Inflammatory Intestinal Diseases

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Increased Intestinal Permeability and Decreased Barrier Function: Does It Really Influence the Risk of Inflammation?

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Key Words

Intestinal permeability · Endotoxemia · Inflammatory bowel disease · Irritable bowel syndrome · Liver disease · Acute pancreatitis · Chronic kidney disease · Chronic heart failure · Depression

Abstract

Background: Increased intestinal permeability due to barrier dysfunction is supposed to cause microbial translocation which may induce low-grade inflammation in various diseases. However, this series of events has not been comprehensively evaluated yet. Summary: Intestinal epithelial barrier dysfunction and increased permeability have been described in patients with inflammatory bowel disease (IBD), irritable bowel syndrome (IBS), alcoholic liver disease, nonalcoholic steatohepatitis (NASH), liver cirrhosis, acute pancreatitis, primary biliary cholangitis (PBC), type 1 and type 2 diabetes, chronic kidney disease, chronic heart failure (CHF), depression, and other diseases. Most clinical reports used either permeability assays of challenge tests or measurement of circulating bacterial markers like endotoxin for assessment of 'the leaky gut'. The intestinal permeability assessed by the challenge tests has often been related to the changes of tight junction proteins in the epithelium or circulating endotoxin levels. In patients with IBD, alcoholic liver disease, NASH, liver cirrhosis, PBC, obstructive jaundice, severe acute pancreatitis, and CHF, endotoxemia and proinflammatory

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E-Mail karger@karger.com www.karger.com/iid cytokinemia have been found in addition to increased permeability. In the serum of patients with IBS and depression, antiflagellin antibodies and antilipid A antibodies were detected, respectively, together with increased permeability and proinflammatory cytokinemia. The site of infection, which is localized to the intestine in IBD and IBS, includes various extraintestinal organs in other diseases. The relation of gut dysbiosis to intestinal barrier dysfunction has gradually been clarified. Key Messages: Although no direct causeand-effect relationship has been confirmed, all clinical and experimental data suggest the importance of intestinal hyperpermeability in the inflammatory changes of various diseases. Increased intestinal permeability is a new target for disease prevention and therapy. Considering the close relationship of 'the leaky gut' and gut dysbiosis to the major diseases, we can conclude that meticulous dietetic and probiotic approaches to recover healthy microbiota have the potential to make a breakthrough in the management of these diseases tomorrow. © 2016 S. Karger AG, Basel

Introduction

Intestinal barrier prevents the entry of pathogenic microorganisms and toxic luminal substances while regulating the absorption of nutrients, electrolytes and water from the lumen into the circulation [1]. These functions

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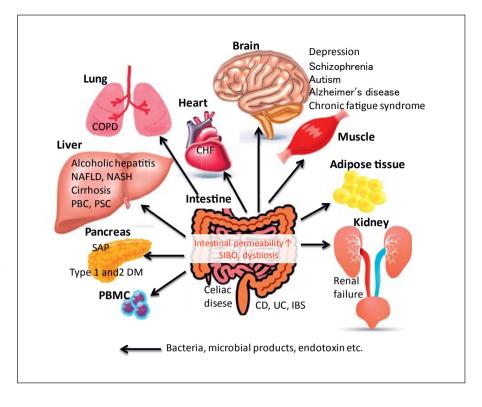


Fig. 1. The diseases in which increased intestinal permeability has been reported in the literature. Intestinal epithelial barrier dysfunction and increased permeability allow the translocation of bacteria and microbial products, which may induce inflammatory changes in the target organs. The 'leaky gut hypothesis' seems to be a reasonable explanation of the pathophysiological background of various diseases. PSC = Primary sclerosing cholangitis.

are preserved by a complex multilayer system, consisting of an external physical barrier and an inner functional immunological barrier [2]. From a structural perspective, this multilayer system includes a mucus layer and a monolayer of epithelial cells interconnected by tight junctions (TJs). Intestinal permeability is a functional feature of the intestinal barrier measurable by analyzing flux rates of inert molecules across the intestinal wall, which was precisely defined by the consensus in an expert panel in Frankfurt/Germany in June 2012 [2]. An intact intestinal barrier prevents the permeation of antigens, endotoxins, pathogens, and other proinflammatory substances into the human body, whereas intestinal disintegrity allows their entry, which may trigger local or systemic inflammation and disease [3]. Assessment of intestinal barrier function and permeability in humans is currently possible by using intestinal permeability assays, and by the assessment of biomarkers of epithelial integrity such as soluble adhesion molecules, other biomarkers of immunity or inflammation, or bacterial markers like circulating endotoxin. Most clinical reports used either permeability assays of challenge tests or measurement of circulating bacterial markers like endotoxin [2]. It should be noted that the mechanisms determining the flux of challenged substance and the translocation of bacteria or their products are different, which presents a

limitation to the intestinal permeability assays. Therefore, the detection of gut-derived microbial products in the circulation is considered as a definite evidence of increased intestinal permeability. In addition, histological approaches and scanning electron microscopy analyses of the intestinal mucosa have been used in experimental settings [2]. The minute mechanism of this gastrointestinal barrier and variable evaluation methods of gut permeability have been described in detail in a previous review [2].

Intestinal epithelial barrier dysfunction and increased permeability have been described in many human diseases, including inflammatory bowel disease (IBD), irritable bowel syndrome (IBS), alcoholic liver disease, nonalcoholic fatty liver and steatohepatitis, liver cirrhosis, severe acute pancreatitis (SAP), primary biliary cholangitis (PBC), type 1 and type 2 diabetes, depression, and more, as presented in figure 1. The 'leaky gut hypothesis' explains that the intestinal barrier dysfunction induces the chronic low-grade inflammation in various target organs by virtue of microbial products. This review discusses clinical and experimental evidence linking gut permeability and inflammation in various diseases to find out whether increased intestinal permeability and decreased barrier function really influence the risk of inflammation. Clinical evidence on the permeability tests, intestinal ep-

Diseases	Permeability test	Intestinal epithelial changes	Microbial products in the blood	Inflammatory changes in various organs
IBD	IP↑ in CD (L/M test) [4] IP↑ in CD (active > inactive) (L/M test) [6] IP↑ in inactive UC (L/M test) [7] IP↑ in 50% CD and 31% UC (iohexol test) [8]	Expression of PrP(c) ↓ in the colonic epithelia from CD and UC patients [9]	Endotoxemia (+) in 28~88% active UC and 48~94% active CD [10, 11] Endotoxemia (+) correlated with disease activity, disease extent and plasma TNF [10]	Serum LBP and sCD141 correlated with disease activity and proinflammatory cytokines [11]
IBS	IP† especially in PI-IBS and D-IBS [15] IP† associated with IBS interference with activities and work, anxiety and depression (L/M test) [16]	ZO-1 expression ↓ correlated with increased paracellular IP [17] ZO-1 and occludin expression↓ [18]	Anti-flagellin antibodies (+) in ca. 30% IBS (mostly in PI-IBS) [23]	Serum TNF-α, IL-1β, IL-6† especially in D-IBS [19] LPS-induced TNF-α production by PBMCs† associated with anxiety [19]
Alcoholic liver disease	IP↑ (⁵¹ Cr-EDTA test) [25] Increased IP was associated with dysbiosis (⁵¹ Cr-EDTA test) [27] IP↑↑ (L/M test) [26]	ZO-1 and occludin in the duodenum↓ [32]	Endotoxemia (+) correlated with ID (PEG Mr 4000 method) [29]	Serum IL-8↑, granulocytosis correlated with endotoxemia in alcoholic hepatitis [37] Plasma TNF-α↑ correlated with endotoxemia in alcoholic cirrhosis [38]
NAFLD	IP† (⁵¹ Cr-EDTA test) [44] IP† (L/M test) in child steatohepatitis correlated with the severity of steatohepatitis [45]	ZO-1 in the villi↓ Nuclear ZO-1 in the crypt cell↓ correlated with IP [44]	Endotoxemia (+) related with SIBO [47]	Liver TLR4 protein expression [↑] in NASH [23] Liver TLR4 mRNA and serum TNF-α [↑] in NAFLD [47] Liver TLR4 mRNA [↑] (NASH > NAFL) [48]
Liver cirrhosis	IP↑ (L/M test, ⁵¹ Cr-EDTA test, PEG test) [54]	Occludin and claudin-1↓ in the duodenum negatively correlated with serum endotoxin levels [55]	Endotoxemia (+) related with Child-Pugh grades, renal disturbance, GI bleeding, and infection [28]	TNF, IL-2, IL-4, IL-6, IL-8 and IFN-γ [↑] especially in those with massive ascites [59]
PBC	IP† (L/M test) [63]	IgA secretion ↓ [64]	Endotoxemia (+) [65] IgM antibodies against LTA (+) [67]	Liver TLR4 and CD14 protein↑ [65]
Obstructive jaundice	IP↑ (L/M test)[69] IP↑ (PEG test) [70]	Occludin, claudin-1, -7 proteins↓ in the duodenum [75]	Endotoxemia (+) [73]	TNF, IL-6, IL-8† [72]
SAP	IP↑ (L/M test) [77, 80] IP↑ (PEG test) [79, 111]	TJPs in the colon↓ occludin and ZO-1↓ in patients with BT[82]	Endotoxemia (+) correlated with IP [77, 79]	Inflammatory cytokines† [81] Serum IL-6, TNF-α, CRP, CT severity index correlated with IP [80]

Table 1. Clinical evidence on the permeability tests, intestinal epithelial changes, circulating microbial products and inflammatory changes of the target organs in various diseases

ithelial changes, circulating microbial products and inflammatory changes of the target organs in various diseases is summarized in table 1.

Inflammatory Bowel Diseases

A critical etiological factor in IBD is that the mucus layer becomes more permeable to bacteria and bacterial products [2]. Patients with Crohn's disease (CD) exhibit marked increases in intestinal permeability assessed by the lactulose-mannitol (L/M) test [4]. Paracellular intestinal permeability estimated by the Ussing chamber method was increased even in patients with inactive IBD [5], although the maximum blood lactulose concentrations were higher in patients with active diseases [6]. A significant correlation was noted between the maximum blood lactulose concentrations and serum CRP levels [6]. Increased intestinal permeability was also detected in patients with ulcerative colitis (UC) in remission [7]. Intestinal permeability estimated by the serum level of iohexol, a radiographic contrast media, was increased in 50% of CD and in 31% of UC patients, and was also related to the disease activity judged by the endoscopy [8]. Disrupted barrier function comprises alterations in epithelial TJ, i.e. a reduced number of horizontal TJ strands and an altered TJ protein (TJP) expression, and subcellular distribution with decreased expression of cellular prion protein in colonic epithelia of CD and UC patients [2, 9]. These barrier defects are attributed to the enhanced activity of proinflammatory cytokines like TNF- α , INF- γ , IL-1 β and IL-13, which are highly expressed in the chronically inflamed intestine [2].

Systemic endotoxemia was reported to be present in 28~88% of patients with UC and 48~94% with CD during clinical relapse [10, 11]. Endotoxemia, its correlation with disease activity, disease extent, and circulating TNF support a pathogenic role of endotoxin in IBD [10]. Serum endotoxin, lipopolysaccharide (LPS)-binding protein (LBP), and soluble CD14 levels were correlated with disease activity and paralleled to a rise in proinflammatory cytokines, suggesting a contribution of bacterial products to the inflammatory cascade in IBD patients [11].

As for the changes in the gut microbiome, a decreased abundance of butyrate-producing *Faecalibacterium prausnitzii* (Firmicutes phylum, Clostridiales order) may lead butyrate deficiency and intestinal inflammation in IBD [12]. Moreover, an increase in the number of sulfatereducing bacteria, which produce toxic hydrogen sulfide, may provoke epithelial cell injury and inflammation in UC [13]. Metabolome analysis further suggests that bile acid dysregulation due to gut dysbiosis may result in increased intestinal permeability and inflammation in patients with IBD [14].

Irritable Bowel Syndrome

Increased small bowel and colonic permeability has been noted in both adult and pediatric patients, primarily with postinfectious IBS (PI-IBS) and diarrhea predominant IBS (D-IBS) [15]. The L/M ratio is significantly correlated with IBS interference with activities and work, anxiety, and depression [16]. Increased paracellular permeability was associated with the expression and distribution of TJPs, lower levels of the protein zonula occludens (ZO)-1 and occludin in intestinal tissue [17, 18].

Concerning inflammatory changes, Liebregts et al. [19] studied the cytokine production in peripheral blood mononuclear cells (PBMCs) and reported that IBS, especially D-IBS patients, showed high baseline serum TNF- α , IL-1 β , and IL-6 levels. PBMCs in patients with D-IBS further showed enhanced TNF- α release by endotoxin (LPS) stimulation, which was correlated with anxiety [19]. Mast cells, which increase in the colon of IBS patients [20], degranulate to release inflammatory and immune mediators promoting the recruitment of other inflammatory cells [21].

Expression of Toll-like receptor 4 (TLR4; recognizes bacterial LPS) and TLR5 (recognizes flagellin, a common bacterial antigen) is increased in colons of IBS patients compared with controls [22]. Antiflagellin antibodies were found in almost 30% of IBS (mostly PI-IBS) patients as opposed to only 7% of healthy controls [23], suggesting the importance of the interaction between the intestinal microbiota and the immune system in these IBS patients [22]. Several susceptibility genes for IBS involved in the innate immunity, recognition of bacteria, or maintenance of intestinal barrier integrity have been identified [24].

Alcoholic Liver Disease

Noncirrhotic alcoholic patients abstaining from alcohol for less than 4 days almost invariably showed increased intestinal permeability by way of a ⁵¹Cr-EDTA absorption test [25]. In many patients this abnormality persisted for up to 2 weeks after cessation of drinking [25]. Alcoholics with chronic liver disease showed a markedly increased intestinal permeability by the L/M test, whereas the increase in intestinal permeability was slight in alcoholics with no liver disease [26]. Only patients with increased intestinal permeability had an altered fecal microbiota composition: a drastic decrease in the abundance of Ruminococcus, Faecalibacterium, Subdoligranulum, Oscillibacter, and Anaerofilum belonging to the Ruminococcaceae family [27]. The total amount of bacteria and those belonging to the Ruminococcaceae family, especially for Faecalibacterium prausnitzii was negatively correlated with intestinal permeability, while the genera Dorea and Blautia were positively correlated with intestinal permeability [27].

The passage of viable bacteria from the intestinal lumen through the mesenteric lymph nodes and other sites are defined as bacterial translocation (BT). The concept of BT was later broadened to microbial products or their fragments, such as endotoxin, peptidoglycan, lipopeptides, and bacterial DNA [28]. The liver receives portal blood containing these microbial products and acts as the initial site of their filtration and detoxication [28]. Parlesak et al. [29] found a significant correlation between the plasma endotoxin concentrations and the intestinal permeability of polyethylene glycol (PEG) Mr 4000. They discussed PEG as an appropriate probe for the assessment of gut-derived endotoxin translocation on the basis of its homogeneous chemical properties, appropriately adaptable molecular mass and linear, chain-like shape mimicking the structure of endotoxin [29]. Alcohol is known to disrupt gastrointestinal epithelial barrier integrity [30], resulting in the translocation of potentially harmful bacteria and their products such as endotoxins [31, 32]. Alcohol and its metabolites, acetaldehyde and fatty acid ethyl esters, may contribute to the disruption of TJs, mainly through nitric oxide-mediated oxidative tissue damage and alterations in the cytoskeleton, but also through direct cell damage [33-35].

There is strong evidence to support the concept that gut-derived endotoxin as a marker of BT plays a central role in the initiation and progression of alcohol-induced liver injury [36]. Once endotoxin reaches various organs, it powerfully stimulates TLR4 both in hepatic macrophages (Kupffer cells - KCs) and extrahepatic macrophages, which activate downstream signaling pathways responsible for overproduction of proinflammatory cytokines such as TNF-a, IL-6, and IL-8. Plasma endotoxin levels were increased with the progression of alcoholic liver injury and reached the maximal level in patients with alcoholic cirrhosis and severe alcoholic hepatitis who showed marked hypercytokinemia [37]. In these patients, plasma endotoxin levels were positively correlated with serum IL-8 levels and peripheral neutrophil counts [37]. In another study on alcoholic cirrhosis, plasma endotoxin levels were strongly correlated with the plasma levels of TNF-a and its receptors -p55 and -p75 [38]. Alcohol induces the LBP and TLR4, and increases responsiveness to gut-derived endotoxin. The binding of LPS to CD14/ TLR4 on KCs stimulates the production of cytokines and oxidants, which leads to hepatic stellate cell activation and collagen synthesis in the liver of patients with alcoholic steatohepatitis [28, 39].

Lower median abundances of Bacteroidetes and higher ones of Proteobacteria in the colonic microbiome were proved to be correlated with high levels of serum endotoxin in alcoholics [40]. Endotoxemia and hepatic inflammation are considered as consequences of the expansion of the Gram-negative bacteria from the Proteobacteria phylum, which occur in response to chronic ethanol consumption [41]. A decrease in commensal probiotics could contribute to the loss of protective TJ barrier [42].

Nonalcoholic Fatty Liver Diseases

Nonalcoholic fatty liver disease (NAFLD) is the hepatic manifestation of the metabolic syndrome. It includes a spectrum of pathological changes ranging from the simple accumulation of fat (NAFL) in the liver through nonalcoholic steatohepatitis (NASH) to fibrosis and cirrhosis [43]. NAFLD patients present increased gut permeability characterized by disruption of the intercellular TJs with decreased TJP ZO-1 expression, which is likely to allow translocations of bacteria and their products [44]. Intestinal permeability is increased in children with NAFLD, and correlates with the severity of steatohepatitis [45]. On the other hand, NASH patients were found to have endotoxemia and overexpression of TLR4 protein in the liver [46, 47] associated with proinflammatory cytokine release and systemic inflammation. Hepatic TLR4 mRNA expression and plasma endotoxin levels were proved to be increased in NASH patients compared with NAFL patients [48]. Induction of an intestinal inflammation by dextran sulfate sodium in experimental NASH promotes LPS translocation, hepatic inflammation, and fibrogenesis [49]. Our group reported enhanced a-SMA expression (suggesting hepatic stellate cell activation), elevated liver LBP mRNA levels, increased intestinal permeability, and decreased intestinal TJP expression in the rat NASH model fed a choline-deficient L-amino-aciddefined diet [50]. We also proved that oral administration of poorly absorbable antibiotics improved all of these intestinal and liver events and inhibited the progression of liver fibrosis [50].

Small intestinal bacterial overgrowth (SIBO) relevant in NASH patients is also associated with enhanced hepatic expression of TLR4 and release of IL-8 [51]. In the experimental condition of high-fat diet, not only bacterial products but also complete living bacteria can be translocated from the intestinal lumen towards adipose tissues [52]. Adipose tissues in NASH patients are infiltrated by a large number of macrophages, and this recruitment is linked to systemic inflammation and insulin resistance [53].

Liver Cirrhosis

Many authors have reported that patients with liver cirrhosis revealed intestinal hyperpermeability [54]. Structural and functional changes in the intestinal mucosa that increase intestinal permeability have been regarded as an important pathogenetic factor for several

complications of liver cirrhosis including bacterial infections. Reduced expression of duodenal occludin and claudin-1 has been found especially in patients with decompensated cirrhosis. Negative correlation was proved between these expressions and serum endotoxin levels [55]. Bacterial products, such as endotoxin, or bacterial DNA can translocate to extraintestinal sites and promote an immunological response similar to that produced by viable bacteria. Pathological BT is a contributing factor in the development of infections and other complications in cirrhosis, by exerting a profound inflammatory state and exacerbating the hemodynamic derangement [54, 56]. SIBO in cirrhosis showed a high correlation with the presence of bacterial DNA fragments in peripheral blood, suggesting that SIBO could be a major risk factor for BT, especially in ascitic patients [57].

Endotoxin binds to TLR4 with the co-receptors CD14 and MD-2. TLR2 heterodimerizes with TLR1 or TLR6 to recognize lipoprotein and peptidoglycan derived from Gram-positive bacteria. Bacterial flagellin is recognized by TLR5. Intracellular TLR3 and TLR9 are activated by microbe-derived nucleic acids including double-stranded RNA and CpG motif containing unmethylated DNA, respectively [43, 58]. Translocated microbial products thus activate KCs through TLRs, which activate innate immune responses including cytokine production [39]. The circulating levels of inflammatory cytokines TNF, IL-2, IL-4, IL-6, IL-8, and IFN-γ were especially high in cirrhotic patients with massive ascites [59]. Depressed elimination of endotoxin by KCs causes spillover endotoxemia. Decreased endotoxin inactivation in the blood is considered to enhance the processing of endotoxin by extrahepatic macrophages which secrete larger amounts of TNF than KCs [28, 60, 61]. The excessive cytokine response to endotoxin by splenic and pulmonary alveolar macrophages may be important in the pathogenesis of acute respiratory distress syndrome and multiple organ failure in advanced liver cirrhosis [28, 61]. Hepatic encephalopathy is also closely related to inflammatory reaction attributable to leaky gut and endotoxemia [28].

Primary Biliary Cholangitis

PBC is a chronic inflammatory cholestatic disease of unknown etiology that affects small and medium intrahepatic bile ducts [62]. The permeability of both the stomach and small intestine was increased in patients with PBC estimated by the sugar test and the L/M test [63]. The majority (66.6%) of patients with abnormal permeability did not have evidence of portal hypertension, and some patients with very early-stage PBC had increased permeability [63]. So far, no structural changes in the intestinal mucosa have been reported, although an IgA secretion defect in the intestinal epithelium was proposed [64]. Significant endotoxemia was found in patients with early PBC, and enhanced immunohistochemical expression of TLR4 and CD14 was found in the liver tissues of PBC patients [65]. TLR4 expression is significantly elevated in biliary epithelial cells and periportal hepatocytes of PBC patients [66]. PBC sera were also positive for IgM antibodies against lipoteichoic acid (LTA), the Gram-positive bacterial cell wall component [67]. LTA was localized around the sites of chronic nonsuppurative destructive cholangitis in the portal area in stage 1-2 PBC and was detected around the sites of ductular proliferation at the periphery of portal tracts in stage 3-4 PBC [67].

Obstructive Jaundice

Surgery for relief of obstructive jaundice has a risk of sepsis and renal dysfunction [68]. Increased intestinal permeability was confirmed in these patients both by the L/M test [69] and the polyethylene glycol test [70], which may be related to frequently observed BT [71]. Malignant obstructive jaundice caused increased blood concentrations of endotoxin and inflammatory cytokines [72, 73]. Portal endotoxemia resulted in an increase in TNF-a, IL-6, and IL-10 in the bile duct-ligated rats [74]. Altered TJPs, decreased expression of occludin, claudin-1, and -7, were found in the duodenal epithelium of patients with obstructive jaundice [75]. Bile duct ligation in animals also resulted in the regional loss of occludin expression in the intestinal epithelium, which was improved by bile feeding [76]. Bile acids inhibit the growth of Bacteroides, Clostridia, Lactobacillus and Streptococcus, and the absence of bile acids results in a disturbed intestinal bacterial balance with overgrowth of Gram-negative bacteria [68].

Acute Pancreatitis

Increased intestinal permeability was confirmed by several methods in patients with acute pancreatitis [77–80]. The intestinal permeability was correlated with plasma endotoxin [77, 79, 80], serum TNF- α [77, 80], IL-6, CRP, and the severity index estimated by computed tomography [80]. The urinary concentration of intestinal

fatty acid-binding protein, a sensitive marker of intestinal ischemia, correlated positively with intestinal permeability, which suggests that splanchnic hypoperfusion induces the loss of intestinal mucosal integrity [78]. Overwhelming systemic production of inflammatory mediators and early organ failure are characteristics of SAP [81]. The expression of TJPs in the colonic mucosal tissue was decreased in patients with SAP, 62% of which showed BT (positive bacterial DNA in the peripheral blood) [82]. Further, patients with BT showed a lower level of occludin and ZO-1 expression [82]. The failure of intestinal barrier is associated with translocation of bacteria and inflammatory products through the intestinal wall, which can be responsible for the infection of the necrotic pancreas and systemic inflammatory response [83]. Higher rates of multiple organ failure and infectious complications were observed in patients with SAP and intestinal dysbiosis: the increase in Enterococcus and decrease in Bifidobacterium [84]. Serum IL-6 levels were positively correlated with the abundance of Enterobacteriaceae and Enterococcus and negatively correlated with that of Bifidobacterium, whereas plasma endotoxin was positively correlated with the abundance of Enterococcus, which suggests that the intestinal dysbiosis may be involved in the progression of acute pancreatitis [84].

Chronic Kidney Disease

Increased intestinal permeability in patients with chronic kidney disease (CKD) was reported in the early 1990s [85] after the studies on the intestinal mucosal changes showing shortening of the villi, elongation of the crypts, and infiltration of lamina propria [86, 87]. An in vitro study using TJ-forming human enterocytes revealed that exposure to plasma from patients with end-stage renal disease damages the epithelial TJ and impairs its barrier function [88]. Uremia-induced disruption of intestinal TJ and barrier function is, in part, mediated by urea [89]. This leaky barrier allows the translocation of endotoxin, bacterial DNA, and uremic toxins from the gut [90]. Bacterial DNA from the colon was detected in the mesenteric lymph nodes, liver, spleen, and blood of CKD rats [91]. Circulating endotoxin levels, which increase along the stages of CKD and are highest in patients on hemodialysis or peritoneal dialysis [92, 93], are correlated with serum CRP levels [92-94]. The gut microbiota in patients with end-stage renal disease exhibited significant expansion of bacterial families possessing urease, uricase and indole, and p-cresol-forming enzymes, and contraction of families possessing butyrate-forming enzymes [95]. Given the deleterious effects of indoxyl sulfate, pcresol sulfate, and urea-derived ammonia, and beneficial actions of short chain fatty acid (SCFA) butyrate, these changes in intestinal microbial metabolism may contribute to uremic toxicity and inflammation [95].

Chronic Heart Failure

Patients with chronic heart failure (CHF) showed a 35% increase in small intestinal permeability by the L/M test and a 210% increase in large intestinal permeability by the sucralose test [96]. Increased wall thickness of both small and large intestines and larger amounts of adherent bacteria within mucus were also noted in CHF patients [96]. Increased intestinal permeability was associated with clinical disease severity, venous blood congestion, and serum CRP [97]. Raised plasma endotoxin and cytokine levels were found in patients with CHF during acute edematous exacerbation [98]. Adults with congenital heart disease had elevated levels of inflammatory cytokines and endotoxin, which were related to the functional status [99]. It was also reported that CHF patients with abnormal endotoxin levels had higher concentrations of TNF and sTNF-R1 [100]. The above results all indicate that disturbed intestinal microcirculation and barrier function in CHF seem to induce translocations of bacteria and their products and to trigger cytokine generation, thereby contributing to impaired cardiac function [101]. An analysis of gut microbiota in CHF patients revealed that they had massive quantities of pathogenic bacteria such as Campylobacter, Shigella, Salmonella, Yersinia enterocolitica, and Candida species compared with normal controls, which may be related to increased intestinal permeability, and have intestinal overgrowth of pathogenic bacteria and clinical disease severity [97].

Depression

There is now full evidence that major depression is accompanied by an activation of the inflammatory response system and that proinflammatory cytokines and endotoxin may induce depressive symptoms [102]. The prevalence and median values of serum IgM and IgA against the LPS of Gram-negative enterobacteria, i.e. *Hafnia alvei*, *Pseudomonas aeruginosa*, *Morganella morganii*, *Pseudomonas putida*, *Citrobacter koseri*, and *Klebsiella pneumoniae* were significantly higher in depressed patients

than in controls [103]. Deranged intestinal permeability may underpin the chronic low-grade inflammation observed in depression, and the gut microbiome plays a critical role in regulating intestinal permeability [1]. Bacterial DNA is present in whole serum from depressed patients who also display increased TLR4 expression on PBMC [1]. Patients with major depression exhibit inflammatory responses, including increased expression of proinflammatory cytokines and their receptors and increased levels of acute-phase reactants, chemokines, and soluble adhesion molecules in peripheral blood and cerebrospinal fluid [104]. Peripheral blood gene expression profiles revealed an overexpression of proinflammatory IL-6, IL-8, and type I IFN-induced signaling pathways [104-107]. Further, increased expression of a variety of innate immune genes and proteins, including IL-1 β , IL-6, TNF-a, TLR3, and TLR4, has been found in postmortem brain samples from suicide victims that had depression [108, 109].

Conclusions

We have presented here the clinical and experimental evidence in several diseases in which increased intestinal permeability owing to gut barrier dysfunction may be responsible for inflammatory changes, although we are not convinced that it *really* influences the risk of inflammation. Direct cause-and-effect relationships have not been demonstrated anywhere. There has been no study to compare the grade of inflammation between those with and without increased intestinal permeability. No experimental study has been directed to confirm this hypothesis. Nevertheless, when we have collected references using the keywords 'intestinal permeability', 'gut barrier', 'dysbiosis', 'endotoxin', 'bacterial translocation', 'Tolllike receptor', 'cytokine' and so on, a common story consistent with the thesis has surprisingly appeared concerning diseases all over the body. Although there still remain lot of issues to be confirmed concerning the results and interpretations, this 'leaky gut hypothesis' seems to be a reasonable explanation of the pathophysiological background of these diseases. The hypothesis contains broad and deep clinical implications and suggests the importance of the intestine in human health and disease. The whole story on 'leaky gut' in the diversity of human diseases could not be covered in this short review, but may be presented in an entire book.

Recently, a marked technological progress in the studies of gut microbiota has opened a new research field. Accumulating lines of evidence support the close relationship of gut microbiota and intestinal functions [43]. The 'leaky gut hypothesis' may be coupled with the 'dysbiosis hypothesis' when the host reactions to gut dysbiosis are more clearly defined in association with intestinal changes. It is true that intestinal permeability is a new target for disease prevention and therapy [2]. Although probiotics, prebiotics, antibiotics and their combinations have been extensively tried until now, safe and useful medication to repair 'leaky gut' is still difficult to produce. Anyway, it should be noted that food intake affects the intestinal microbiome composition and intestinal permeability [2]. Remarkable increases in IBD patients and diabetic patients over the past 30 years in Japan may be related to the dramatic changes in the environment, especially the changes in dietary habits: from the traditional Japanese foods to high-fat, high-calorie Western foods [110]. The meaningful advice by a Japanese clinician in 1970s with a word of 'Ishoku-Dogen', which proposes a restricted balanced diet for prevention and treatment of illness, is expected to improve the situation. Further investigations on the relationship of diet to gut microbiome and intestinal functions may support that an ideal traditional diet from the old East may help the 'leaky gut syndrome' of modern people in the West and East. There is a great possibility that meticulous management of gut microbiota and intestinal functions improve general human health by virtue of lifestyle improvement combined with dietary and pharmaceutical approaches.

Disclosure Statement

I have no conflicts of interest to declare that are relevant to the subject of this review paper and any of the statements in it.

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