiopathologically important mechanisms.

There is much folklore about the leaky gut and its relationship to microbial balance within the gut. One of the first “hits” in searching information on leaky gut on the internet provides comprehensive advice, contrasting what happens when the balance is “right” and when “out of whack”, and advice on how to get the gut microbes back into balance (BOX 1).

There is no controversy regarding barrier dysfunction in diseases resulting in intestinal inflammation and damage such as celiac or Crohn’s disease, or ulceration from nonsteroidal anti-inflammatory drugs (NSAIDs) resulting in structural abnormalities of the epithelium. The “leaky gut” is a simplistic term reflecting intestinal permeability, a function that was extensively studied in these diseases and reported in the scientific literature from 1970–1990.[1–3]

There may be several reasons for this resurgence of interest in the “leaky gut”. First, there is frustration about the lack of perceived advances in the management of common gastrointestinal symptoms such as pain, diarrhea and bloating; thus, a cause, such as leaky gut, is sought. Second, the scientific literature has promulgated “dysbiosis” in diverse states, from obesity to autism, despite evidence that this documented parabiosis[4] does not necessarily result in any metabolic or other changes in mucosal functions, including barrier function, or the role in pathogenesis of these diseases.[5] Third, there is scientific research using diverse methods that documents alterations in human intestinal barrier function in disorders such as irritable bowel syndrome or food allergy. However, there is no current gold standard with clear performance characteristics of the tests for barrier function, the diverse methods available actually measure very different endpoints, and the clinical significance and relevance are unclear. Fourth, as explained by Quigley,[6] there are popular perceptions of the barrier as a single cell thick, the epithelial layer having disruptions of intercellular connections leading to increased permeability and consequent access to the blood stream for various noxious chemicals, intact bacteria and a host of dietary and microbial components, and designation as the primary abnormality in diverse diseases such as food intolerance, fibromyalgia, chronic fatigue syndrome, and autism (all unsupported by any data).

Despite the recommendation to not self-treat, there are many resources, books and articles with recommendations on restoring the “healthy balance” including eating dirt, curing with candida, and education about the microbiome, the “human super organism” and “the good gut”.

Given the current perceptions on “leaky gut” that appear informed mostly by folklore or overreaching conclusions based on limited data, it is important to provide a balanced view of

the scientific data to facilitate the role of clinicians in addressing the nature, diagnosis and treatment of abnormal intestinal barrier function in humans. To address this, it is important to characterize the intestinal barrier, the pathways between and through epithelial cells, and measurement of intestinal permeability in humans. This provides the basis for examination of non-intestinal diseases characterized as “stress” can really break the barrier, and to review possible treatments including diet, natural substances, and medications.

The Intestinal Barrier

The intestinal barrier is a dynamic entity interacting with and responding to various stimuli. It consists of multiple elements. In the lumen, there is degradation of bacteria and antigens by bile, gastric acid, and pancreatic juice, and commensal bacteria that inhibit the colonization of pathogens by production of antimicrobial substances. The next element of the barrier is the microclimate consisting of the unstirred water layer, glycocalyx, and mucus layer which prevent bacterial adhesion by immunoglobulin A (IgA) secretion and by the physical barrier provided by the glycocalyx and mucus. Epithelial cells, connected by apical junctional complexes, have the ability to transport luminal content, but they also react to noxious stimuli by secretion of chloride and antimicrobial peptides. The Paneth cells in the epithelial layer, where they are most numerous in the crypts, also produce high quantities of defensins and several other antibiotic peptides and proteins when exposed to Gram positive and negative bacteria or bacterial products such as lipopolysaccharide. Beyond the epithelium, the lamina propria provides defense based on innate and acquired immunity cells secreting IgA, cytokines, chemokines, and mast cell proteases, as well as endocrine and secretomotor mechanisms mediated by the enteric nervous system which result in intestinal propulsive motility.[7] Some of the important transmitters are serotonin (5-HT), histamine, and cannabinoids.

The mucus layer consists of two components: an inner firmly adherent layer where bacteria are sparse and secreted peptides are protective with antibacterial functions (e.g., defensins, lysozyme); and a thicker and loosely adherent outer layer where bacteria and bacterial products are abundant. The mucus layer is thicker in the colon than in the small bowel and may reach a depth of over 800 microns, which is not much less than the height of an entire villus (range 500–1600 microns). There is regional variation in the barrier along the gut; in the small intestine, pore size increases from 4–5 Å at the villus tip to over 20 Å at the base of the crypt. In addition, the microbiota influences the barrier, and elements of the barrier impact the microbiota.[6] There are several examples in the literature demonstrating diverse effects of bacteria and their products on intestinal barrier structure or function. Thus, *Bifidobacteria* enhance barrier function in experimental necrotizing enterocolitis in mice,[8] the yeast *S. boulardii* has beneficial effects on altered intestinal microbiota and epithelial barrier defects in different pathologies,[9] and different strains of *E. coli* have opposite effects on the barrier: *E. coli* Nissle 1917 stimulates the TJ protein ZO-2;[10, 11] whereas, a prototypic translocating bacterium, *E. coli* strain C25, increases permeability.[12] Products of the bacteria such as bacterial toxins, or bacterial dehydroxylation producing secondary bile acids and short chain fatty acids produced by bacterial fermentation protect against bacteria or enhance the barrier function.[13–15]

At the level of the epithelial cells, from the apical to the basal domains of enterocytes, there are three sets of intercellular junctions: tight junction [zonula occludens (ZO)], adherens junction (zonula adherens), and desmosome. Together they comprise the apical junctional complex, which supports the dense microvillus brush border and regulates epithelial barrier function and intercellular transport.[16] The anatomy and composition of the mucosal barrier and its intercellular junctions are shown in Figure 1.[17]

In general, it is recognized that there are three distinct paracellular epithelial permeability pathways: 'leak' and 'pore' pathways which are regulated by tight junctions and define intestinal permeability; and an 'unrestricted' pathway which is associated with apoptotic leaks in pathological states, is independent of tight junctions, and provides access of luminal antigens to the lamina propria. In the presence of erosions or ulcers, bacteria gain access to the mucosa (Figure 2).[16,18]

The Importance of the Mucus Component of the Barrier

Mucus is secreted by the goblet cells and serves as the first physical defense in the barrier, preventing antigens, toxins and bacteria from directly contacting the epithelial cells. The elements of the mucus layer are highly glycosylated mucin proteins with a central protein core [abundant in serine (Ser), threonine (Thr), and proline (Pro) amino acid residues] and O-glycosylation with hexoses and hexosamines oriented almost perpendicular to the protein core like a bottle brush, forming a gel-like sieve overlying the intestinal epithelium.[19]

In the small and large intestine, mucin 2 (MUC2) is the most abundant mucus protein secreted by goblet cells. Intestinal epithelial cells (IECs) also express transmembrane mucins (MUC1, MUC3, MUC4, MUC12, MUC13, and MUC17) that remain attached to the apical surface and form the glycocalyx together with glycolipids. Other major mucus proteins secreted by goblet cells are chloride channel regulator, calcium-activated-1 (CLCA1), Fc globulin binding protein (FCGBP) which covalently binds and cross-links mucus proteins, zymogen granule protein 16 (ZG16, a small lectin-like protein that binds to Gram+ organisms), and antibodies, especially IgA. Secreted mucus mixes with Paneth cell secretions containing antibacterial peptides, lysozyme, deleted in malignant brain tumors 1 (DMBT1), and also MUC2.[19]

Immune regulators, such as antimicrobial proteins (AMPs) and IgA molecules, are released in the mucus gel in a gradient from the epithelium to the lumen, thereby reinforcing the defense against the luminal microbes.[20] The composition of the mucus layer can affect the microbiota in the gut, while the microbiota also determines the properties of the mucus gel. [21] Muc2 knock-out mice spontaneously develop colitis.[22]

Surfactants, including bile salts [chenodeoxycholate (10 mM) and hyodeoxycholate (10 mM), but not cholate (10 mM), ursodeoxycholate (10mM), or Tween-20] induce secretion of mucus.[23] In fact, bile salts impact the ability of mucus to serve as a barrier to hydrophilic and lipophilic compounds.[24] Secretion of mucus is a prelude to the epithelial damage that leads to colonic secretion by the secretagogue bile acids, chenodeoxycholic acid[25, 26] and deoxycholic acid,[27, 28] and these effects can be partly inhibited by prostaglandins. Intra-

arterial prostaglandin E2 (PGE2) evokes the secretion of intestinal mucus,[29] and PGE2 reverses NSAID-enteropathy, in part, by inducing mucus secretion.[30] Other goblet cell secretagogues are cholinergic agonists, histamine, peptide tyrosine tyrosine (peptide YY), and serotonin. Overall, there is a role for the enteric nervous system, the enteroendocrine cells, and resident immune cells in mediation of colonic mucus release.[31] Cholinergic inhibition of mucin secretion with intra-arterial atropine reduced the epithelial damage and fluid secretion secondary to 5mM sodium chenodeoxycholate in the rabbit colon in vivo.[32]

Dietary emulsifiers, like bile acids, are amphipathic, that is, they are molecules with hydrophilic and lipophilic sections that maintain fat molecules in liquid suspension or water-soluble components in a hydrophobic environment. Dietary emulsifiers interact with the multilayered endogenous mucus secretions that coat the luminal surfaces of the intestinal tract and may compromise the ability of human mucus to prevent contact between microorganisms and intestinal epithelial cells.[33] Numerous synthetic surfactant food additives (anionic, cationic or non-ionic) are used in the food industry [such as mono- and diglycerides or esters of fatty acids (E471, E473, E475)], as reviewed elsewhere.[34, 35] Some of these have been shown to increase intestinal permeability through paracellular and/or transcellular mechanisms, and some of them were also shown to inhibit P-glycoprotein or have mucolytic activity, such as the two emulsifiers, carboxymethylcellulose (CMC) and polysorbate 80 (Tween).[36]

Additionally, based on the general characteristics of surfactants, it can be predicted that they decrease the hydrophobicity of the mucus layer, which has also been shown to be associated with increased intestinal permeability.[34] Dietary emulsifiers may interact with gut microbiota and altered mucus thickness to promote colitis in *I110* and *Tlr5* knock-out mice, which are predisposed to development of spontaneous colitis[37] and increase translocation of *E. coli* across intestinal epithelial cells.[38]

Example of Increased Transcellular Permeability

The examples provided above focused predominantly on the intercellular barrier and mucus barrier. However, there is evidence that there may be altered expression of cellular transport mechanisms that may ultimately lead to intercellular barrier dysfunction and systemic inflammation. An excellent example of the potential impact of increased transcellular permeability without apoptosis or intestinal ulceration is provided by the demonstration, in mouse models of obesity and diabetes, that hyperglycemia drives intestinal barrier permeability through glucose transporter 2 (GLUT2)-dependent transcriptional reprogramming of intestinal epithelial cells and alteration of tight and adherence junction integrity.[39] These findings were demonstrated by reduced ZO-1 expression and increased fluorescein isothiocyanate (FITC)-dextran in serum after oral administration, indicating increased intestinal permeability, as well as an increased short-circuit current measured across the epithelial layer in Ussing chambers. There were also increased intestinal bacteria at systemic sites. Control experiments showed that these effects were due to hyperglycemia rather than obesity or alterations in leptin signaling.

Intestinal Permeability: Pathophysiological Mechanisms and Methods of Analysis

In this section, diverse methods of measurement and analysis are discussed, and the strengths and weaknesses are addressed. It is also important to note that intestinal permeability is influenced by several factors, which are not always considered in the publications, including the circadian cycle[40] and stress.[41]

Orally administered probe molecules

Intestinal permeability is most commonly measured indirectly in humans by the fractional urinary excretion of orally ingested probes which cross the intestinal epithelium by the paracellular pathway, enter the bloodstream, are filtered by the glomerulus, and excreted in the urine without active reabsorption in the kidney.[42] Fractional urinary excretion can, therefore, be used as an indirect measure of intestinal permeability.

The most commonly used probe molecules are saccharides. Although regional differences for preferred absorption sites have been suggested,[43] it is important to note that sucrose is only useful in the first hour post ingestion, as it is rapidly metabolized (to glucose and fructose) and, therefore, at best, provides information about gastric and duodenal permeability. Moreover, other monosaccharides, such as mannitol and rhamnose, and disaccharides, such as lactulose and sucralose, are all absorbed in the small bowel and colon, and the timing of urinary excretion provides the best way to differentiate regional analyses: 0–2 hours reflects predominantly small intestinal permeability; and 8–24 hours reflects almost exclusively colonic permeability.[44] Also, among these saccharides, sucralose is the one not metabolized by colonic bacteria; similarly, polyethylene glycol (PEG) 400 and radioactive chromium complexed with ethylene diamine tetracetic acid (^{51}Cr -EDTA) are not degraded by colonic bacteria. However, the utility of these probe molecules is somewhat compromised by the “background” ingestion of some of the sugars, particularly mannitol and sucralose.

Until relatively recently, interpretation of permeability tests was based on the following assumptions. Lactulose is a relatively large molecule, can only cross via the leak pathway or at sites of epithelial damage, and is considered a marker of barrier integrity. Mannitol, which is reputedly one-third as large as lactulose, is assumed to cross the pore pathway, which allows passage of sodium ions, water and small solutes; therefore, mannitol and other monosaccharides such as rhamnose used to be regarded as measures of surface area.[42] The inference was that the lactulose: mannitol ratio might measure the sum of leak pathway permeability and epithelial damage normalized to surface area. However, a review of the molecular sizes of the sugar probe molecules (Table 1) suggests that there is no relevant difference in the reported or estimated molecular diameters and, therefore, it appears unlikely that they traverse the epithelium through different pathways.

In fact, the mass of saccharides that is absorbed during one hour in the healthy gut following ingestion orally is typically up to 2% of rhamnose and 0.07% of lactulose administered in children in USA.[45] In healthy adults, the fractional excretion over 24 hours is $31.2 \pm 3.4\%$

(SEM) for ^{13}C -mannitol, and $0.32\pm 0.03\%$ for lactulose.[46] Therefore, despite the similarity in molecular diameters, there is a 100-fold difference in the percent of recovery of the mono- and disaccharides; however, if the larger molecule traverses the “leak” pathway, the latter might also allow passage of the small molecule, and there is no compelling evidence that the different sugars actually traverse the intestinal barrier via different pathways.

Other factors, such as tertiary molecular structure, are likely relevant to explain the marked difference in absorption ratios of monosaccharides and disaccharides which is 30- to 100-fold for the monosaccharide, mannitol, compared to the disaccharide, lactulose. In practice, modern methods of assay based on liquid chromatography-mass spectrometry accurately measure the saccharides, and the most commonly used combinations are therefore lactulose or sucralose (as disaccharides) with mannitol or rhamnose (as monosaccharides). Because of potential “contamination” by environmental exposures to mannitol and sucralose, the lactulose and rhamnose or ^{13}C -mannitol saccharides are increasingly used for *in vivo* permeability measurements. In addition, as indicated above, 0–2-hour urine collection reflects predominantly small intestinal permeability, and 8–24 hour collection almost exclusively colonic permeability.

Overall, these tests still have limited validity based on uncertainty of the normal values, lack of standardization of test procedure, and lack of validation including responsiveness of a standardized test to treatment.

***In vitro* or tissue measurements of intestinal barrier**

Several methods are used to assess intestinal permeability on biopsies taken from the human gut and measured by the transfer of probe molecules across mucosal biopsies in Ussing chambers (in association with measurements of transepithelial resistance and short circuit current measurements). Other approaches quantitate the tight junction proteins in the mucosal biopsies, or they assess the fecal supernatant in cellular monolayers or rat or mouse colonic mucosa *in vitro*.

There are differences in *in vivo* compared to *in vitro* measurements of barrier functions. The molecular size of probe molecules that can cross the epithelial barrier in humans *in vivo* is at least 10-fold smaller than *in vitro*, which shows that molecules of approximately 4–40 kDa (e.g., dextran 4 or 40) easily traverse the intestinal mucosa in a Ussing chamber *in vitro*. There are also differences that may reflect additional functional barriers *in vivo* that are excluded in the *in vitro* studies, including the lamina propria, innervation by submucosal neurons, and permeability of end-capillaries that constitute other potential barriers impeding passage of the probe molecules into the circulation *in vivo*. For example, intercellular complexes are under neurohumoral control, such as from VIP and cholinergic neurons, from the submucosal plexus,[47, 48] and these are lost in biopsied mucosa.

Endoscopic measurements of intestinal barriers in humans

Two techniques are available: first, confocal endomicroscopy, which shows leaks of intravenously administered fluorescein into the gut lumen during endoscopy[49] (e.g., in response to food-associated changes in the intestinal mucosa of patients with diarrhea-predominant irritable bowel syndrome); and second, endoscopic mucosal impedance, in

which a 2-mm diameter catheter is passed through an endoscope and placed in contact with the duodenal mucosa under direct visualization, and two circumferential sensors, placed 2mm apart on the mucosa for 0.10s, in the 4 quadrants of the duodenum with a decompressed lumen, and all fluid aspirated.[50] The studies of food-associated changes during these challenge tests provide some evidence that there can be barrier changes that may indeed support the concept of transient “leakiness” of the gut.

Abnormal Barrier Function in Intestinal Disease States

It is clear that inflammatory or ulcerating diseases result in abnormal intestinal barrier function. However, this is not the category of disease that is being associated with leaky gut, as discussed in the next section. The abnormal barrier function is well described for conditions such as inflammatory bowel disease (IBD), as well as in first degree relatives of patients with IBD,[3, 51] celiac disease and gluten sensitivity without overt celiac disease in patients with HLA-DQ2/8 (genotype associated with celiac disease,[52, 53] intestinal graft versus host disease, enteric infections and infestations, and human immunodeficiency virus infection and acquired immune deficiency syndrome (HIV/AIDs).[42] There is also extensive literature[54] documenting abnormal intestinal permeability in irritable bowel syndrome and the association of abnormal permeability with pain in IBS, although the degree of altered barrier function is clearly lower than in inflammatory bowel or celiac diseases. Longitudinal studies in patients with IBD suggest that increased intestinal permeability preceded relapsed of Crohn’s disease,[55] suggesting a pathogenetic role of the epithelial barrier in the pathogenesis of gut inflammation; in addition, IBS is highly prevalent in first degree relatives of IBD patients,[56] suggesting that intestinal permeability could be a relevant factor in the determination of symptoms.

Several other non-gastrointestinal diseases have been associated with leaky gut, based on limited or no supporting data,[6,42] including asthma, autism, Parkinson’s disease, multiple sclerosis, eczema, psoriasis, eosinophilic esophagitis, environmental enteropathy, kwashiorkor, fibromyalgia, depression, chronic fatigue syndrome, multi-organ failure syndrome (shock, burns, trauma), non-alcoholic fatty liver disease (NAFLD), alcoholic cirrhosis, obesity, metabolic syndrome, pancreatitis, and rheumatoid arthritis. Two separate groups independently studied small bowel permeability in patients with eosinophilic esophagitis; one group documented increased small bowel permeability, though the mechanism whereby this results in the eosinophilic infiltration of the esophagus is unclear; [57] the second group did not document increased small bowel permeability, but they reported improvement in the eosinophilic esophagitis with an elemental diet.[58] All of these diseases and disorders associated with possibly altered intestinal barrier function are pathological diseases, not what is usually associated with nonspecific “leaky gut”.

Leaky Gut: The Pro Arguments

The concept of a leaky gut in non-gastrointestinal diseases is supported by evidence of dysfunctional gut mucosal barrier in stress-associated conditions and the response of the altered barrier to non-drug interventions; associations of disease states with altered intestinal permeability and microbiome; and alterations in intestinal permeability as a result of gut-

directed therapy in diverse conditions including healthy people or diverse groups of children in central African countries.

Dysfunctional gut mucosal barrier due to endurance exercise and effects of non-drug intervention

Table 2 summarizes the literature on two types of conditions that result in stress to the intestinal barrier, with documented effects on intestinal permeability or biochemical evidence of mucosal damage, and restoration with a dietary, non-pharmacological intervention. These studies suggest that there are “stress” states in which there is documentation of altered barrier function and examples of normalization that would support the concept of a transient leakiness of the gut barrier. There is also recent evidence suggesting that changes in intestinal microbiota composition (characterized by increased α -diversity and changes in the relative abundance of >50% of identified genera, including increased abundance of less dominant taxa at the expense of more dominant taxa such as *Bacteroides*) and metabolism (reduced serum IL-6 and reduced stool cysteine) coincide with increased intestinal permeability (documented by increased sucralose excretion in urine after oral load) in young adults under prolonged physiological stress in the form of a 4-day cross-country ski march.[59] Discussion of the role of the microbiota in intestinal barrier function is beyond the scope of the current article. Further studies are required to explore the hypothesis that epithelial barrier dysfunction associated with mucosal enrichment of specific bacterial strains may predispose to a shift to disease-associated microbiota that eventually leads to pathological consequences such as inflammatory bowel disease in individuals with genetic predisposition.[60]

Associations of disease states with altered intestinal permeability and microbiome

From a detailed review of the literature, it is clear that animal models of disease have documented a three-point relationship: disease phenotype, barrier change, and altered microbiota. As exemplified best with chronic liver disease, the directionality of the relationship is controversial, even from animal studies. For example, one hypothesis argues that increased endogenous production of ethanol by gut bacteria (e.g., *E. coli*) caused by small intestinal bacterial overgrowth results in increased intestinal permeability, bacterial translocation and hepatic inflammation due to the translocated bacteria or their products.[61] An alternative hypothesis is that the liver disease causes a systemic inflammatory response that leads to increased intestinal permeability, with bacterial translocation and further hepatic damage.[62, 63]

In many studies in the literature, particularly in human studies, the three focal points of the relationship are not all examined and “triangulation” is therefore based on hypothesis or inference based on the association between two of these three factors. Table 3 summarizes information garnered from studies in aging, food allergy, liver disease, parenteral nutrition (or enteral exclusion), and neuropsychiatric diseases in humans or in animal models where there are no human data available. In general, the data should be regarded as hypothesis-generating, that is, the leaky gut may be a cause or an effect of the disease (as in the case of liver disease), and there may be either normal or dysbiotic microbiota that lead to inflammatory or other consequences that have impact on the disease. In some situations,

alteration of the microbiota may result in reduced severity of the disease in humans, as demonstrated with hepatic cirrhosis in response to treatment with *Lactobacillus casei* strain Shirota or VSL#3 which contains eight lyophilized bacterial strains.[64]

Alterations in intestinal permeability as a result of gut-directed therapy

A third pro argument is provided by examples from the literature of *in vivo* human studies showing alterations in intestinal permeability as a result of gut-directed therapy, as summarized in Table 4. One study worthy of specific comment is the randomized, placebo controlled trial of oral glutamine in about 100 patients showing normalization of LMR in association with improvement in the IBS-SSS (symptom severity score), stool frequency and consistency [65]. However, most of the other studies are small, used different nutrients and diverse methods, and require replication.

There is also hope that probiotics or commensal organisms improve intestinal barrier function;[66] however, the evidence to date is sparse, often based on animal models rather than human studies, the beneficial effect may be through the effects of butyrate, and the documented effects have been reported for the organism *Akkermansia muciphila*. [67] There is a micro-integral membrane protein (MIMP) that is the smallest domain of surface layer protein from *Lactobacillus plantarum*. MIMP had a significant anti-inflammatory effect in an experimental model of dextran sodium sulphate (DSS)-induced colitis.[68] This was achieved through multiple mechanisms: regulating the gut barrier (appearance of FITC-dextran in serum after oral gavage, as well as up-regulation of the expression of junctional adhesion molecule (JAM)-1, occludin and ZO-1 in the colon tissues), microbiota (increased richness and diversity, including increased *Leuconostocaceae* and *Leuconostoc*, instead of *Firmicutes* and *Clostridia* which were abundant in the DSS group), and inflammatory cytokines through the toll-like receptor (TLR)4-related pathway.[68]

These cumulated observations suggest that there are non-pathological situations that may be associated with increased permeability, and these relatively minor perturbations can be reversed with dietary, nonpharmacological approaches. Further studies of such approaches, including prebiotics and probiotics, are eagerly awaited.

Leaky Gut: The Con Arguments

As indicated by other authors,[6, 16] there are, however, important pitfalls that need to be considered and precautions to be taken in attributing biological or clinical relevance to “leakiness” of the barrier. First, altered permeability may be an epiphenomenon. For example, any inflammatory process may impair barrier integrity, and other factors such as dietary components or intraluminal factors such as bile acids can independently influence barrier function. Second, although allergens, stress, and physical activity may indeed alter intestinal barrier function, it is unclear how this predisposes to clinical consequences. Third, impaired barrier function (e.g., genetically determined defects in barrier components) does not, in isolation, lead to disease phenotype in experimental animal models of disease. Fourth, increased permeability is not necessarily deleterious, and there is no convincing evidence that an intervention that restores or improves barrier function in humans can alter the natural history of disease. Thus, for example, whereas anti-tumor necrosis factor- α

(TNF- α) therapy reduces mucosal inflammation and restores intestinal permeability in patients with inflammatory bowel disease (IBD), and butyrate, zinc, and some probiotics also ameliorate mucosal barrier dysfunction, it is still unproven that permeability manipulation should be considered as a therapeutic target in IBD.[69]

Conclusions

Although the ultrastructure and function of the epithelial barrier have been well characterized, the role of and interactions with other components of the barrier, especially the mucus layer and its perturbation, remain unclear. The role of gut barrier function is deemed to be important, but there are many unresolved questions as there are no validated clinical diagnostic tests. Although chemicals, nutrients, prebiotics, and even plant extracts (e.g., indigo naturalis) improve barrier function, there are no validated drug treatments yet, and the impact of restoring barrier function to ameliorate clinical manifestations in local gastrointestinal disease or systemic diseases is as yet unproven. Clinicians should be aware of the potential of barrier dysfunction in gastrointestinal diseases, and the potential as a target for future therapy.

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Abbreviations used:

5-HT	serotonin
⁵¹Cr-EDTA	radioactive chromium complexed with ethylene diamine tetracetic acid
AMPs	antimicrobial proteins
CLCA1	chloride channel regulator, calcium-activated-1
CMC	carboxymethylcellulose
Da	dalton
DAO	diamine oxidase
DMBT1	deleted in malignant brain tumors 1
DSS	dextran sodium sulphate
EoE	eosinophilic esophagitis
FABP	intestinal fatty-acid binding protein
FCGBP	Fc globulin binding protein

FITC	fluorescein isothiocyanate
GLUT2	glucose transporter 2
HIV/AIDSs	human immunodeficiency virus infection and acquired immune deficiency syndrome
IBD	inflammatory bowel disease
IECs	intestinal epithelial cells
I-FABP	intestinal fatty-acid binding protein
IgA	immunoglobulin A
JAM	junctional adhesion molecule
LC-PUFA	long chain polyunsaturated fatty acids
LMR	lactulose mannitol excretion ratio
LPS	lipopolysaccharide
LRR	lactulose-rhamnose ratio
MIMP	micro-integral membrane protein
MUC	mucin
NAFLD	non-alcoholic fatty liver disease
NASH	non-alcoholic steatohepatitis
NSAIDs	nonsteroidal anti-inflammatory drugs
PEG	polyethylene glycol
PUFA	polyunsaturated fatty acids
Pro	proline amino acid
SEM	standard error of the mean
Ser	serine amino acid
SIBO	small intestinal bacterial overgrowth
SLR	sucralose to lactulose ratio
Thr	threonine amino acid
TJ	tight junction
TLR	toll-like receptor
TNF-α	tumor necrosis factor- α

TPN	total parenteral nutrition
Tween	polysorbate 80
ZG16	zymogen granule protein 16
ZO	zonula occludens

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BOX 1.**Leaky Gut**

<https://www.healthyway.com/content/the-truth-about-leaky-gut-syndrome-what-it-is-and-why-you-want-to-avoid-it>

BALANCE When the microbial balance in your gut is right, your whole body functions the way it's supposed to. But when that balance gets out of whack—say because of chronic stress, chronic constipation, exposure to environmental toxins like pesticides, eating a poor diet, or taking an antibiotic that wipes out a lot those microbes—the “bad” bacteria cut holes in the fence and some of them, along with food particles and toxins, leak into the bloodstream. When your immune system sees organisms where they don't belong, it attacks, causing irritation and inflammation.

CAUSES: Leaky gut has so many possible causes – so many possibly symptoms

CONSEQUENCES: “The leakage in leaky gut may be responsible for a huge variety of health issues, ranging from minor (bloating, cramps, fatigue, food allergies and sensitivities, gas, and headaches) to “bigger things”: autoimmune conditions, depression and other mood disorders, diabetes, inflammatory bowel disease, and multiple sclerosis”.

TREATMENT: “Functional medicine”: get gut microbes back into balance: *Multi-Step program*

- Remove foods that create problems e.g. gluten, sugar, and dairy.
- Replace with foods less likely to irritate gut: Fermented foods e.g. sauerkraut, kimchi, yogurt, kefir, and pickles, are healing foods
- Repair the damage with supplements: L-glutamine (heal the intestinal lining), vitamin D, zinc, and omega-3 fatty acids (such as fish oil).
- Repopulate your good gut bacteria: probiotics or get a transplant from another person
- One of the biggest leaky gut red flags is having issues with a variety of foods.
- Talk with your healthcare provider: You might have leaky gut syndrome.
- Do not, however, try to treat it yourself

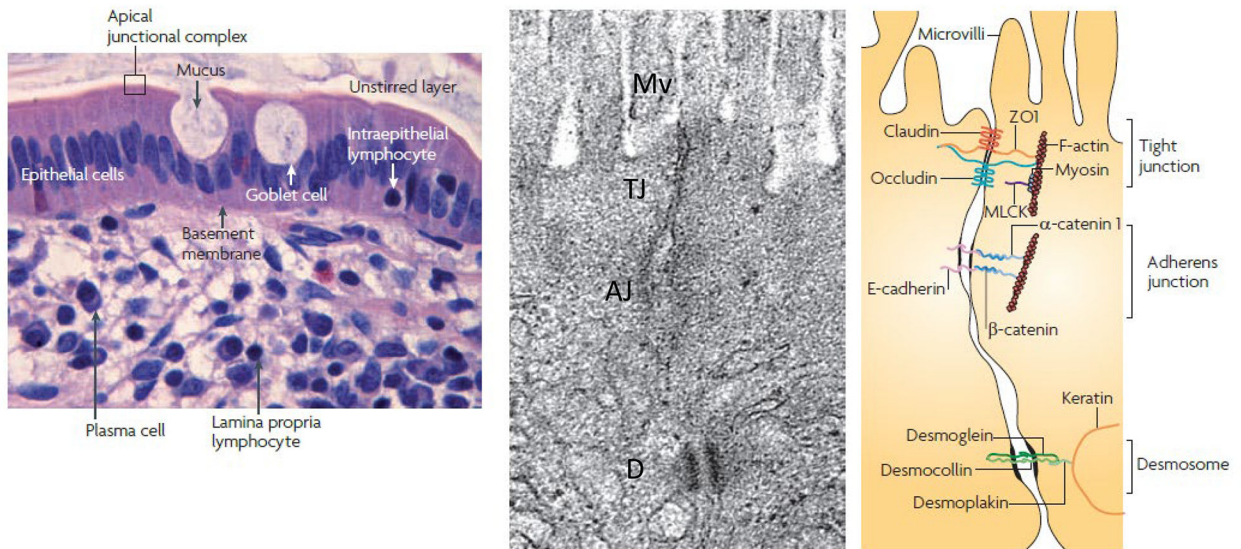


Figure 1: Anatomy of the mucosal barrier.

Left panel: In the human intestinal mucosa, composed of columnar epithelial cells, lamina propria (with its immune cells) and muscular mucosa, the goblet cells synthesize and release mucin, and the unstirred layer is immediately above the epithelial cells. The tight junction is a component of the apical junctional complex, and it seals paracellular spaces between epithelial cells.

Middle and Right Panels show an electron micrograph and the corresponding line drawing of the junctional complex of an intestinal epithelial cell. The key elements of the tight junction are the zona occludens and zona adherens, each of which is made up of different components. Just below the base of the microvilli (Mv), the plasma membranes of adjacent cells seem to fuse at the tight junction (TJ), where claudins, zonula occludens 1 (ZO1), occludin and F-actin interact. E-cadherin, α -catenin 1, β -catenin, catenin δ 1 (also known as p120 catenin; not shown) and F-actin interact to form the adherens junction (AJ). Myosin light chain kinase (MLCK) is associated with the peri-junctional actomyosin ring. Desmosomes, located beneath the apical junctional complex, are formed by interactions between desmoglein, desmocollin, desmoplakin and keratin filaments.

In general, diffusion through claudins and occludin within the membrane is energy-independent, whereas ZO-1 facilitates exchange between tight junction and cytosolic pools through energy –dependent mechanisms

Reproduced from ref. 17.

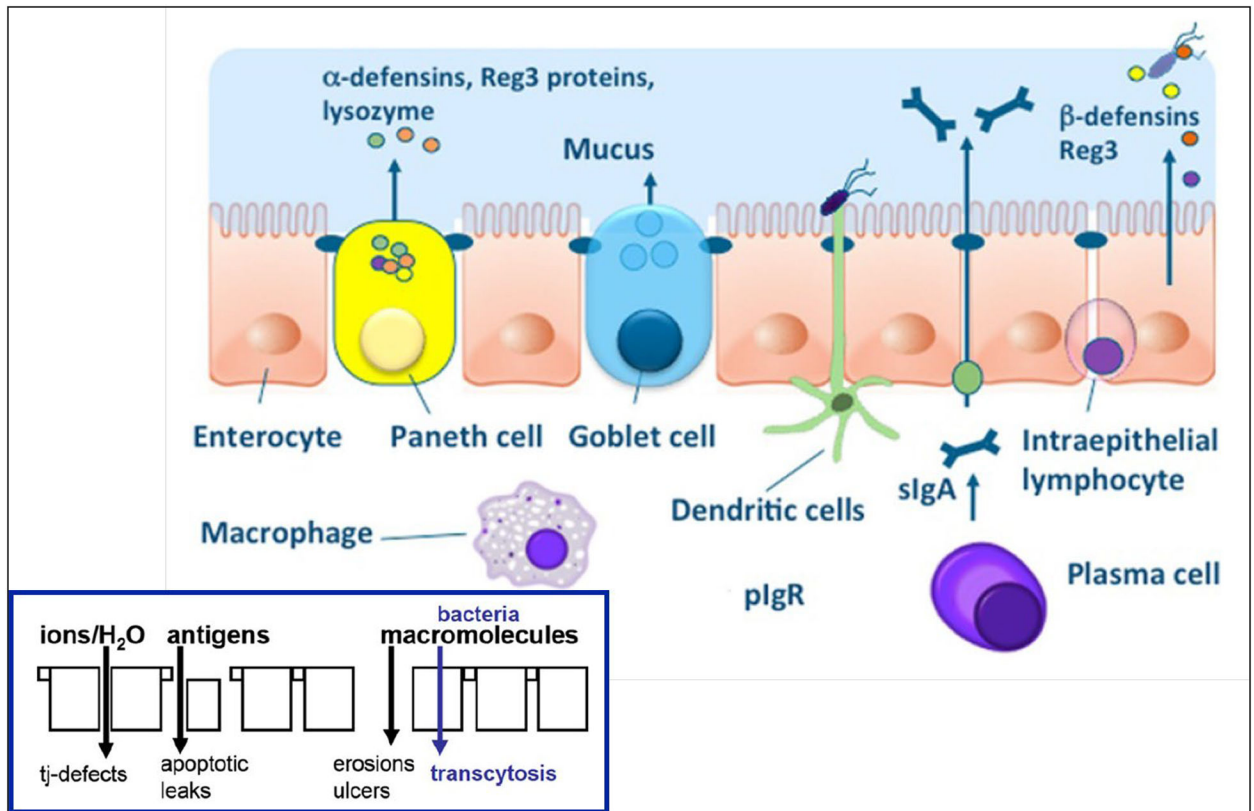


Figure 2.

Intestinal barrier and its dysfunctions. The intestinal barrier includes the mucus layer preventing bacterial adhesion by secretion of chemicals such as α -defensins and IgA secretion, epithelial cells, connected at the tight junctions (tj) by junctional complexes, having the ability to transport luminal content and react to noxious stimuli by secretion of chloride and antimicrobial peptides, and the lamina propria innate and acquired immunity cells secreting Ig and cytokines. Intestinal permeability measurements are determined by the marker molecules used for measurement, since the type of molecules that pass the intestinal barrier depends on the type of lesion. Reproduced from ref. 18.

Table 1.

Molecular sizes of the sugar and other probe molecules

Probe molecule	Mol. weight, Da	Mol. diameter, Å, reported	Mol. diameter, Å, estimated*
Rhamnose	164	8.2	6.9
Mannitol	182	6.7	7.2
Lactulose	342	9.5	9.7
Cellobiose	342	10.5	9.7
Sucralose	398	NA	10.4
Cr-EDTA	340	10.5	9.6
Dextran 4 kDa	4,000	NA	30.0

* calculated based on the formula: $radius = 0.33 * (MM^{0.46})$, where MM=molecular mass

Å= ångström; Cr-EDTA=chromium complexed with ethylene diamine tetracetic acid, Da=dalton

Summary of the literature on conditions that result in stress to the intestinal barrier, with documented effects on intestinal permeability or biochemical evidence of mucosal damage, and restoration with a dietary, non-pharmacological intervention

Table 2.

Barrier stressor; Clinical scenario	Specific Study	Effects on Barrier Function		Dietary Intervention and Its Effects	Reference #
		Intestinal permeability	Mucosal Damage		
Endurance exercise Marathon runners with fecal occult blood or bloody diarrhea	Biking challenge	Urine iohexol (MW 821Da) ↑	Serum I-FABP ↑, zonulin ↓		70
	Running challenge	LRR ↑ and correlated with core temperature (e.g. >39°C)	ND		71
	Biking challenge	LRR ↑	Serum I-FABP ↑	Citrulline (vs alanine) reversed ↑ serum I-FABP and gastric hypoperfusion without effect on LRR	72
	Biking challenge	ND	Serum I-FABP ↑	Sucrose (vs. nitrate) reversed ↑ serum I-FABP no gastric hypoperfusion	73
	7 runners, 2 boxers, 3 rugby	LRR ↑	ND	Nutriceutical colostrum vs. placebo reduced LRR ↑ and apoptosis of cell lines in vitro	74
	Diverse NSAIDs e.g. indomethacin	⁵¹ CrEDTA, saccharides	ND		75-77
	Indomethacin	LRR ↑	ND	Zinc carnosine (vs. placebo) reduced LRR and increased HT29 cell proliferation (vs. ZnSO4)	78
Pregnancy with or without obesity	Aspirin	increased colon permeability; urine SLR and sucralose	ND	Bifidobacterium BB-12 and adolscentsis (IVS-1) and galacto-oligosaccharide prebiotic reduce colon permeability	79
		Increased serum zonulin and increased serum LPS, associated with metabolic risk markers	ND		80
		High serum zonulin	ND	n-3 PUFAs, fiber, and a range of vitamins and minerals; reduced high serum zonulin; greater richness of gut microbiota	81
	No effects on serum zonulin or LPS		ND	Probiotics and/or LC-PUFA	82

Da=dalton; I-FABP=intestinal fatty-acid binding protein; LC-PUFA=long chain polyunsaturated fatty acids; LPS=lipopolysaccharide; LRR=lactulose-rhamnose ratio; ND=not determined; NSAIDs=nonsteroidal anti-inflammatory drugs; PUFA=polyunsaturated fatty acids; SLR=sucralose to lactulose ratio

Table 3.

Summary of diseases or disorders with increased intestinal permeability and altered microbiota. In each category, it is infrequent for the altered barrier dysfunction and microbiota to be documented in the same human study.

Condition	Small Intestinal or Colonic Barrier Function		Microbiota Changes	Other Effects	Effects of Treatment
	IP probe molecules or epithelial damage	Serum biomarkers			
Aging	No difference in LMR or most TJ protein expression, but increased claudin 2 expression and decreased transepithelial resistance in ileal biopsies ex-vivo[83]	↑ zonulin[84]	↓ <i>Firmicutes</i> , <i>Bifidobacteria</i> , <i>Faecalibacterium prausnitzii</i> ↑ <i>Bacteroidetes</i> , <i>Clostridia</i> and <i> facultative anaerobes</i> [85]		
Food allergy	↑ LMR 3 fold vs health[86] ↑ LMR 38% in children with food allergy[87]			Postulated mast cell and IgE-mediated increase in inflammatory cytokines[88]	Increased LMR in children with food allergy despite dietary exclusion[87]
Eosinophilic esophagitis	Increased small bowel IP based on lactulose absorption[89] but not LMR in adults[89, 90] or in children;[91] ex-vivo assessment of duodenal mucosal integrity was normal[90]		Esophageal microbiome: increased hemophilus[92] or <i>Neisseria</i> and <i>Corynebacterium</i> in active EoE[93]	Bacterial load and TLR1, TLR2, TLR4, and TLR9 were overexpressed and mucin genes under-expressed on biopsies with active EoE[94]	No effect of elemental diet on duodenal mucosa or LMR or tight junction protein expression;[90] No effect of exclusion diets on esophageal microbiome[93]
Liver Diseases					
NAFLD/ NASH	↑ LMR or ⁵¹ Ci-EDTA in 39% of 139 pts with NAFLD (SRMA 5 studies)[62]	↑ LPS in 42% of NASH;[95] ↑ LPS in NAFLD associated with SIBO[96]	↑ SIBO (37.5% in pts with NAFLD, especially gram -ve bacteria and <i>E.coli</i>)[96] Review documents show diverse microbiota changes (variable in different studies)[97]	Increased endogenous ethanol production by gut bacteria in NAFLD[61]	
Cirrhosis			Significant microbiota change in liver cirrhosis[98]		Reduced cirrhosis severity with <i>Lactobacillus</i> and VSL#3 probiotics[64]
Sclerosing cholangitis	LRR normal [83% (19/22) with quiescent IBD][99]	Higher serum I-FABP associated with IgA antibodies against F-actin[100]	1/22 had SIBO (<i>Enterobacter</i>);[94] Enhanced mucosal immune response to various microbial antigens associated with IgA antibodies against F-actin[99]	IgA antibodies against F-actin, independent predictor of poor disease outcome [100]	
TPN or enteral deprivation	↑ FITC-Dextran I.P ex-vivo and ↓ ZO-1, E-cadherin, and claudin-4 in unfed segments in pediatric patients:[101] ↓ ZO-1 and villus height in mice[102]		Wide variability in microbial diversity in patients with small bowel resections;[103] Patients with short bowel on TPN have “lactobacteria” enriched in the <i>Lactobacillus/Leuconostoc</i> group, depleted in anaerobic micro-organisms (especially <i>Clostridium</i> and <i>Bacteroides</i>)[104]	In TPN-liver disease, microbes or LPS reaching liver and activating Kupffer cells;[105] <i>Lactobacota</i> fermentation leads to increased risk of d-encephalopathy[104]	Successful use of fecal microbial transplant for the treatment of recurrent D-lactic acidosis[106]
Neurological Diseases					

Condition	Small Intestinal or Colonic Barrier Function		Microbiota Changes	Other Effects	Effects of Treatment
	IP probe molecules or epithelial damage	Serum biomarkers			
<i>Alzheimer</i>			Differences in <i>Bacteroides</i> , <i>Faecalibacterium</i> , <i>Anaerostipes</i> , <i>Prevotella</i> , <i>Escherichia</i> , and <i>Lachnospira</i> at genus level and decreased <i>Firmicutes</i> / <i>Bacteroidetes</i> ratio at phylum level[107]		Only 2/6 trials of omega-3 FA showed any benefit on cognitive decline, typically in those with mild cognitive impairment[108]
<i>Parkinson</i>	Down-regulation of occludin not ZO-1 in colonic mucosa; however, flux of sulfonic acid and horseradish peroxidase not abnormal with or without Lewy bodies:[109] LMR normal, but ↑ 24h urinary sucralose (marker of total intestinal permeability)[110]	Lower plasma levels of LPS binding protein, an indirect measure of systemic endotoxin exposure[110]	Significantly more intense staining of <i>E. coli</i> in epithelium and lamina propria of sigmoid mucosa:[110] Reduced butyrate-producing bacteria from the genera <i>Blautia</i> , <i>Coprococcus</i> and <i>Roseburia</i> ; putative "proinflammatory" <i>Proteobacteria</i> of the genus <i>Ralstonia</i> significantly more abundant in mucosa of Parkinson's patients[111]	Correlation of increased intestinal permeability in Parkinson disease with intestinal alpha-synuclein:[109] Relative abundance of <i>Enterobacteriaceae</i> positively associated with severity of postural instability and gait difficulty[112]	
<i>ALS</i>	↑ LPS in most severe amyotrophic lateral sclerosis[113]		Low diversity of the microbiome compared to healthy cohorts; low <i>Ruminococcus</i> spp. in 3/5 patients with low <i>Firmicutes</i> / <i>Bacteroidetes</i> (F/B) ratio[114]	Decreased levels of butyrate-producing bacteria; decreased levels of micro-organisms of the genera <i>Oscillibacter</i> , <i>Anaerostipes</i> , and <i>Lachnospira</i> :[115] 3/5 patients had elevated inflammatory markers in stool[114]	
<i>Psychiatric diseases</i>	Plasma levels of LPS, zonulin and FABP2 were each significantly elevated in depression/anxiety patients compared to non-depressed or anxious controls.[116]		A review documents extensive literature on cross-sectional and longitudinal studies documenting association between stool microbiota and anxiety and depression.[117] A review documents studies of the microbiome and microbial translocation in patients with schizophrenia and bipolar disorder.[118]	Elevated serum IgM and IgA against LPS in depression:[119] Psychological stress increases pro-inflammatory cytokines (extensive literature reviewed in ref. 120).	Probiotics reduce depression scores in 6 randomized, placebo-controlled trials (reviewed in ref. 121).

EoE=eosinophilic esophagitis; FABP= fatty-acid binding protein; FITC=fluorescein isothiocyanate; IBD=inflammatory bowel disease; I-FABP=intestinal fatty-acid binding protein; IgA=immunoglobulin A; IP=intestinal permeability; LMR=lactulose mannitol excretion ratio; LPS=lipopolysaccharide; LRR=lactulose-rhamnose ratio; NAFLD=non-alcoholic fatty liver disease; NASH=non-alcoholic steatohepatitis; SIBO=small intestinal bacterial overgrowth; TJ=tight junction; TLR=oll-like receptor; TPN=total parenteral nutrition; ZO=zonula occludens

Table 4.

Examples from the literature of *in vivo* human studies showing alterations in intestinal permeability as a result of gut-directed therapy with nutrient, supplement or fiber

Therapy	Patients with	Comments	Reference #
Glutamine i.v.	Nutritional depletion pre-surgery	Patients with increased intestinal permeability did not improve after glutamine-enriched total parenteral nutrition	122
Enteral glutamine	50–80% burns	Urinary LMR in enteral glutamine group lower than standard enteral formula	123
Enteral glutamine	30–75% burns	Plasma DAO activity and urinary LMR in enteral glutamine group were lower than in untreated burn group	124
Enteral glutamine	Crohn's disease patients	Glutamine and active control (whey) both reduced LMR	125
Oral glutamine	Post-infectious IBS	Elevated urinary LMR was normalised in the glutamine but not the control group	65
Galactooligo-saccharides	Obesity	Post-aspirin sucralose:lactulose ratios and sucralose excretion reduced indicating improvements in colonic permeability	79
Inulin- enriched pasta	Healthy young volunteers	Reduced serum zonulin and urine LMR but not mannitol excretion	126
Psyllium	Children with IBS	No effect on sucrose and sucralose recovery, and LMR	127
Fiber	NAFLD patients undergoing diet Rx for obesity	Increasing nutritional fiber from 19 to 29g/day reduced serum ZO levels, and improved hepatic steatosis	128
Rhubarb	Day 3 post-burns	Plasma DAO activity in rhubarb-treated group were lower than in controls	129
Gelling complex of tara gum + exopolysaccharides of <i>Strep. Thermophilus</i> ST10	Healthy participants	Reduced LMR and reduced sucralose concentration suggesting improvement in both small intestinal and colonic permeability	130
Ascorbic acid	Healthy female participants	Excretion of lactulose over 6-hr augmented after consumption of either aspirin or ascorbic acid compared with that after consumption of placebo	131
Fermented and amylase-digested weaning foods	Tanzanian children aged 6–25 months with acute diarrhea	Reduction in L/M ratio between admission and day 3 of hospitalization was significantly greater in the fermented and amylase-digested weaning group (89%) than the conventional or high-energy density amylase digested porridge groups (44% and 75%, respectively)	132
Common beans	Rural children in second year of life (Malawi)	Additional beans in diet reduced lactulose excretion without effect on LMR	133
Micronutrient-fortified complementary/replacement food	Zambian infants from 6 to 18 months old	LMR (adjusted for baseline urinary L:M ratio, socio-economic status, mother's education, season of birth and baseline stunting, and stratified by maternal antenatal HIV status, child's sex, concurrent breast-feeding status and anemia at baseline) worse in group with fortified diet especially among boys and, among the infants of HIV-negative mothers	134

DAO=diamine oxidase; LMR=lactulose mannitol excretion ratio