Role of Probiotics in Human Gut Microbiome-Associated Diseases

Seon-Kyun Kim†, Robin B. Guevarra‡, You-Tae Kim, Joongi Kwon, Hyeri Kim, Jae Hyoung Cho, Hyeun Bum Kim, and Ju-Hoon Lee

1Department of Food Science and Biotechnology, Kyung Hee University, Yongin 17104, Republic of Korea
2Department of Animal Resources Science, Dankook University, Cheonan 31116, Republic of Korea

Probiotics, including bacteria and yeast, are live microorganisms that have demonstrated beneficial effects on human health. Recently, probiotic bacteria are constantly being studied and their applications are also being considered in promising adjuvant treatments for various intestinal diseases. Clinical trials and in vivo experiments have extended our current understanding of the important roles that probiotics play in human gut microbiome-associated diseases. It has been documented through many clinical trials that probiotics could shape the intestinal microbiota leading to potential control of multiple bowel diseases and promotion of overall wellness. In this review, we focused on the relationship between probiotics and the human gut microbiota and its roles in gut microbiome-associated diseases. Here, we also discuss future directions and research areas that need further elucidation in order to better understand the roles of probiotics in the treatment of intestinal diseases.

Keywords: Human gut microbiota, intestinal disease, probiotics

Introduction

In modern times, probiotics have attracted public interest in the course of promoting human intestinal health, and the addition of prebiotics to probiotics, as synbiotics, has also been recommended. Therefore, the global probiotics market has grown rapidly, which makes probiotics not only as a supplement but also a food [1]. The intake of probiotics has been suggested to enhance overall health and immunity. It is now recognized that the human gut microbiota may play a role in the development of metabolic diseases including obesity, diabetes and inflammatory bowel disease [2]. Interestingly, the administration of probiotics has been shown to improve the prognosis and management of those diseases [3]. In addition, although genomic and microbiological characteristics of individual probiotic strains are important, it is necessary to further understand the effect on human gut microbiota during ingestion. This review provides a brief summary of the concept and function of probiotics, and their relationship with intestinal microbiome-associated diseases [4].

Concept of Probiotics and Their Potential Benefits

The Food and Agriculture Organization (FAO) of the United Nations and the World Health Organization (WHO) define probiotics as “live microorganisms which when administered in adequate amounts confer a health benefit on the host.” Probiotics are used constantly to improve the homeostasis of internal microbiota to maintain the human intestinal health [5]. Consequently, the number of harmful bacteria that cannot survive in an acidic environment is decreased, and the beneficial bacteria that grow well in an acidic state proliferate, thereby balancing the intestinal microbiota. The relationship among probiotics, gut microbiota, and disease is shown in Fig. 1.

Probiotics are suggested to have both nutritional and
beneficial immune modulatory effects in patients with inflammatory bowel disease (IBD). In a previous study, the anti-inflammatory effects of probiotics in patients with IBD revealed that probiotics could have desirable effects resulting in clinical benefits [6]. These health-benefiting effects of probiotics include the prevention of constipation, antibiotic-associated diarrhea (ADD), and enhancement of the immune system [7, 8]. Because intestinal microbial communities are not permanent and can be altered by various factors such as lifestyle, diet, and antibiotics, probiotics may be used to treat and prevent a variety of intestinal disorders. The beneficial effects of probiotics and their relationship with intestinal diseases have been demonstrated in a variety of studies (Table 1).

The Homeostasis of Human Gut Microbiota and Its Potential Roles in Human Intestinal Health

The human gut microbiota is composed of complex and diverse microbial communities that are associated with human intestinal health [9]. The gut microbiota is believed to be composed of about 100 trillion microbial cells that provide a wide range of metabolic functions to the host [10]. For example, the intestinal microbiota plays important roles in the degradation of plant-derived complex carbohydrates, which are not digested by the host, due to the lack of enzymes needed to degrade the structural polysaccharides present in plant material [11]. The large intestine contains mostly strict anaerobic bacteria, which can be categorized as either harmful or beneficial [12]. Additionally, the human intestinal microbiota can degrade and produce organic acids and short-chain fatty acids (SCFAs) such as propionate, acetate and butyrate, which have been shown to control the composition of the microbiota. Moreover, human intestinal microbiota are known to perform various functions in the host including intestinal development, homeostasis and protection against pathogenic bacteria. Furthermore, numerous studies have indicated that the dysbiosis of the intestinal microbiota contributes to the development of metabolic diseases including obesity and diabetes, and various intestinal diseases including AAD, IBD, Crohn’s disease (CD),

Fig. 1. Role of probiotics relevant to intestinal diseases via recovery of dysbiosis in the intestinal microbiota. Various intestinal diseases such as AAD, IBD, CD, and CRC are known to be closely associated with dysbiosis of the intestinal microbiota between host-beneficial and host-harmful intestinal bacteria. Once this balanced composition of intestinal microbiota is disrupted, it can cause diarrhea, inflammation, and even colon cancer. Administration of probiotics has been suggested as a therapeutic approach to prevent these intestinal diseases. Probiotics are recognized to have antimicrobial, anti-inflammatory, and even anti-carcinogenic activities, so they may assist in recovering the unbalanced intestinal microbiota. This recovery can also reduce these intestinal diseases or alleviate their symptoms. On the basis of this information, it is now widely believed that there is a strong connection between intestinal microbiota, probiotics, and intestinal health and disease.
ulcerative colitis (UC), and colorectal cancer (CRC) [3, 13].

A well-balanced (homeostatic) gut microbial population is essential for the host and the microbiota to coexist in a mutually beneficial relationship. Failure to maintain gut homeostasis leads to negative changes in the host metabolism which are linked to chronic diseases such as IBD, cancer, cardiovascular disease and metabolic syndrome [14]. It has been recognized that imbalance of the

<table>
<thead>
<tr>
<th>Disease</th>
<th>Used probiotics</th>
<th>Study summary</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>Antibiotic-associated diarrhea (AAD)</td>
<td><em>L. acidophilus</em></td>
<td>Clinical trials on adults (avg. age, 39) who were tested for 12 days in three different groups: bio yogurt (<em>n</em> = 131), commercial yogurt (<em>N</em> = 118), and no yogurt (<em>N</em> = 120). The percentages of participants suffering AAD during this study are 6.9% (bio yogurt), 11.0% (commercial yogurt), and 14.2% (no yogurt), respectively.</td>
<td>[15]</td>
</tr>
<tr>
<td></td>
<td><em>B. animalis</em> subsp. <em>lactis</em></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>L. delbrueckii</em> subsp. <em>bulgaricus</em></td>
<td>202 seniors (avg. age, 70) volunteered for clinical trials (105 yogurt group and 97 control group) during 8 days. This study showed that the percentage of participants suffering AAD was 12.4% in yogurt group and 23.7% in control group, respectively.</td>
<td>[29]</td>
</tr>
<tr>
<td></td>
<td><em>S. thermophilus</em></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Inflammatory bowel disease (IBD)</td>
<td><em>L. acidophilus</em> La-5</td>
<td>176 IBD patients (86 in probiotic yogurt group and 90 in placebo group with general yogurt) and 84 healthy people as a control group with probiotic yogurt volunteered for clinical trials during 8 weeks. The results showed that the numbers of <em>Bifidobacterium</em> (B) and <em>Lactobacillus</em> (L) increased in both probiotic yogurt group (B, 43.8%; L, 36.1%) and control group (B, 11.0%; L, 16.2%), comparing to placebo group (B, -4.3%; L, 3.3%).</td>
<td>[18]</td>
</tr>
<tr>
<td></td>
<td><em>B. animalis</em> subsp. <em>lactis</em> Bb-12</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>B. longum</em> With fructooligosaccharide/inulin mixture for synbiotics</td>
<td>16 IBD patients (avg. age, 43) were tested for 4 weeks with two different groups: probiotic group (<em>n</em> = 8) and placebo group (<em>n</em> = 8). mRNA levels of inflammatory cytokines (TNF-α and IL-1α) were significantly reduced and subsequent biopsies in the test group confirmed reduction of inflammation.</td>
<td>[20]</td>
</tr>
<tr>
<td>Crohn’s disease (CD)</td>
<td><em>L. rhamnosus</em> NCIMB 30174</td>
<td>62 CD patients volunteered for 4 weeks as probiotic group (<em>N</em> = 33) and placebo group (<em>N</em> = 29). However, there was no significant statistical difference between the two groups and inflammation reduction was not observed.</td>
<td>[17]</td>
</tr>
<tr>
<td></td>
<td><em>L. plantarum</em> NCIMB 30173</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>L. acidophilus</em> NCIMB 30175</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>E. faecium</em> NCIMB 30176</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Crohn’s disease (CD) continued</td>
<td>VSL#3 (Lactobacillus, <em>Bifidobacterium</em>, <em>S. thermophilus</em>)</td>
<td>119 CD patients were tested for 1 year as VSL# group (<em>N</em> = 59) and placebo group (<em>N</em> = 60). VSL# group was divided into three subgroups: a screening phase (Day 0), a double-blind treatment phase (Day 1-90), and an open-label treatment phase (Day 91-365). While late VSL#3 group (open-label treatment phase, Day 91-365) showed 42.1% severe recurrence, early VSL#3 group (Day 1-365) showed only 20.5% severe recurrence. All CD patients receiving VSL#3 showed reduced mucosal inflammatory cytokine levels, comparing to the placebo group.</td>
<td>[23]</td>
</tr>
<tr>
<td>Colorectal cancer (CRC)</td>
<td><em>L. paracasei</em></td>
<td>54 CRC patients (avg. age, 64) were tested for three days before surgery in three groups: group A (synbiotics), B (prebiotics) and C (placebo with bowel preparation). Quantitative Real-Time PCR showed that <em>L. paracasei</em> was detected with the highest number in group A (53.8-fold), comparing to groups B (2.5-fold) and C. However, <em>Leuconostoc mesenteroides</em>, <em>P. pentosaceus</em>, and <em>L. plantarum</em> were not detected in group B and C, and was only detected with high number in group A.</td>
<td>[27]</td>
</tr>
<tr>
<td></td>
<td><em>Leu. mesenteroides</em></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>L. plantarum</em></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>P. pentosaceus</em> with prebiotic mixture (beta-glucan, inulin, pectin, and resistant starch) for synbiotics</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>B. longum</em></td>
<td>60 CRC patients were tested for 12 days in a probiotics group (<em>N</em> = 30) and a placebo group (<em>N</em> = 30). The incidence of diarrhea was significantly lower in the probiotics group (26.67%, 8/30) than in the placebo group (53.33%, 16/30).</td>
<td>[30]</td>
</tr>
<tr>
<td></td>
<td><em>L. acidophilus</em></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>E. faecalis</em></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>
microbiota also plays a role in the development of asthma [15]. Therefore, various factors involved in the regulation of the gut microbiome are key in the prevention and treatment of diseases.

**Human Gut Microbiome-Associated Diseases and Probiotics Applications**

**Dysbiosis of the Human Gut Microbiota**

Microbial dysbiosis is defined as the imbalance of the structure and function of the gut intestinal microbiota [16]. It is a common problem in the current era, caused by bacterial infections, changes in diet, and antibiotics. Recent research has demonstrated that dysbiosis of the gut microbiota is associated with the pathogenesis of intestinal disorders such as IBD, irritable bowel syndrome (IBS) and coeliac disease. Beneficial bacteria in the intestinal tract inhibit infiltration and growth of pathogenic bacteria via their competition for space and resources [17]. Interestingly, probiotics are one of the key solutions to restore the gut microbial balance and to prevent infections in patients following antibiotic treatment, which induces gut microbiota dysbiosis.

**Antibiotic-Associated Diarrhea (AAD)**

AAD is a common adverse effect of antibiotic treatment, which results from the disruption of the intestinal microbiota. One of the causes of AAD is a disease-causing bacterium, *Clostridioides* (formerly *Clostridium*) *difficile*, that can cause infection of the large intestine as a result of reduced resistance to antibiotics [18]. Interestingly, it has been suggested that probiotics may be beneficial and safe in the prevention of AAD as demonstrated in several randomized controlled trials (RCT). For example, Goldenberg *et al.* reported that probiotics are effective in the prevention of *C. difficile*-associated diarrhea in adults and children [19, 20]. A previous review and meta-analysis suggested that the protective effects of probiotics as adjunct therapy may be used in the prevention of AAD in outpatients of all ages [18]. In this study, it has been reported that probiotic intervention may reduce the risk of AAD by 51% with no apparent increase in the risk of side effects [18]. Moreover, combined results in various trials showed that *Lactobacillus rhamnosus* and *Saccharomyces boulardii* were the most protective against AAD. Further clinical trials for ADD are summarized in Table 1. However, the exact cause of AAD has not yet been determined, thus further studies are needed to elucidate the underlying mechanisms of AAD as caused by bacterial infection.

**Inflammatory Bowel Disease (IBD)**

IBD is a chronic inflammatory disease of the gastrointestinal tract. IBD encompasses Crohn’s disease (CD), ulcerative colitis (UC) and indeterminate colitis (IC), which can be differentiated by the location of the inflammation in the gastrointestinal (GI) tract [21]. Although the cause of IBD is not yet well understood, it is thought to be an abnormal reaction of the body’s immune system. Unbalanced diet and stress are considered to be potential factors for IBD. Some evidence suggests that intestinal microbiota may be involved in the pathogenesis of IBD. Furthermore, a number of studies have revealed the difference in the composition of the microbiota between healthy individuals and IBD patients [22]. It has also been suggested that maintaining the balance of intestinal microbiota may be important in preventing IBD [23]. In the past decades, probiotics have gained much attention as a potential therapy to modulate the microbiota and its beneficial effects in IBD. For example, probiotics were used for inducing remission and maintenance therapy in ulcerative colitis [24]. However, a recent study indicated that probiotic supplementation in patients with inflammatory bowel disease is a promising adjuvant treatment in UC but not in CD [21]. Therefore, there is still insufficient data available regarding the efficacy of probiotics to provide generalized recommendations for use in patients with CD.

**Crohn’s Disease (CD)**

CD is an inflammatory bowel disease that affects the entire GIT, which is accompanied by symptoms such as abdominal pain, diarrhea, fever, fatigue, and weight loss. Although the cause of CD is still unknown, it has been hypothesized that various factors such as microbiological, genetic and environmental factors play a role in the development of CD [25]. There is no definite cure yet for CD, but there are several ways to alleviate the symptoms. For example, steroid medications may be used to reduce intestinal inflammation, and immunosuppressants may be taken to reduce the activity of the immune system [26]. Similarly, probiotics may offer an alternative approach in addition to conventional therapy. For example, Fedorak *et al.* demonstrated that supplementation of probiotics in CD patients was beneficial after surgery as compared with the late supplementation [27]. However, another recent study demonstrated that multi-strain probiotic adjuvant treatment had no significant changes in intestinal inflammation following the probiotic therapy in patients with CD [21]. Therefore, due to inconsistencies in the results of clinical trials involving probiotic supplementation in the treatment...
of CD, further research is needed to identify the roles of probiotics.

**Colorectal Cancer (CRC)**  
CRC, also known as bowel cancer or colon cancer, is a cancer in any part of the large intestine including the colon or rectum. The incidence rate of colon cancer has been increasing in developed countries such as Europe, the United States, and Australia [28]. Symptoms of the disease include bloody excrement and weight loss. It is mostly caused by aging, lifestyle, but genetic cases are few. However, in developing countries where regular health screening is not possible, the incidence of colon cancer in young people is high [23]. Moreover, people with inflammatory bowel disease such as UC or CD have an increased risk of developing colorectal cancer [29]. Interestingly, recent studies have explored the roles of probiotics in the prevention of colorectal cancer through alteration of the intestinal microbiota and its possible immunomodulatory mechanisms. However, the effectiveness of cancer therapy using probiotics warrants further investigations including clinical trials and animal model studies since not all probiotics have the same effects [30].

Even though the effectiveness of probiotics has been proven through a great many clinical studies as well as animal experiments, much remains to be learned. The study of clinical trials showed that the intake of synbiotics resulted in a significant change in intestinal microbiota in colorectal cancer patients, with an increase in *Lactobacillus* [31]. In some patients who underwent colonic resection, improvement of mucosal structure in the way of better epithelial barrier function was confirmed. Therefore, ingestion of synbiotics may also alter the metabolic activity in the intestines. These findings suggest that probiotics are effective not only in animals but also in humans in the treatment and prevention of intestinal disorders such as colorectal cancer [14].

**Future Prospects**

Previous studies evaluated human intestinal microbiota following ingestion of probiotics have shown promising results to treat human gut microbiome-associated diseases. According to rapid development of NGS technologies, omics studies of the selected probiotics, probably associated with homeostasis of human gut microbiota and even regulation of intestinal diseases, have been recently made available to further understand their effects for maintenance and promotion of human health. These various omics technologies including microbial genomics, transcriptomics, metagenomics, and even metabolomics have extended our knowledge on isolation and selection of new probiotic strains, verification of their real probiotic effects and even molecular mechanisms, and evaluation of their health-promoting effects in vitro as well as in vivo experiments. Therefore, this extended information of probiotics and their effects on human health on an omics level would be an essential key in the development of new probiotics as well as even pharmabiotics. However, overuse of specific probiotics such as *L. rhamnosus* has been reported to cause bacteremia, sepsis, or endocarditis in patients with severe inflammations in digestive organs [32]. Therefore, it is important to use proper amounts of probiotics when they are used on specific patients for a therapeutic purpose.

**Acknowledgments**

The present study was supported by the research fund (19162MFDS037) from the Ministry of Food and Drug Safety, Republic of Korea. This research was also supported by Kyung Hee University in the sabbatical year of 2018 for J-H. Lee.

**Conflict of Interest**

The authors have no financial conflicts of interest to declare.

**References**