

Bacteriocin-Producing Probiotic Lactic Acid Bacteria in Controlling **Dysbiosis of the Gut Microbiota**

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Several strains of lactic acid bacteria are potent probiotics and can cure a variety of diseases using different modes of actions. These bacteria produce antimicrobial peptides, bacteriocins, which inhibit or kill generally closely related bacterial strains and other pathogenic bacteria such as Listeria, Clostridium, and Salmonella. Bacteriocins are cationic peptides that kill the target cells by pore formation and the dissipation of cytosolic contents, leading to cell death. Bacteriocins are also known to modulate native microbiota and host immunity, affecting several health-promoting functions of the host. In this review, we have discussed the ability of bacteriocin-producing probiotic lactic acid bacteria in the modulation of gut microbiota correcting dysbiosis and treatment/ maintenance of a few important human disorders such as chronic infections, inflammatory bowel diseases, obesity, and cancer.

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INTRODUCTION

Gut microbiota is crucial in maintaining the host defense system and homoeostasis, protecting against pathogens, and strengthening the gut integrity. The majority of gut bacteria belong to Bacteroidetes (e.g., Porphyromonas, Prevotella), Firmicutes (e.g., Enterococcus, Lactobacillus, Streptococcus, Ruminococcus, Clostridium), Actinobacteria (e.g., Bifidobacteria), and Proteobacteria (e.g., Escherichia coli) (Zhang et al., 2015). The perturbation in the commensal gut microbial flora causes dysbiosis, which happens due to several factors like imbalanced diet, infection, or the use of antibiotics, which can cause a long-term shift in the gut commensal microflora, promoting a large number of deadly diseases (Lange et al., 2016). There are several diseases associated with the dysbiosis of intestinal microbiota such as viral infections, inflammatory bowel disease (IBD), Crohn's disease (CD), colorectal cancer, and obesity (Kim et al., 2019). Dysbiosis results in the development of diseases related to immune deregulation such as allergy and autoimmune and inflammatory disorders (D'amelio and Sassi, 2017). The colonization of gut by bacteriocin-producing probiotic strains inhibits the adhesion of pathogen to intestinal epithelial cells through competition, clearing niche, and spatial segregation (Heilbronner et al., 2021). Probiotics are living microorganisms that, upon ingestion in an adequate amount, provides a health benefit to a host by improving the intestinal microflora (Binda et al., 2020). Bacteriocins are antimicrobial peptides produced by these bacteria that generally inhibit/kill pathogenic bacteria in the gut and change the composition of gut microbiota in animal models such as mice, pigs, and chickens (Gillor et al., 2008; Yang et al., 2014). The mode of

Bacteriocins in Modulation of Gut Microbiota

action of bacteriocins is different from antibiotics, which kill the target cells by pore formation and membrane disruption. Moreover, bacteriocins, being ribosomally synthesized proteins, are degraded by proteolytic enzymes, and therefore, the pathogens are not able to develop resistance in the gut (Epand et al., 2016; Umu et al., 2017). Further, bacteriocins have a simpler biosynthetic mechanism and are easy to increase their activity against target microorganisms with the help of bioengineering as compared to conventional antibiotics. In addition, a higher specific activity against multidrug-resistant pathogens offers advantages for their applications in therapeutics (Perez et al., 2014). Therefore, the use of bacteriocins and/or bacteriocin-producing probiotics is a novel approach for the treatment of several diseases including enteric infections and the restoration of health-promoting microbial community (Fong et al., 2020).

LACTIC ACID BACTERIA

Probiotic lactic acid bacteria (LAB) are a nonpathogenic heterogeneous group of catalase-negative, Gram-positive, nonsporulating bacteria. They produce lactic acid as a main product from glucose and several growth-inhibiting substances like bacteriocins, bacteriocin-like inhibitory substances (BLISs), hydrogen peroxide, diacetyls, and carbon dioxide. These bacteria need complex nutritional substances for growth such as amino acids, peptides, nucleotide bases, vitamins, fatty acids, and carbohydrates (Mokoena, 2017). They are found in dairy products, fermented meats, fishes, beverages, pickled vegetables, and cereals and in the cavities of human and animals. Important genera include *Lactococcus, Enterococcus, Streptococcus, Pediococcus, Aerococcus, Alliococcus, Carnobacterium, Dolosigranulum, Oenococcus, Tetragenococcus, Vagococcus, Weissella*, and *Lactobacillus* being the largest genus (Bintsis, 2018).

Lactobacilli are the most common probiotics found in humans and other animals. The major species of lactobacilli found in the gut are L. gasseri, L. crispatus, Limnosilactobacillus reuteri, Ligilactobacillus salivarius, and L. ruminis (Walter, 2008; Zheng et al., 2020). A metagenomic analysis suggests that therapy using a combination of different species of Bifidobacterium and Lactobacillus remarkably changes the composition of intestine microbiota in mice (Azad et al., 2018). Other than LAB, Bifidobacterium is considered as the first gut-colonizing microbe that exerts health benefits to the host (O'Callaghan and Sinderen, 2016). In breastfeeding infants, the species of Bifidobacterium are present in a wide range that gradually change with age. B. longum, B. bifidum, and B. breve are generally dominant in the gut of infants, whereas B. catenulatum, B. adolescents, and B. longum are present in adults. They may be used as remedy for the treatment/ maintenance of various gastrointestinal (GI) diseases and restrict the deleterious microorganisms, enhance the GI fence, and inhibit proinflammatory cytokines (Xue et al., 2017).

Bacteria other than LAB also dominate the gut and play crucial functions. For example, harmless *E. coli* Nissle, found in the gut, is a widely utilized probiotics used for the balance of intestine microbiota. It has been revealed that it can restore the production of human β -defensin 2 that could save an intestinal hurdle in opposition to the adherence and capture by pathogenic *E. coli* (Schlee et al., 2007; Liu et al., 2017). Thus, LAB are a major component of gut microbiota that play an important role in maintaining the balance of the total microbial community.

BACTERIOCINS

Bacteriocins are multifunctional, antimicrobial peptides produced by different bacteria that act at low concentrations and generally inhibit the growth of closely related species (narrow spectrum), but recent findings also suggest the occurrence of broad-spectrum bacteriocins (Chi and Holo, 2018; Goval et al., 2018). Bacteriocin-producing cells are resistant to these antimicrobial peptides due to the presence of immunity proteins on the cell membrane of the producer bacteria. Nisin, a bacteriocin produced by several strains of Lactococcus lactis, has received GRAS (generally regarded as safe) status by the American Food and Drug Administration (FDA) and is generally used in food safety (Negash and Tsehai, 2020). According to Zacharof and Lovitt (2012), bacteriocins have been classified into three classes on the basis of biochemical and genetic characteristics: class I bacteriocins are lantibiotics with molecular weight < 5 kDa; they are posttranslationally modified, leading to the formation of methyllanthionine and lanthionine. Class II bacteriocins are non-lantibiotics with molecular weight <10 kDa, non-modified, heat stable, and further divided into three subclasses: class IIa peptide with anti-listerial activities such as pediocinPA1/AcH from Pediococcus species contain the N-terminal conserve sequence YGNGVXC (Nishei et al., 2012); class IIb consists of two peptide bacteriocins such as lactococcin G from L. lactis; in Class IIc, Nand C-terminals are linked by peptide bond forming cyclic bacteriocins, e.g., enterocin AS-48 (Van Belkum et al., 2011). Class III consists of heat-stable and large-sized bacteriocins with molecular weight > 30 kDa, e.g., enterolysin and helviticin (Yang et al., 2014).

Bacteriocins are effective against various human infectious diseases due to their efficacy against several pathogens. For example, pneumonia, meningitis, and sepsis caused by Streptococcus pneumonia can be treated with nisin (Goldstein et al., 1998). The cyclic bacteriocin griselimycin was able to cure tuberculosis in mice (Kling et al., 2015). It has been reported that a few bacteriocins such as pediocin PA-1 and lactocin AL705 show anticancer and anti-inflammatory activities (Huang et al., 2021). For example, nisin inhibits the proliferation of cancer cells by the formation of ion channels on the cell membrane, releasing lactate dehydrogenase, increasing the number of reactive oxygen species, and obstructing the mitochondrial respiration of cancer cells. It was also reported that nisin, in combination with cancer drugs, shows synergistic activity in the clearance of a tumor (Preet et al., 2015). Bacteriocins increase the anti-inflammatory cytokine level and decrease the pro-inflammatory cytokine level by various signaling pathways such as mitogen-activated protein

Bacteriocins in Modulation of Gut Microbiota

kinase and Toll-like receptor (Sassone-Corsi et al., 2016). Thus, bacteriocins are important probiotic metabolites of LAB that can be used for different health-promoting activities of the host.

BACTERIOCIN-PRODUCING LACTIC ACID BACTERIA

There are sufficient studies describing the effect of bacteriocinproducing LAB on changing gut microbiota in animals and humans (Yang et al., 2014; Hernandez-Gonzalez et al., 2021). For example, bacteriocin Abp118, produced by L. salivarius UCC118 isolated from the terminal ileum of a human intestine, shows antilisterial activity in the gut of murine and porcine (Riboult-Bisson et al., 2012). There is change in fecal bacteria community in humans caused by L. plantarum P-8, which is due to the production of plantaricin (Kwok et al., 2015). In another study, intraperitoneally injected nisin F produced by L. lactis ssp. lactis F10 showed a stabilizing effect on bacterial community in the gut of mice (Umu et al., 2017). Another bacteriocin, thuricin CD composed of two peptides, Trna and $Trn\beta$, is secreted by *Bacillus thuringiensis* DPC6431, which kills a wide range of C. difficile isolates without affecting commensal microbiota in a distal colon model (Rea et al., 2010).

Bacteriocin-producing LAB are effective against infections caused by foodborne pathogens like Listeria monocytogenes and several enterococci present in human intestine (Harris et al., 1989; Millete et al., 2008). Pediococcus acidilactici UL5 produces pediocin PA-1, which showed anti-listerial activity in a mouse model without affecting the native intestinal microflora (Dabour et al., 2009). The administration of enterocin CRL35 produced by E. mundtii CRL35 in pregnant mice inhibited the transfer of L. monocytogenes to vital organs (Salvucci et al., 2012). Plantaricin PJ4 produced by L. helveticus PJ4 isolated from the gut of rat showed potent results in reducing weight in obese mice (Bai et al., 2020). Similarly, plantaricin EF produced by L. plantarum NCMIB8826 shows a beneficial effect in diet- induced obese mice (Heeney et al., 2019). The immunomodulatory and immunostimulatory effects of nisin (in the form of commercial preparation)-containing diet was evaluated and increased the CD4 and CD8 T lymphocytes and reduced the B-lymphocyte cell count (Pablo et al., 1999). In another in vivo study, a reduction in the colonization of vancomycinresistant enterococci (VRE) in the intestine of the mice model was reported by administering the bacteriocin producer, L. lactis MM19 and P. acidilactici MM33 isolated from the fecal sample of a human (Millete et al., 2008). When rats were administered S. aureus K followed by treatment with nisin F intranasally, immunosuppressed rats showed pneumonia symptoms that had not been administered with nisin F, while the rats colonized by S. aureus K and treated with nisin F showed a healthy trachea and lungs (De Kwaadsteniet et al., 2009).

Umu et al. (2016) demonstrated the effect of bacteriocinproducing LAB and their isogenic mutants on the modulation of gut microbiota. The bacteriocins used in this study were sakacin A, pediocin PA-1, enterocin P, Q, and L50. They demonstrated that the oral administration of a bacteriocin producer does not change the overall structure, but some beneficial changes occur at a lower taxonomic level in the mice gut, whereas some changes were reversed back after treatment. It was interesting to know that the isogenic mutant of respective strains did not cause such changes, suggesting the role of bacteriocin in the modulation of microbiota (Umu et al., 2016). Oral administration of probiotics such as Lacticaseibacillus casei, L. acidophilus, L. plantarum, and Streptococcus thermophiles in a dose-dependent manner enhanced the number of Ig-A- and Ig-G-producing cells (Azad et al., 2018). The administration of nisin Z and pediocin AcH reduced the colonization of the pathogen when given 8 days prior to infection with vancomycin-resistant Enterococcus (Millette et al., 2008). There is alteration in the composition of gut microbiota when administered with a combination of probiotics, e.g., L. ramnosus, L. acidophilus, and B. bifidum, in mice fed with a high-fat diet (Azad et al., 2018). Thus, there is enough evidence suggesting the role of bacteriocin-producing probiotic LAB in modulating gut microbiota and maintaining host health. Bacteriocin-producing LAB used for the treatment of several diseases are mentioned in Table 1.

GUT MICROBIOTA AND IMMUNE MODULATION

The mucosal immune system protects GI tract from evading pathogens. Mucosa associated lymphoid tissue, epithelial layer and lamina propria are the main parts of the immune system. *L. fermentum, L. crispatus,* and *L. gasseri* are known to interact with dendritic, enterocytes and Treg cells in human GI tract and adaptive immunity is activated to release pro- and anti-inflammatory cytokines (Azad et al., 2018). There is modulation in an innate and adaptive immune system by the antigenic fragments of probiotic strains as they are capable to enter into the intestinal epithelial cells and M cells of Peyer's patches. Cytokines such as interleukin (IL), tumor necrosis factor (TNF), and interferon (IFN) regulate the innate immune system. Similarly, differentiation of CD8 + T-lymphocyte cells into cytotoxic T-lymphocytes kills the virus-infected cells and activates natural killer cells and macrophages, destroying pathogens (Singh and Rao, 2021).

It was reported that fermentation products of the probiotic *Bifidobacterium breve* C-50 trigger the maturation of dendritic cells and promote the survival of dendritic cells (DCs) and IL-10, which show an anti-inflammatory response. Prolonged survival of DC is caused by increased levels of antiapoptotic protein; BCL-xl triggers PBK/Akt phosphorylation, causing maturation by elevating the effect of CD86 and CD83 maturation markers (Hoaru et al., 2006). DC protects feasible gut microflora and dispatches microorganisms to "mesenteric lymph nodes" and results in the production of IgA antibodies to defend the opposition of mucosal invasion (Macpherson and Uhr, 2004; Macpherson et al., 2005). The differentiation of naive T cells into

S. No.	Lactic acid bacteria	Bacteriocins	Diseases/target pathogens	Model	References
1	Lactococcus lactis DPC3147	Lacticin3147,	Clostridium difficile associated diarrhea (CDAD)	in vitro	(Rea et al., 2007)
2	L. garvieae	Garvicin ML	Active Streptococcus pneumonia.	in vitro	(Borrero et al., 2011)
3	L. lactis	Nisin Z	Immunomodulatory effect	Murine model	(Millette et al., 2008)
4	L. lactis	Nisin F	Respiratory infection	Murine model	(De Kwaadsteniet et al., 2009)
5	L. lactis	Nisin	Meniningitis, sepsis, pneumonia	in vitro	(Goldstein et al., 1998)
6	L. lactis	Nisin Z	Enteric pathogens	Mouse model	(Millette et al., 2008)
7	L. lactis	Nisin A	Colorectal cancer	in vitro	(Norouzi et al., 2018)
8	L. lactis	Nisin	Stress reduction	Mice model	(Jia et al., 2018)
9	Lactobacillus salivarius	Bacteriocin Abp118	Listeriosis	Murine model	(Riboult-Bisson et al., 2012)
10	L. salivarius NRRLB	Bacteriocin OR-7	Campylobacter jejuni	Chicken model	(Ilinskaya et al., 2017)
11	L. salivarius	Bactofencin A	Antilisterial, antistaphylococcal	in vitro	(O'Conner et al., 2018)
12	L. curvatus	Lactocin AL705	Listeriosis	in vitro	(Huang et al., 2021)
13	L. rhamnosus	Lactocin 160	Escherichia coli, Bordetella pertussis	in vitro	(Belfiore et al., 2007)
14	Pediococcus acidilactici	Pediocin PA1	Listeriosis	Murine model	(Dabour et al., 2009)
15	P. acidilactici	Pediocin AcH	Enteric pathogens	Mouse model	(Millette et al., 2008)
16	P. acidilactici	Pediocin	Colorectal cancer	in vitro	(Kaur and Kaur, 2015)
17	P. acidilactici K2a2-3	Pediocin PA-1	Anti-cancerActivity	in vitro	(Huang et al., 2021)
18	Enterococcus mundtii RL35	Enterocin CRL35	Listeriosis	Murine model	(Salvucci et al., 2012)
19	E. avium	Avicin	Listeriosis	in vitro	(Birri et al., 2010)
20	<i>E. faecium</i> P13	Enterocin P	Enteric pathogens	in vitro	(De Kwaadsteniet et al., 2006)
21	E. mundtii RL35	Enterocin CRL35	Herpes virus	in vitro	(Wachsman et al., 1999)
22	E. faecium ST5Ha	Bacteriocin ST5Ha	Herpes virus	in vitro	(Todorov et al., 2010)
23	Carnobacterium maltaromaticum	Piscicolin 126, carnobacteriocin	Listeriosis	Pork model	(Martin-Visscher et al., 2008)
24	Streptomyces spp.	Griselimycin	M. tuberculosis	in vivo	(Kling et al., 2015)
25	Leuconostoc citreum GJ7	Kimchichin	Salmonella typhi	in vitro	(Chang and Chang, 2011)
26	Erwinia carotovora NA4	Erwinaocin NA4	Coliphage	in vitro	(Dey et al., 2021)

different types of cell lines like TH-17, TH-2, TH-1, CD8+ repressor, and regulatory T cell depends upon the interaction of DC with specific pattern recognition factors. A study states that cytophage-bacteroides required for development of TH17 cells in lamina propria, which, in turn, maintains the balance between regulatory T-cell populations and TH-17 (Foligne et al., 2007; Delcenserie et al., 2008; Zeuthen et al., 2008). Evidence proved that when germ-free mice were colonized with Bacteroides fragilis NCTC9343, an immense restoration was observed in the number of CD4+ CD45Rb T-cell populations. This restoration was caused by B. fragilis polysaccharide A (Mazmanian et al., 2008). Interestingly, it was observed that mutant strain lacking this polysaccharide A failed to restore the number of CD+ CD45Rb T-cell populations. In a model of colitis, when L. paracasei was administered intragastrically, it rendered a protective effect by reducing the severity of diseases and delaying their progression (Mileti et al., 2009). It was observed that naïve T cells acquired suppressor functions when DC was treated with any one of these probiotics: Streptococcus thermophilus DN-001 621, Bacteroides adolescentesis DN-150 017, and Bifidobacterium animalis DN173 016. The suppressive effect was caused by the decrease in the proliferation of differentiated T cells and IFNy production by CD4+ effector T cells (Baba et al., 2008). Nisin showed an immunomodulatory effect in the mice model, resulting in an increase in CD4+ and CD8 T lymphocytes, with decreased B lymphocytes and its

administration for a long duration might balance the level of B and T lymphocytes. Nisin Z was effective in modulating the innate immune response by lowering the level of proinflammatory cytokines in human peripheral blood mononuclear cells (PBNCs). It can be used in periodontal disease in which there is an initial burst of neutrophils and in its later stages, B- and T-cell-related immune response was shown by the immune system (Shin et al., 2016).

GUT MICROBIOTA AND GUT-BRAIN AXIS

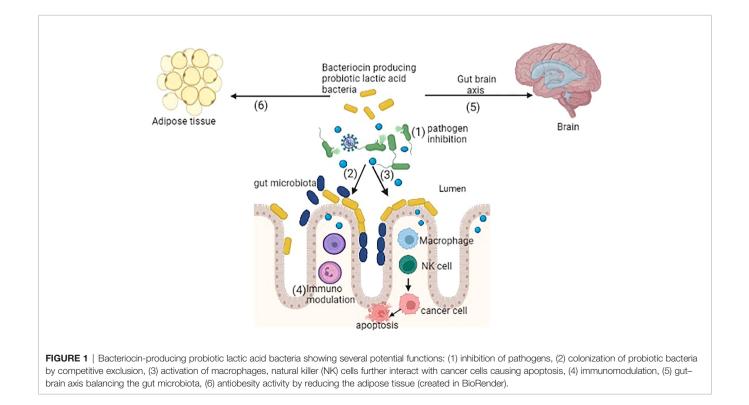
Gut microbes generally secrete amino acids that interact with the ganglion cells (Dinan and Cryan, 2017) in response to the central nervous system (CNS) pattern of chemical messengers (Sanders et al., 2011; Briguglio et al., 2018). Various passages of interaction in the middle of the intestine and CNS have been studied (Dinan and Cryan, 2017). The vagus nerve performs a relationship in the middle of the intestine and spinal cord, which is terminated in the brain stem nuclei and is tactile to deviating fibers (Bonaz et al., 2018),. Thus, the brain stem nuclei may influence numerous bowel roles and convey gestures to more CNS zones, such as the midbrain along with the cerebral cortex (Wang and Wang, 2016). An interchange in the middle of intestine and CNS can also take place through systemic blood flow (Gibson and Mehler, 2019).

The microbiota-gut-brain axis can be noticed as a web with various functions, where midway and sideways, the immune and endocrine systems take part into duplex transmission (Borre et al., 2014). Initially, microbes are capable to substitute, combine, and break neurotransmitters along with transmodulators, like acetate, propionate, butyrate histamine, other pyrimidines, and glutathione (Dinan and Cryan, 2017). These substances act as a neurotransmitter in the brain and stabilize the neuronic venture. However, there is a need of a detailed study to show the direct effect (Angelucci et al., 2019). Furthermore, the gut microbiota produce other proteins that are the deleterious substitute of CNS, being robust irritant cytokines and inborn response activator B cells in the host (Alam et al., 2017). Thus, native microbiota can influence the microbiota-gut-brain axis through antibodymediated nervous and endocrine systems along with various pathways (Gibson and Mehler, 2019). The outcome to these neural changes in the brain can guide to destruction, hypertension, and other coherent diseases (Saunders et al., 2002; Galland, 2014; Johnson and Foster, 2018; Angelucci et al., 2019). Alteration in the gut microbiota is connected to several neurological disorders (Cox and Weiner, 2018), which involve not only hypertension and stress (Nagpal et al., 2018) but also neurodegenerative diseases (Quigley, 2017) and refractory epilepsy (Braakman and Van Ingen, 2018).

Till date, there is no direct evidence available suggesting the role of bacteriocins or bacteriocin-producing LAB in the gutbrain axis. However, gut microbiota may be modulated with producer strains and may indirectly influence the gut-brain axis. In an *in silico* study, the beneficial effects of nisin on neurotransmitter, aquaporin, and commensal gut microbiota were analyzed using high-throughput sequencing, which provided the relationship between the gut microbiota and the neurochemicals used in the gut-brain axis. Nisin showed the highest expression of norepinephrine in the brain as compared to the control group and ciprofloxacin-treated group. Further, it was found that mice treated with nisin showed an increased level of *Lactobacillus*, Bacteroides, and *Bifidobacterium* and decrease in pathogenic *E. coli* and enterococci in the cecum sample. Thus, there is a strong relation between nisin, gut bacterial flora, and reduction in stress triggered by *E. coli* in the mice model (Jia et al., 2018).

BACTERIOCIN-PRODUCING LACTIC ACID BACTERIA AND THEIR ROLE IN DISEASES

Alteration in the normal microbiota of gut causes several chronic diseases, like joint pain, immune-related diseases, metabolic disorders, liver diseases, and various GI diseases (Carding et al., 2015). Bacteriocins may play a role in shaping the host microbiota and indirectly play an important role in correcting dysbiosis and the improvement of host health. Here, we have discussed a few important diseases that occur during the dysbiosis of the gut and their possible cure using bacteriocin-producing probiotic lactic acid bacteria. For clarity, a diagrammatic presentation of the same is depicted in **Figure 1**.



Colonic Infections

Bacteriocins and/or the bacteriocin-producing strains of LAB are documented for the inhibition of several foodborne and clinical pathogens causing severe infections. Most of the LAB bacteriocins are pore formers and interact with the cell membrane to kill the target bacteria through the dissipation of membrane potential and ATP efflux, leading to cell death. Thus, the bacteriocins of LAB may serve as an alternative to clinical antibiotics and can be applied to treat bacterial infections (Pérez-Ramos et al., 2021; Sheoran and Tiwari, 2021; Li et al., 2022).

Clostridium difficile is the main causative bacterium for colonic infection. Using an ex vivo model, it was found that purified nisin was selectively able to deplete C. difficile in a fecal microbial environment without affecting native gut microbiota. The other pathogenic bacteria are E. coli, Salmonella typhi, Campylobacter jejuni, Shigella, and Yersinia enterocolitica (Papaconstantinou and Thomas, 2007). Such infections were found to be minimized by increasing the population of commensal probiotic L. acidophilus (Yun et al., 2014). The combination of lactocin and a chelating agent, ethylenediamine tetraacetic acid, was found effective against E. coli (Belfiore et al., 2007). L. salivarius NRRLB produces the bacteriocin OR-7, which is active against an enteric pathogen, C. jejuni, in the human GI tract (Ilinskaya et al., 2017). Enterocin P inhibits Staphylococcus, Clostridium, L. monocytogenes, Enterococcus faecium, and E. faecalis (De Kwaadsteniet et al., 2006). The inhibitory effect of the bacteriocin-producer L. casei against E. coli and L. monocytogenes was found in the mice model (Soltani et al., 2021). Kimchicin GJ7 produced by L. citreum GJ7 inhibited S. typhi in vitro (Chang and Chang, 2011). Bacteriocin BM1829 produced by L. crustorum MN047 inhibited E. coli, S. typhi, and S. aureus by arresting the cell cycle at the G₁/S checkpoint or destructing the membrane integrity (Yan et al., 2021).

During the ongoing pandemic, viral infections have caused severe diseases and mortality. There are many antiviral agents proposed and tested that have recently been proven successful in treating such infections. However, these therapies showed toxicity and were not able to reduce the symptoms completely. Therefore, it is important to find a safe alternative for the treatment of viral infections (Lehtoranta et al., 2020; Villena et al., 2020). Probiotic LAB exert their antiviral effect by direct interaction with viruses, producing bacteriocins, or enhancing the innate immunity of the host (Al Kassaa et al., 2014; Tiwari et al., 2020). Bacteriocins show antiviral activity against a number of viruses by blocking the synthesis of glycoprotein in the late stage of virus replication (Huang et al., 2021). Acute gastroenteritis is mainly caused by rotavirus, norovirus, and adenovirus in children below 5 years of age. Rotavirus is a double-stranded RNA-lacking envelope that causes the destruction of epithelial cell lining in infants, causing diarrhea (Li et al., 2021). Lacticaseibacillus rhamnosus GG suppresses human rotavirus and induced autophagy in the intestine of piglets by lowering the amount of autophagy proteins, p-mTor, and VPS34-positive cells, Beclin 1 and ATG16L1. L. rhamnosus GG also increases the level of p53

proteins and induces the apoptosis of infected intestinal cells (Wu et al., 2013).

A few enzymatic reactions important for viral infection are inhibited by bacteriocin or bacteriocin-like substances (Salman et al., 2020). It was observed that oral administration of nisin increases the level of CD4+ and CD8+ T lymphocytes and reduces the B cells in mice (Dey et al., 2021). Enterocin CRL35 produced by Enterococcus faecium CRL35 inhibits the replication of the herpes simplex virus, which causes gutrelated ulcerative diseases in humans (Wachsman et al., 1999). Erwiniaocin NA4 produced by Erwinia carotovora NA4 kills the coliphage HSA, and enterocin NKR-5-3C produced by Enterococcus faecium NKR-5-3 shows antagonistic activity against HSV type 1 (Dey et al., 2021). Nisin and sakacin A are effective against the non-enveloped murine norovirus, and bacteriocin ST5Ha produced by Enterococcus faecium ST5Ha shows antiviral activity against the herpes simplex virus (Todorov et al., 2010; Lange-stark et al., 2014). Norovirus is an enteric virus with non-enveloped, single-stranded RNA, which belongs to the family Calciviridae. There is an increase in the number of proteobacteria and a decrease in Bacteroidetes in a norovirus-infected person. The direct binding of norovirus to fecal isolated proteobacteria indicates the modulation of gut microbiota. Further, the attachment of P-particles present on the virus to epithelial cells can be inhibited by L. casei BL23 (Salman et al., 2020). This evidence suggests that bacteriocins and/or the bacteriocinproducing strains of LAB have a potential in preventing the viral infection (Cavicchioli et al., 2018) and therefore, further research is essential to find out the exact mechanism of action before application in therapeutics.

Inflammatory Bowel Disease

IBD is a long-term erythrogenic disease of the gastrointestinal tract that comprises ulcerative colitis (UC) and CD (Bjarnason et al., 2019). The etiology of IBD is not determined yet. It is generally aggravated by improper diet, the disruption of microbiota, and depression. However, a few intestinal microorganisms such as E. coli, C. concisus, and Mycobacterium avium are also involved in the pathophysiology of the IBD (Ryma et al., 2021). UC and CD illnesses have constant provocative states of etiology with various components including genetic susceptibility, hereditary inclination, ecological triggers, changes in the immune system, and an unusual response of gut microbiota. In these diseases, microbial imbalance occurs in a patient, which is characterized by dysbiosis (Sidhu and Vander Poorten, 2017). Fecal microbiota transplant is a potential treatment for IBD, but its success rate is low (Colman and Rubin, 2014). The efficacy of Lactobacillus GG as an adjuvant has been studied in maintaining the remission in CD patients (Scaldaferri et al., 2013). Probiotics seem to be effective and well tolerated by IBD patients, but the role of bacteriocin and its mechanism is still unknown (Ryma et al., 2021).

The microbiota of patients related to IBD is distinct from healthy individuals (Shadnoush et al., 2015). It was observed that the number of Firmicutes like *Faecalibacterium prausnitzi*, which is one of the most abundant gut bacteria, and Bacteroidetes decreased and Proteobacteria and Actinobacteria were increased during IBD. Thus, it is necessary to stabilize the gut microbiota to overcome such diseases where bacteriocin-producing probiotics can play a significant role by promoting the growth of healthy microbiota (Furrie et al., 2005), although a study has shown that probiotic supplementation in IBD is favorable for the cure of ulcerative colitis but not CD (Bjarnason et al., 2019). Alterations in the gut microbiota cause a defect in the mucus laver that increases the intestinal permeability to pathogens and triggers an immune response, causing intestinal inflammation (Michielan, and D'Incà, 2015). Bacteriocin can maintain the integrity of gut epithelium by directly inhibiting/killing the pathogen or can act as a colonizing peptide promoting LAB to occupy niches in the intestine. L. reuteri is a commensal bacterium of gut secretion reuterin, which inhibits several enteropathogens such as yeast, fungi, protozoa, and viruses and promotes the growth of beneficial Gram-positive bacteria (Liu et al., 2020). It was found that probiotics and their metabolites such as short-chain fatty acids play an important role in intestinal dysbiosis and the immunopathogenesis of IBD (Ryma et al., 2021). In a recent study, it was reported that the bacteriocin-producing strains of L. casei, L. plantarum, L. rhamnosus, and L. acidophilus isolated from breast milk competed with intestinal pathogens, reduced the human colorectal adenocarcinoma cell line (HT-29), lowered cholesterol levels, and improved IBD in the mice model (Abdi et al., 2021).

Colorectal Cancer

Colorectal cancer affects the rectum and colon of the large intestine with major symptoms of bloody stool and reduced body weight. It depends on various factors such as diet, lifestyle, and aging (Center et al., 2009). The efficacy of potent LAB was demonstrated through prominent clinical investigation and animal model experiments (Krebs, 2016). The study on the mice model provides evidence that Bifidobacterium with bacteriocin-producing probiotic combination reduces the chances of colorectal cancer (O'Callaghan and Sinderen, 2016). L. acidophilus alone or in combination was found to boost the immunity against colorectal cancer (Zhong et al., 2014). Probiotic bacteria secrete numerous substances with anticancer activity including bacteriocins, toxins, and enzymes. Nisin A produced by L. lactis inhibits tumor cell growth and changes the membrane integrity of liver hepatocellular carcinoma (HepG2). Nisin forms pores in the cell membrane and induces apoptosis through an intrinsic pathway and also acts as an antimetastatic agent by lowering the proliferation of melanoma cells (Norouzi et al., 2018). In addition, pediocin produced by P. acidolactici K2a2-3 inhibits the proliferation of human colon adenocarcinoma cells (HT29) (Soleimanpour et al., 2020).

The *in vitro* effect of colicin E7 produced by *E. coli* on the HT-29 cell line was evaluated for the expression of p53, and bcl-2 shows a decrease in bcl-2 and increase in p53 gene expression (Taherikalani and Ghafourian, 2021). Microcin causes cell membrane depolarization, the fragmentation of DNA, release of phosphatidylserine, and caspase activity (Baindara et al.,

2018). In an *in vitro* study, pediocin produced by *Pediococcus acidilactici* K2a2-3 shows an anticancer activity on HT-29 and DLD-1 cell lines in a dose-dependent manner (Kaur and Kaur, 2015). This evidence suggests the role of bacteriocin either directly or indirectly for the cure of colorectal cancer.

Obesity

Obesity is a metabolic disorder closely related to dysbiosis in the gut microbiota. Probiotics are helpful in modulating the gut microbiota to combat such disorders. The gut microbiota are involved in balancing energy intake and satiety through gut peptide signaling or altering the nervous system. The balance of the regulatory signaling peptide is altered if there is a change in gut microbiota. Hence, obesity can be cured by restoring the gut microbiota. There is change in the ratio of Firmicutes/ Bacteroidetes in obese people (Mazloom et al., 2019). The imbalance was identified by a decrease in the number of Gram-negative aerobes and anaerobe Bacteroidetes and increase in Gram-positive Firmicutes (Sze and Schloss, 2016). However, weight gain and fit metabolic physiology in mice could be passed on via fecal/stool microbiota transplant (Turnbaugh et al., 2008; Liou et al., 2013). Bacteriocinproducing probiotics decrease the absorption of fatty acids and reduce the size of adipocytes and also increases the expression of genes related to oxidation of fatty acids (Wicinski et al., 2020). L. plantarum stimulates the production of TNFa and also regulates the production of leptin hormones (Behrouz et al., 2017).

Probiotics indirectly affect obesity by the production of bacteriocin, which modulates the bacterial content (Million et al., 2013). Treatment with *L. mali* APS isolated from kefir reduces obesity in the mice model. Bacteriocin PJ4 produced by *L. helveticus* is proven to be effective in reducing the inflammation and body weight in the mice model (Bai et al., 2020). Heeney et al. (2019) investigated those mice fed with plantaricin EF-producing *L. plantarum* NCMIB8826 reduced the consumption of high-fat diet and exhibited approximately 10% reduction in weight gain. The same was absent in the group supplemented with the isolgenic ($\Delta plnEFI$) mutant strain LM0419.

CONCLUSIONS AND FUTURE PERSPECTIVE

The bacteriocins produced by probiotic lactic acid bacteria are generally small cationic peptides that kill the target cells by pore formation. These peptides show antimicrobial activity against related strains and pathogenic bacteria such as *Salmonella*, *Staphylococcus*, *Listeria*, *Clostridium*, and *Enterococcus*. Bacteriocins are also effective against viral infections caused by rotavirus, norovirus, adenoviruses, etc. Gut microbiota is an important part of human body and play a key role in stabilizing several body functions. Probiotics and their bacteriocins have potential in modulating the gut microbiota through antimicrobial action and immune modulation and are thus helpful in restoring the balanced microbial community in the gut and host immunity. In addition, the role of bacteriocins has also been demonstrated in colorectal cancer, IBD, and obesity. Thus, there is further need to characterize probiotic bacteria in the gut for their bacteriocin profiling and their role in the establishment of ecological niche of the gut using advanced techniques such as metagenomics, proteomics, and metabolomics. Such inventions will lead the discovery of nature-derived novel products and strategies for the cure of several chronic disorders.

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