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Potential probiotics for regulation of the gut-lung axis to prevent or alleviate influenza in vulnerable populations

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ABSTRACT

Influenza, also known as "flu", is an infectious disease caused by influenza viruses. Three types of influenza virus, A, B, and C, are able to infect humans. In most people, influenza causes mild symptoms, but it can also induce severe complications and death. Annual influenza vaccines are currently the main intervention used to minimize mortality and morbidity. However, vaccination frequently fails to provide adequate protection, especially in the elderly. Traditional flu vaccine targets hemagglutinin to prevent virus infection, but the constant mutation of hemagglutinin means that it is a challenge to develop vaccines quickly enough to keep up with mutations. Thus, other methods of curbing influenza incidence would be welcomed, especially for vulnerable populations. Although influenza viruses primarily infect the respiratory tract, influenza virus infection also induces intestinal dysbiosis. Through gut microbiotaderived secreted products and the circulating immune cells, gut microbiota can affect pulmonary immunity. The crosstalk between the respiratory tract and gut microbiota, termed the "gut-lung axis", is observed in the regulation of immune responses against influenza virus infection or inflammationinduced lung damage, indicating the possibility of using probiotics to prevent influenza virus infection or alleviate respiratory symptoms. In this review, we summarize the current findings on the antiviral functions of particular probiotics and/or combinations and discuss the antiviral mechanisms and immunomodulatory activities of probiotics in vitro, in mice, and in humans. Clinical studies show probiotic supplements can provide health benefits, not only to the elderly or children with compromised immune systems, but also to young- and middle-aged adults.

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Abbreviations: gastrointestinal (GI), antigen-presenting cells (APCs); Shortchain fatty acids (SCFAs), dendritic cells (DCs); nature killer (NK), T-helper 1 (Th1); T-helper 2 (Th2), T-helper 17 (Th17); lipopolysaccharide (LPS), Toll-like receptor (TLR); CC-chemokine receptor 9 (CCR9), C-C motif ligand 25 (CCL25); hemagglutinin (HA), neuraminidase (NA); interferons (IFNs), IFN-stimulated genes (ISGs); myxovirus resistance (Mx), oligoadenylate synthetases (OAS); ribonuclease L (RNase L), macrophage colony-stimulating factor (M-CSF); regulated on activation, normal T cell expressed and secreted (RANTES); upper respiratory tract infections (URTIs), tumor necrosis factor (TNF); interleukin (IL), real-time polymerase chain reaction (RT-PCR); bronchoalveolar lavage fluid (BALF), macrophage (M Φ).

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1. Introduction

Human influenza viruses A, B, and C are RNA viruses. Influenza A and B viruses can cause seasonal epidemics, but only influenza A viruses are known to cause flu pandemics, i.e., global epidemics of influenza. Influenza C viruses, by contrast, usually cause mild res-piratory symptoms and minor localized outbreaks.^{[1](#page-7-0)} For most people, influenza usually causes mild symptoms, such as cough, sore throat, and rhinitis. However, influenza infection can also cause serious diseases such as bronchitis, pneumonia, myocarditis, and encephalitis, and result in high morbidity and mortality in children, pregnant women, and the elderly. Recent estimates find that 291,243-645,832 global seasonal influenza-associated respiratory deaths $(4.0-8.8$ per 100,000 individuals) occur annually,^{[2](#page-7-1)} suggesting an urgent need to prevent influenza infection, especially in the aforementioned vulnerable populations.

Influenza A viruses can be found in human and animal

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populations. The classification of influenza A viruses is based on two glycoproteins on the surface of the virus: hemagglutinin (HA) and neuraminidase (NA). To date, 18 HA subtypes (H1 to H18) and 11 NA subtypes (N1 to N11) have been identified as antigenically distinct. HA attaches to cell surface sialyloligosaccharide receptors, initiating virus entry into host cells. NA enables the virus to be released from the host cell by cleaving sialyloligosaccharide residues from the host cell surface. 3 Thus, both HA and NA play an important role in the virulence and pathogenesis of the influenza virus.

Annual influenza vaccines are the main intervention used to minimize the mortality and morbidity of influenza. Traditional flu vaccines target the surface HA to prevent the virus from binding to human cells. However, vaccination frequently fails to provide adequate protection from infection, especially in people older than 65 years. Its protective effects only range from 30% to 40% in elderly adults.^{[4](#page-7-3)} In addition to immunosenescence that accounts in part for the reduced vaccine efficacy observed in the elderly, 4 the constant mutation of HA through antigenic drift and reassortment also means that it is challenging to keep vaccine development up to date with the latest mutations. A vaccine formulated for any given year may be ineffective in the following year because the rapid mutation and evolution of influenza viruses mean that different strains may become dominant every year. The antibodies induced by a specific influenza virus strain or subtype cannot usually effectively neutralize other strains or subtypes. Moreover, the traditional method of influenza vaccine production is through culturing the virus using embryonated hens' eggs. Some vaccine virus strains grow poorly in eggs, and can cause safety concerns in people with allergy to chicken eggs. 5 Under these circumstances, probioticbased supplements may provide an alternative choice to protect against influenza virus infection, especially in the elderly or those who are allergic to chicken eggs.

Probiotics are live microorganisms that confer health benefits to the host when administered in adequate amounts. They have long been known for their microbiota-modulatory properties and immunomodulatory activities. 6 Probiotics can stimulate antigenpresenting cells (APCs), modulate cytokines or immunoglobulin release, build a barrier lining on the gut epithelial cells, alter mucus secretion and defend against pathogenic bacteria and virus infection.[7](#page-7-6) This review highlights and provides updated information about current probiotic-based supplements with respect to their biological function in the prevention or alleviation of influenza virus infection and infection-induced tissue damage. Advances made over the past 11 years regarding how probiotics inhibit influenza virus infection, and the underlying modulatory mechanisms in vitro, in vivo, and in clinical studies are summarized. The interplay and communication between the gastrointestinal (GI) tract and respiratory system is also discussed.

2. The gut-lung axis in the pathogenesis or regulation of influenza virus infection

"Microbiota" means the community of microorganisms, including bacteria, viruses, fungi, and protozoans, which live in a host. The gut microbiota constitutes a complex ecosystem inhabited and regulated by microorganisms that have co-evolved in mutualistic relationships in the gut. Commensal microbiota produce a variety of metabolites, hormones, essential vitamins, and bioactive products in the host gut system, 8 and the GI tract is constantly communicating with the immune system. Through the secreted products derived from gut microbiota and the circulating immune cells, gut microbiota can influence the local pulmonary immunity. The "gut-lung axis" refers to the crosstalk between the respiratory tract and GI tract.^{[9](#page-7-8)}

For example, gut microbiota-derived endotoxins, microbiota metabolites, and cytokines were able to reach the lung niche through circulation in the blood stream, forming a bidirectional gut-lung axis crosstalk. Short-chain fatty acids (SCFAs) released from the gut were able to reach the bone marrow through circulation to affect hematopoiesis, resulting in an increase in Ly6c monocytes in the bone marrow. After moving to the lung, these patrolling Ly6c⁻ monocytes differentiated into alternatively activated macrophages expressing lower amounts of C-X-C motif chemokine ligand 1 (CXCL1), a neutrophil chemoattractant.¹⁰ Therefore, neutrophil influx was reduced, limiting influenzatriggered lung immunopathology. Moreover, SCFAs directly promoted the antiviral activity of influenza-specific $CD8⁺$ T cells by enhancing their metabolism. Thus, SCFAs released from the gut balanced innate and adaptive immunity, leading to the resolution of influenza infection and prevention of immune-associated pathology.[10](#page-7-9) Pretreatment with bacterial lipopolysaccharide (LPS), which constitute the exterior surface and are shed by all Gramnegative bacteria, triggered an antiviral response through the Toll-like receptor 4 (TLR4) pathway to protect mice from lethal infection with influenza A virus.¹¹ Antibiotic treatment induced gut dysbiosis and impaired the generation of influenza virus-specific $CD4⁺$ T and $CD8⁺$ T cells and antibody responses following respiratory influenza virus infection. Both antibody and T-cell responses were completely restored by a single inoculation of LPS either intranasally or intrarectally.¹² Rectal inoculation was used to mimic the effect of the high levels of commensal bacteria present in the colon. Furthermore, intrarectal injection of other TLR ligands, such as CpG (TLR9 agonist) or Poly I:C (TLR3 agonist), were also able to restore immunity in response to influenza virus infection. Intact commensal microbiota were required to provide signals that induce inflammasome activation in the lung, leading to migration of dendritic cells (DCs) from the lung to the mesenteric lymph node for T-cell priming.^{[12](#page-7-11)} It has also been reported that antibiotic treatment suppressed lung stromal cells from expressing anti-viral type I interferon (IFN) signature, resulting in the acceleration of influenza virus replication in the lung epithelia. Fecal transplantation from control mice to antibiotic-treated mice reversed the antibiotic-induced reduction of IFN signature in lung stromal cells, suggesting that commensal microbiota could drive the IFN signa-ture expression specifically in lung stroma cells.^{[13](#page-7-12)} These findings together indicated that commensal microbiota were able to regulate adaptive immunity in the respiratory mucosa against influenza virus infection ([Fig. 1](#page-2-0)a).

Although influenza viruses primarily adhere to and replicate within epithelial cells spanning from the upper (nasal cavity, oral cavity, pharynx, larynx) to the lower respiratory tract (trachea, bronchi, bronchioles and alveoli), 1 influenza virus infection also affects the gut microbiota. Influenza A virus infection was found to have minor effects on the global lower respiratory tract microbiome in mice. In comparison, influenza A viruses induced robust depletion of bacterial content and disruption of mucus layer integrity in the intestine.¹⁴ Another study also reported that influenza virus infection resulted in significantly altered gut microbiota diversity in mice.¹⁵ Both studies reported decreases in the richness of Firmicutes (as shown by Lactobacillus) after infection with influenza virus in mice. $14,15$ $14,15$ It was found that the fecal microbiome of the H1N1 patients was enriched by Enterococcus, Prevotella, Finegoldia, and Peptoniphilus, while the microbiome of healthy controls was dominated by Blautia, Romboutsia, Collinsella, Bifidobacterium, and other beneficial bacteria. Six biomarkers (Fusicatenibacter, Romboutsia, Anaerostipes, E. hallii group, Ruminococcus torques group, and Blautia) could be used to differentiate the H1N1 patients and healthy people.¹⁶ Indeed, influenza infection in humans is often accompanied by GI symptoms, such as nausea, Y.-H. Wang, A. Limaye, I.-R. Liu et al. Journal of Traditional and Complementary Medicine 13 (2023) 161–169

Fig. 1. The gut-lung axis in the pathogenesis or regulation of influenza virus infection.

(a) Explanations about how gut microbiota might affect the lung immunity against influenza infection. The probiotics are colonized in the gut after one orally uptakes. Intestinal microbiota can release secreted products, such as bacterial ligands and metabolites, which enter the blood stream to affect the immune cells in the circulation. Immune cells can migrate to the lung through circulation in the blood vessels, further regulating the antiviral response and/or lung inflammation. (b) Explanations about how influenza virus infection might affect the composition of gut microbiota. After influenza virus infection, lung-derived CCR9+CD4+T cells were recruited

by intestinal epithelial cells and induced the outgrowth of E. coli. Type I IFN produced in the lung after influenza infection also influenced the intestinal microbiota, resulting in the enrichment of Proteobacteria. In mice infected with influenza virus, the richness of Lactobacillus were decreased in the gut microbiota. In H1N1 patients, fecal microbiome was enriched by Enterococcus, Prevotella, Finegoldia, and Peptoniphilus, while the microbiome of healthy controls was dominated by Blautia, Romboutsia, Collinsella, Bifidobacterium, and other beneficial bacteria.

vomiting, and diarrhea, although no evidence has shown that the influenza virus replicates in the intestine in humans. Intranasal administration of influenza viruses did not infect the small intestine directly but caused lung and intestinal immune injury in mice. Lung-derived CC-chemokine receptor 9 positive (CCR9⁺) CD4⁺ T cells were recruited by C-C motif ligand 25 (CCL25) expressed on intestinal epithelial cells. After migrating into the GI tract, lungderived CD4⁺ T cells induced the outgrowth of E . coli, aberrant Thelper 17 (Th17)-dependent inflammation, and intestinal damage by secreting IFN- γ .^{[17](#page-8-1)} In addition to IFN- γ , type I IFN produced in the lung of the respiratory infection of influenza virus, also influenced lung after respiratory infection of influenza virus also influenced the intestinal microbiota, resulting in the enrichment of Proteobacteria and the depletion of obligate anaerobic bacteria in the gut. Moreover, type I IFN-mediated dysbiosis inhibited the antimicrobial inflammatory responses in the gut against secondary Salmonella infection, further enhancing Salmonella intestinal colonization and systemic dissemination.^{[18](#page-8-2)} These findings together suggest that respiratory infection by influenza viruses may not only affect the local pulmonary immunity but also induce the distal intestinal dysbiosis [\(Fig. 1](#page-2-0)b).

3. Possible mechanisms through which probiotics may modulate influenza virus infection

The microbiota-modulatory effect of probiotics is achieved mainly through the inhibitory activity of probiotics against pathogens, which increases the populations of beneficial microorganisms and decreases the levels of pathogens in the GI tract.¹⁹ The immunomodulatory effect of probiotics on the innate and adaptive immune systems may provide host protection against a wide variety of pathogens. 20 Several probiotic strains have shown antiinfluenza virus activities. $20,21$ $20,21$ Possible mechanisms for the antiviral activity of probiotics have been proposed [\(Fig. 2\)](#page-3-0), including (1) inhibition of viral entry into host cells; (2) secretion of metabolites with antiviral activity; (3) stimulation of innate and adaptive immunity of the host. 2

(1) Inhibition of influenza virus entry into host cells

Some probiotics have been found to bind to influenza virus, blocking virus adsorption on host cells.^{[23](#page-8-7)} Enterococcus faecium NCIMB 10415 inhibited influenza virus A H1N1 and H3N2 via direct physical interaction between bacterial cells and viral particles.^{[24](#page-8-8)} Besides the direct interaction between probiotics and viruses, some probiotics were able to adhere to cells and compete with viruses for attachment to cell adhesion receptors, thereby preventing viral invasion into host cells. 23 23 23 Pretreatment of DCs with the S-layer protein of Lactobacillus acidophilus ATCC 4356 was able to prevent the invasion of influenza A H9N2 into DCs .^{[25](#page-8-9)}

(2) Secretion of metabolites with anti-influenza virus activity

Some probiotics can synthesize antiviral metabolites, such as organic acids, bacteriocins, and hydrogen peroxide. 22 22 22 The bacteriocin from Lactobacillus delbrueckii subsp. bulgaricus 1043 showed inhibitory effects against influenza virus A H7N1 and H7N7. 26 26 26 The enterocin B originating from Enterococcus faecium L3 reduced the reproduction of influenza virus A H3N2 in cells and had a protective effect against influenza viral infection in mice.^{[27](#page-8-11)}

- (3) Stimulation of innate and adaptive immunity of the host
- (a) Effects of probiotics on innate and adaptive immune response against influenza virus infection

Innate immunity functions as the first line of host defense against viral infection. The magnitude of innate response activated by viral infection may influence the adaptive immunity induction. It is now widely accepted that some highly pathogenic influenza viral strains hyperstimulate the immune system, leading to an event called "cytokine storm", which induces tissue damage and causes systemic disease.^{[28](#page-8-12)} Therefore, the risk of viral infection can be significantly reduced if innate immunity is enhanced but the

3. Stimulation of innate and adaptive immunity of the host

Fig. 2. Mechanisms of probiotics against influenza virus infection.

1. Inhibition of viral entry into host cells: some probiotics directly bind to influenza virus, blocking virus adsorption on host epithelial cells. 2. Secretion of metabolites with antiviral activity: antiviral metabolites such as organic acids, bacteriocins, and hydrogen peroxide produced by some probiotics can disrupt virial protein structure or inhibit virus replication.3. Stimulation of innate and adaptive immunity of the host: administration of some probiotics induced dendritic cells (DCs) or lung stromal cells to produce type I and/or type II IFNs, which further activate NK cells. In addition, microbe-associated molecular patterns such as TLR ligands stimulate activation of antigen-presenting cells, including DCs, macrophages (MΦ), and B cells, further priming CD4⁺ T, and CD8⁺ T cells. Cytokines release from T cells can induce B cells production of secretory IgA to neutralize viral infectivity. Moreover, probiotics may alleviate viral infection induced lung inflammation.

excessive production of inflammatory cytokines is prevented. Numerous studies show that some probiotics, most of them belonging to the Lactobacillus and Bifidobacterium genera, provide host protection against viral infection through their modulatory effects on innate and adaptive immunity. 22

Upon viral infection, innate immunity functions as the first line of host defense. Type I IFNs are induced and secreted upon host cell recognition of viral nucleic acids, and are important mediators that exert their antiviral activities though the induction of hundreds of IFN-stimulated genes (ISGs). The products of ISGs exert numerous antiviral effector functions, including blocking viral transcription, degrading viral nucleic acids, inhibiting translation, and modifying protein function to control all steps of viral replication.^{[29](#page-8-13)} The classical ISGs include myxovirus resistance (Mx), 2ʹ,5ʹ-oligoadenylate synthetases (OAS), IFN-stimulated protein of 15 kDa (ISG15), and so on. 30 Mx proteins are dynamin-like GTPases that appear to target viral nucleocapsids, resulting in the inhibition of viral RNA polymerase activity, effectively blocking both transcription and replication of the virus.^{[30](#page-8-14)} OAS are activated by double-stranded RNA binding and can catalyze the formation of $2^{\prime}-5^{\prime}$ oligoadenylates to activate cellular ribonuclease L (RNase L), which in turn, degrade cellular and viral RNA. 31 ISG15 is a small, ubiquitin-like molecule that has numerous antiviral functions including inhibition of virus release, ISGylation of both viral and host proteins, and immunomodulatory cytokine-like properties in its unconjugated form.[30](#page-8-14) Previous studies reported that administration of probiotics induced the expression of IFNs and ISGs. Prophylactical administration of Lactobacillus brevis KB290 significantly alleviated the clinical symptoms induced by influenza virus infection, which

could be attributed to the enhancement of IFN-a production induced by L. brevis KB290 consumption.^{[32](#page-8-16)} Takeda et al. found that oral administration of Lactobacillus plantarum 06CC2 alleviated influenza symptoms in mice, which was associated with the enhancement of IFN- α production through intestinal immunity, as revealed by the increased expressions of IL-12 receptor and IFN- γ in Peyer's patches.^{[33](#page-8-17)} Nakayama *et al.* reported that oral administration of Lactobacillus gasseri SBT2055 prevented body weight loss and decreased viral load in the lung of influenza virus infected mice by the downregulation of viral replication through the induction of antiviral genes Mx1 and OAS1a expression.^{[34](#page-8-18)} Similarly, pretreatment of DCs with the S-layer protein of Lactobacillus acidophilus ATCC 4356 increased Mx1 and ISG15 mRNA expression to inhibit the influenza virus infection.^{[25](#page-8-9)} Lu et al. suggested that oral administration of Lactobacillus mucosae 1025 and Bifidobacterium breve CCFM1026 reduced the viral load in the lungs of influenza virus infected mice, which could be associated with an increase in MxA expression in the lungs resulting from the alteration of gut microbiota.³⁵ As the ISGs are crucial components of the IFN responses and play a key role in establishing an antiviral state for virus clearance and restriction of spread, the anti-influenza virus effects of probiotics could be attributed to their up-regulatory effect on the expression of ISG genes of host cells.

In addition to the upregulation of type 1 IFN and ISG expression, probiotics may alleviate viral infection through the modulation of the production of certain cytokines by immune cells. Oral administration of heat-killed Lactobacillus plantarum 06CC2 alleviated influenza virus infection in mice in correlation with the enhancement of IL-12 and IFN- γ production and the decrease in TNF- α levels in the lung.^{[33](#page-8-17)} Goto et al. demonstrated that oral administration of Lactobacillus acidophilus L-92 showed protective effects against influenza virus infection in mice, which was associated with an increase in expression of various cytokines and chemokines, including IFN- α , IL-1 β , macrophage colony-stimulating factor (M-CSF), eotaxin, and regulated on activation, normal T cell expressed and secreted (RANTES). 36 Park et al. indicated that intranasal or oral administration of Lactobacillus plantarum DK119 conferred protection against influenza virus infection in mice, probably by enhancing the innate immunity of DCs and macrophages and in turn increasing the production of IL-12 and IFN- γ .^{[37](#page-8-21)}
Moreover, probiotics are also able to improve the

Moreover, probiotics are also able to improve the immune system against viral infection by increasing the production of viralspecific antibodies. Nasal administration of Lactobacillus fermentum CJL-112 upregulated the expression of T-helper 1 (Th1) cytokines such as IL-2 and IFN- γ and reduced or left unchanged the levels of Th2 cytokines like IL-4, IL-5, and IL-10 in the lung of influenza virus infected mice. Moreover, nasal administration of L. fermentum CJL-112 also significantly increased the specific antiinfluenza IgA levels in the lungs of influenza virus infected mice.^{[38](#page-8-22)} In comparison, nasal administration of *Lactobacillus rham*nosus CRL1505 significantly increased the levels of IFN- γ and specific anti-influenza IgA and IgG in the serum and/or respiratory tract of influenza virus infected mice.^{[39](#page-8-23)}

(b) Putative mechanisms for the immunomodulatory activities of probiotics

Numerous putative mechanisms for the immunomodulatory activities of probiotics have been proposed. Some extracellular polysaccharides produced by probiotics possess immunomodulatory activities. For instance, the extracellular polysaccharides produced by Lactobacillus delbrueckii OLL1073R-1 increased the expression levels of IFN- α , IFN- β , MxA, and RNase L in porcine intestinal epithelial cells, 40 and significantly protected against influenza virus infection[.41](#page-8-25) Besides extracellular polysaccharides, some cellular components, such as DNA, RNA, and bacterial cell-wall components, including peptidoglycan, S-layer proteins, teichoic acids, capsule, and pellicle can modulate the innate antiviral immune response. The anti-influenza virus effects of Enterococcus faecalis KH2 and Lactobacillus plantarum SNK12 were mediated by the bacterial RNAs. 42 Gao et al. found that the S-layer protein of L. acidophilus ATCC 4356 helped control the exacerbated inflammation caused by influenza virus infection. Secretion of the antiinflammatory cytokine IL-10 by DCs treated with the S-layer protein was significantly higher than secretion from DCs infected with only the H9N2 virus, whereas the pro-inflammatory cytokine TNF- α secretion exhibited an opposing trend.^{[25](#page-8-9)} Inflammation can promote the recruitment of immune cells, but uncontrolled and exacerbated inflammation will induce "cytokine storm", leading to systemic edema and extensive tissue damage.²⁸

4. Clinical findings regarding preventive or relief effects of probiotics against influenza or upper respiratory tract infections

Probiotics provided in capsules, milk or yogurt are often used for application in humans. The clinical effects of probiotics in the prevention of influenza or upper respiratory tract infections (URTIs) and/or on the alleviation of respiratory symptoms have been examined in humans of different ages. Nursing home residents aged 65 and older received Lactobacillus rhamnosus GG capsules daily for six months. Respiratory viral infections as examined by real-time polymerase chain reaction (RT-PCR) were observed in 15.0% of residents with probiotic treatment and in 22.9% of residents with placebo treatment, suggesting that probiotic uptake might reduce the risk for influenza and other respiratory virus in-fections in residents of long-term and chronic care facilities.^{[43](#page-8-27)} Test yogurt containing L. bulgaricus OLL1073R-1 and S. thermophilus OLS3059 may help prevent infection with influenza A virus subtype H3N2 in the elderly with weakened immunity, probably by increasing the production of H3N2-specific IgA in saliva. 44 44 44 Milk with or without *L. casei* strain Shirota was given to the elderly aged 74–92 years, showing that the total number of acute URTI events and the severity of symptoms were not significantly different between the two test groups. Nevertheless, probiotic intervention significantly reduced the mean duration of infection per infection event as compared to the placebo treatment.⁴⁵ Yogurt supplemented with a selected strain of Lactobacillus paracasei N1115 could protect middle-aged and elderly people (aged > 45 years) from acute URTIs, probably due to the increased percentage of total $CD3⁺$ T cells induced by probiotic intervention. 46

For healthy male office workers aged 30-49 years, daily intake of L. casei strain Shirota-fermented milk significantly reduced the incidence and duration of URTIs. The protective effects of probioticfermented milk might be through modulation of the immune system, as revealed by the significantly enhanced nature killer (NK) cell activity and decreased salivary levels of stress marker cortisol at 6 weeks after probiotic intervention versus control milk supplementation. Both control and probiotic milk was able to increase the salivary IgA secretion at 6 and 12 weeks after intervention, but the secreted levels of IgA did not vary significantly between different treatments. 47 For adults aged 25–45 years, test yogurt drink with probiotics, including L. paracasei, L. casei 431, and L. fermentium PCC, significantly reduced the incidence of URTIs and flu-like symptoms, and increased levels of IFN- γ in the serum and secretory IgA in the gut as compared to placebo yogurt drink. There were no significant differences in the productions of IL-4, IL-10, IgA, IgG and IgM in the serum between the probiotic and placebo treatments. 48 In comparison, L. salivarius administrated in powder form to athletes aged $18-35$ years could not provide health benefit in reducing the frequency of URTIs and did not affect blood leukocyte counts or levels of salivary IgA and lysozyme during a spring period of training and competition. 49 College students aged 18–25 years received a probiotic stick containing L. rhamnosus GG and Bifidobacterium lactis BB-12 for 12 weeks, showing that the duration and severity of URTIs were significantly decreased by the combined probiotic intervention. 50 As compared to placebo treatment, L. plantarum DR7 administrated to adults aged 30-60 years could significantly reduce the duration of nasal symptoms of URTIs, while it marginally reduced the duration of general flu symptoms ($P = 0.062$). L. plantarum DR7 administrated to young adults aged < 30 years marginally reduced the duration of general flu symptoms $(P = 0.066)$. L. plantarum DR7 supplement significantly decreased plasma proinflammatory cytokines [tumor necrosis factor (TNF)-a, IFN- γ , IL-1 β] in middle-aged adults, and significantly increased anti-inflammatory cytokines (IL-10, IL-4) in young adults. For both populations, L. plantarum DR7 intervention significantly reduced plasma peroxidation and oxidative stress levels. Therefore, the protective effects of L. plantarum DR7 might be related to its immunomodulatory activities, exerting differential effects on different aged populations. 51

Combined treatment of vitamin C and probiotics, including Lactobacillus acidophilus CUL21 (NCIMB 30156), Lactobacillus acidophilus CUL60 (NCIMB 30157), Bifidobacterium bifidum CUL20 (NCIMB 30153) and Bifidobacterium animalis subsp. lactis CUL34 (NCIMB 30172), significantly reduced the incidence rate of URTIs and the number of days with URTI symptoms in children attending preschool facilities, even though the children did not receive a flu vaccine during the study period. 52 In comparison, children

Table 1

Probiotics contribute to prevention of influenza virus infection or alleviation of respiratory symptoms observed in cells, mice, and humans.

Table 1 (continued)

Abbreviation: dendritic cells (DCs), bronchoalveolar lavage fluid (BALF), interferon (IFN), interleukin (IL), myxovirus resistance (Mx), nature killer (NK), 2',5'-oligoadenylate synthetase (OAS), regulated on activation, normal T cell expressed and secreted (RANTES), tumor necrosis factor-a (TNF-a), colony-forming unit (CFU), multiplicity of infection (MOI), upper respiratory tract infection (URTI), respiratory tract infections (RTI).

receiving L. rhamnosus GG probiotic supplement in milk had fewer days with respiratory symptoms than those receiving control milk without probiotics. However, probiotic intervention was not effective in reducing the occurrence of respiratory viruses and the number of respiratory symptoms observed during viral episodes.^{[53](#page-8-37)} Another study reported that long-term daily intake of milk containing L. rhamnosus GG could significantly decrease the occurrence of respiratory illness in children with recovery of L. rhamnosus GG in their fecal samples. Strain-specific real-time quantitative PCR assay of fecal samples helped confirm the successful colonization of the uptake probiotic in the intestine, providing useful information through which to analyze the association of respiratory illness and probiotic intake in this particular group of children ($n = 128$).^{[54](#page-8-38)} Tablets with or without B. animalis subsp. lactis BB-12 were administrated to 1 or 2-month-old infants, showing that probiotic intervention in early childhood significantly reduced the number of respiratory infections.^{[55](#page-8-39)}

Based on the clinical findings from populations of different ages ([Table 1](#page-5-0)), we know that specific probiotics or their combinations can reduce the duration or risk of influenza and/or other respiratory virus infections and/or relieve the respiratory symptoms after infection. The protective and relief effects of specific probiotics or their combinations can be observed in the elderly with weakened immunity due to immunosenescence, 4 in infants and children with incomplete development of the immune system, 56 and in office workers and college students. These findings suggest that probiotic supplements can provide health benefits, not only to the elderly or children with compromised immune systems, but also to youngand middle-aged adults.

5. Conclusions and future perspectives

The translocation of bacteria from the GI tract to the respiratory system can be observed in sepsis and acute respiratory distress syndrome when barrier integrity is compromised.^{[57](#page-8-41)} Bacteria from the gut can also travel to the lung through aspiration of vomit or esophageal reflux. 58 There is less evidence to show the direct transfer of microorganisms between the gut and lung under normal conditions. However, there is evidence to demonstrate the crosstalk between the respiratory tract and the GI tract, a phenomenon termed the "gut-lung axis". The mucosal surfaces in the lung and gut regulate the immune responses by combating pathogens and preventing excessive inflammation-induced tissue damage. Both the gut and lung are able to influence each other's immune responses through the secreted products released in the blood and the immune cells circulating between sites, which might explain why we can observe the protective effects of probiotics administrated through the oral route against influenza infection in mice and humans. The in vitro cell assay of influenza virus infection reveals the direct inhibition of viral entry or replication by probiotics, which might explain why the intranasal administration of probiotics, but not oral uptake of probiotics, can reduce viral titers and relieve lung inflammation in mice. These studies illustrate the direct use of the nasal spray technique for possible application of probiotics in humans to prevent influenza virus infection or to alleviate respiratory symptoms after infection. The antiviral effects of probiotics through nasal delivery might be faster than through the oral intake. It is possible that nasal administration of probiotics may affect the nasal microbiota. Actually, influenza infection in humans can affect the nasal microbiota, 59 and the nasal commensal Staphylococcus epidermidis is reported to serve as a defense mechanism against influenza virus infection. 60 The relevant studies about the regulation of nasal microbiota are not mentioned in the current review due to space and word limitations.

Clinical trials in humans show promising data demonstrating

that specific probiotics or their combinations can shorten the duration or reduce the risk of influenza and/or other respiratory virus infections, especially in the elderly and children. The clinical findings indicate that specific probiotics or related combinations can provide health benefits for the elderly and children, probably by strengthening their innate and/or adaptive immunity. This is particularly important for aged adults as immunosenescence can reduce the protection of the influenza vaccine.^{[5](#page-7-4)} There are many clinical studies investigating the effects of probiotics as adjuvants to potentiate influenza vaccine efficacy in the elderly. 61 These studies are not mentioned in the current review as, in this article, we aimed to search for probiotic studies pertinent to those who are not suitable to receive vaccination or have a poor response to vaccination. Based on this review, specific probiotics or combinations for which clinical findings are available, can also offer the general public, not only the elderly and children, alternative choices when the seasonal flu vaccine does not prevent against the currently circulating strain of epidemic influenza or cannot meet everyone's need due to vaccine production limit. In addition, the antiviral mechanisms of specific probiotics or combinations and possible explanations for the immunomodulatory activities of probiotics observed in cells, mice and humans have also been addressed, providing valuable information for understanding the mechanistic action of probiotics for further application of probiotics in the future.

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Declaration of competing interest

The authors declare no conflict of interest to report this work.

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