

Can Fortified Bifidobacterium with Mycosporin-like Amino Acid be a New Insight for Neurological Diseases' Treatment?

Hüseyin Sancar Bozkurt^{1*} and Banu Kara²

¹Medical Park Private Tarsus Hospital, Clinic of Gastroenterology, Mersin, Turkey

²University of Health Sciences, Adana Numune Research and Education Hospital, Clinic of Gastroenterology, Adana, Turkey

*Corresponding author: Hüseyin Sancar Bozkurt, Medical Park Private Tarsus Hospital, Gastroenterology, Mersin, Turkey, Tel: 00905054482291; E-mail: sancarb79@gmail.com

Received date: October 05, 2017; Accepted date: December 18, 2017; Published date: December 25, 2017

Copyright: © 2017 Bozkurt HS, et al. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution and reproduction in any medium, provided the original author and source are credited.

Abstract

Gastrointestinal microbiota includes bacteria, archaea, viruses, fungi and other eukaryotes. The genus Bifidobacterium is considered the dominant once; it has an important role on immunologic, hormonal and metabolic homeostasis of the host. Recent studies demonstrated that the Mycosporin-like Amino Acids (MAAs) had prebiotic effects and they modulated host immunity by regulating the proliferation and differentiation of intestinal epithelial cells, macrophage and lymphocytes. Also MAAs modulate NF-kB and tryptophan metabolism. Modulation NF-kB and tryptophan metabolism induced a beneficial effect on central nervous cascade. The safety of Bifidobacterium species is known; although they do not produce MAAs, their presence is required for immunological response continuity of intestine. Thereby we hypothesize that if we could create Bifidobacteria species producing MAAs via genetic engineering; they might have stronger immuno-stimulatory properties and might be used as more potent pharmacological agents in neurological diseases secondary to impaired microbiota.

Keywords Mycosporin-like amino acids; Bifidobacteria; Neurological diseases

Introduction

Microbial organisms live in close association with each other at human's body. The gut microbiota contains many different ecological community of commensal, symbiotic and pathogenic microorganisms [1,2]. The gut microbiota has anti-inflammatory, antioxidant, antioncogenic effects and it contribute to the immunological, hormonal and metabolic homeostasis of the host [3,4]. The genus Bifidobacterium belonging to Actinobacteria phylum and it consist of Gram-positive, non-motile, often branched anaerobic bacteria [5]. The Bifidobacteria are one of the major species of human colon microbiota and they are frequently used as probiotic agents [6]. Bifidobacterium species have immune modulatory, metabolic and anti-inflammatory effects which are not seen in other gastro intestinal flora species [5,7-8]. Bifidobacterium species have the highest level of intrinsic hydrogen peroxide resistance causing antioxidant activity. Bifidobacterium's oligosaccharides metabolism has been shown in many studies; it is separated from others by its fermentation ability [5,9]. Bifidobacteria use the fructose-6-phosphate phosphoketolase pathway to ferment carbohydrates; by this pathway indigestible fructans turn into Short Chain Fatty Acids (SCFA) as butyrate, which have beneficial effect on intestinal immunity and metabolism [10]. Bifidobacteria are the main source of butyric acid production and they are used as probiotic ingredient in many foods [11,12]. Recently diminished production of SCFAs is considered as the cause of antibiotic-associated diarrhoea, inflammatory bowel disease and neurological disorders [13].

Mycosporin-like amino acids

MAAs are low molecular weight (<400 Da) amino acids. They have an ampholyte nature and high denaturation temperature with water soluble property [14]. MAAs acts as UV-absorbers and photoprotectants [14-16]. They improve the growth of beneficial bacteria in the environment of harmful microbiota. MAAs is unique components of red seaweeds and seaweed components are known as supportive reinforcement of the microbiota in intestinal diseases [17-20]. MAAs had been shown to regulate intestinal epithelial cell differentiation and cytokine (IL-1β, IL-6) production [21]. Increased cytokine production, NF-KB activation induced a proliferative effect on epithelial intestinal cells and protect them in experimental colitis secondary to dextran sulphate sodium [22-26]. In vivo experiments showed antiinflammatory effect of MAAs, indicating that they can reinforce membrane barrier function [27-28]. There are two biosynthesis pathways of MAAs. First MAAs biosynthesis pathway is the Shikimate pathway [29] which is known as the synthesis way of aromatic amino acids. Second MAAs biosynthesis pathway is pentose phosphate pathway [30]. In both pathways, 4-deoxygadusol is the common precursor used to produce all MAAs. Trans-aldolase is an enzyme of the non-oxidative phase of the pentose phosphate pathway and Bifidobacterium strains have transaldolase enzyme. Confirmation of a biosynthetic gene cluster for MAAs from Gram-positive bacteria has been showed [31]. Anabaena variabilist PCC 7937 (Cyanobacterium) is able to synthesize MAAs [32]. Genome studies identified a combination of genes, YP_324358 (predicted DHQ synthase) and YP 324357 (O-methyl transferase), which were present only in A. variabilis PCC 7937 and missing in the other Cyanobacteria Anabaena sp. PCC 7120 started to produce MAAs after genomic transfer (YP_324358 and YP_324357 genes) from Anabaena variabilis PCC 7937 [32]. The Bifidobacterium animalis subsp. lactis KLDS 2.0603 strain was demonstrated to have the highest survival rate and adhesion ability in simulated gastrointestinal tract treatments [33]. The

comparative genome analysis revealed that the KLDS 2.0603 has most similar whole genome sequence compared with BB-12 strain. It seems that Cyanobacterium is the source of MAAs and we hypothesize that the genes of Cyanobacterium involved in MAAs biosynthesis could be transferred to the strain *Bifidobacterium animalis* subsp. *lactis* BB-12.

Bifidobacteria's role in gut microbiata and neurological system

The human intestinal microbiota influence neuro development, modulate behavior and contribute to neurological disorders. A functional link between gut microbiota and neurological diseases remains unexplored. Sampson et al. showed that gut bacteria regulate movement disorders in mice and suggest that alterations in the human microbiome represent a risk factor for Parkinson Disease (PD) [34]. Also they showed SCFAs modulate microglia and enhance PD pathophysiology. An increase in inflammatory cytokines has been reported previously in autism spectrum disorders and increased cytokines have been reported in association with a regressive autism phenotype with significant communication challenges and aberrant behaviours [34]. Luna et al. have showed distinctive mucosal microbial signatures in children with autism spectrum disorder and functional gastrointestinal disorders that correlated with cytokine, SCFA s and tryptophan homeostasis [35]. Bifidobacterium species provide a lot SCFA as butyrate which has beneficial effect on immunity, metabolism and oncogenesis.

MAAs and gut-brain axis

In central nervous system, NF-κB signaling is as a pro-inflammatory pathway, as it regulates pro-inflammatory cytokine production, leukocyte recruitment, which are important contributors of the inflammatory response. However, the activation of NF-kB signaling in response to stress can also be a strategy of cytoprotection, as several survival pathways can be activated [25]. NF-kB has apoptotic and antiapoptotic properties in central nervous system. Tryptophan is an essential amino acid for neurotransmitter serotonin (5hydroxytryptamine, 5-HT) products. Impaired tryptophan metabolism has been implicated in the pathophysiology of conditions such as acquired immunodeficiency syndrome-related dementia, Huntington's disease and Alzheimer's disease [36]. The effect of MAAs on central signalling cascades has been investigated in human myelo monocytic THP-1 and THP-1-Blue cells [25]. It seems that modulation of NF-κB and tryptophan metabolism via MAAs has a beneficial effect on central nervous cascade. Beside these properties MAAs also inhibit thiobarbituric acid reactive oxygen species [37].

Conclusion

Significant progress has been made over the last years in recognizing the importance of gut microbiota to brain function. Key findings show that several neurobiological mechanisms link with gut microbiome alterations that potentially contribute to neurological dysfunction, including reduced SCFA production, neuro-inflammation. Creating Bifdobacteria species producing MAAs via genetic engineering could make better quality microbiota and more helpful to human health. MAAs produced via genetic engineering can be used not only as a probiotic, also as a pharmacological agent in intestinal and neurological disorders.

References

- NIH HMP Working Group; Peterson J, Garges S, Giovanni M, McInnes P, et al. (2009) The NIH Human Microbiome Project. Genome Res 19: 2317-2323.
- 2. Sherwood L, Willey J, Woolverton CJ (2013) Prescott's Microbiology, pp: 713-721.
- 3. Cahenzli J, Balmer ML, McCoy KD (2012) Microbial-immune cross-talk and regulation of the immune system. Immunology 138: 12-22.
- Virili C, Centanni M (2017) The role of microbiota in thyroid hormone metabolism and enterohepatic recycling. Mol Cell Endocrinol 17: 30075-30078.
- 5. Mayo B, Sinderen D (2010) Bifidobacteria: Genomics and molecular aspects. Caister Academic Press.
- Ghouri YA, Richards DM, Rahimi EF, Krill JT, Jelinek KA, et al. (2014) Systematic review of randomized controlled trials of probiotics, prebiotics and synbiotics in inflammatory bowel disease. Clin Exp Gastroenterol 7: 473-487.
- Sagar S, Vos AP, Morgan ME, Garssen J, Georgiou NA, et al. (2014) The combination of *Bifidobacterium breves* with non-digestible oligosaccharides suppresses airway inflammation in a murine model for chronic asthma. Biochim Biophys Acta 1842: 573-583.
- Sagar S, Morgan ME, Chen S, Vos AP, Garssen J, et al. (2014) *Bifidobacterium breves* and *Lactobacillus rhamnosus* treatment is as effective as budesonide at reducing inflammation in a murine model for chronic asthma. Respir Res 16: 46.
- 9. Berggren AM, Nyman EM, Lundquist, Björck ME (1996) Influence of orally and rectally administered propionate on cholesterol and glucose metabolism in obese rats. Br J Nutr 76: 287-294.
- Scheithauer TP, Dallinga-Thie GM, de Vos WM, Nieuwdorp M, van Raalte DH (2016) Causality of small and large intestinal microbiota in weight regulation and insulin resistance. Mol Metab 5: 759-770.
- 11. Macfarlane GT, Macfarlane S, Gibson GR (1995) Co-culture of *Bifidobacterium adolescentis* and *Bacteroides thetaiotaomicron* in arabinogalactan-limited chemostats: Effects of dilution rate and pH. Anaerobe 1: 275-281.
- Kato S, Hamouda N, Kano Y, Oikawa Y, Tanaka Y, et al. (2017) Probiotic *Bifidobacterium bifidum* G9-1 attenuates 5-fluorouracil-induced intestinal mucositis in mice via suppression of dysbiosis-related secondary inflammatory responses. Clin Exp Pharmacol Physiol 44: 1017-1025.
- Mortensen PB, Clausen MR (1996) Short-chain fatty acids in the human colon: relation to gastrointestinal health and disease. Scand J Gastroenterol 216: 132-148.
- 14. Llewellyn CA, Airs RL (2010) Distribution and abundance of MAAs in 33 species of microalgae across 13 classes. Mar Drugs 8: 1273-1291.
- 15. Häder DP, Williamson CE, Wängberg SA, Rautio M, Rose KC, et al. (2015) Effects of UV radiation on aquatic ecosystems and interactions with other environmental factors. Photochem Photobiol Sci 14: 108-126.
- Korbee N, Figueroa FL, Aguilera J (2006) Accumulation of mycosporinelike amino acids (MAAs): Biosynthesis, photocontrol and ecophysiological functions. Rev Chil Hist Nat 79: 119-132.
- 17. Kulshreshtha G, Rathgeber B, Stratton G, Thomas N, Evans F, et al. (2014) Feed supplementation with red seaweeds, *Chondrus crispus* and *Sarcodiotheca gaudichaudii*, affects performance, egg quality and gut microbiota of layer hens. Poult Sci 93: 2991-3001.
- Muraoka T, Ishihara K, Oyamada C, Kunitake H, Hirayama I, et al. (2008) Fermentation properties of low-quality red alga Susabinori *Porphyra yezoensis* by intestinal bacteria. Biosci Biotechnol Biochem 72: 1731-1739.
- Walsh AM, Sweeney T, O'Shea CJ, Doyle DN, O'Doherty JV (2013) Effect of dietary laminarin and fucoidan on selected microbiota, intestinal morphology and immune status of the newly weaned pig. Br J Nutr 110: 1630-1638.
- McDonnell P, Figat S, O'Doherty JV (2010) The effect of dietary laminarin and fucoidan in the diet of the weanling piglet on performance,

Page 3 of 3

selected faecal microbial populations and volatile fatty acid concentrations. Animal 4: 579-585.

- 21. Martinez-Augustin O, Rivero-Gutierrez B, Mascaraque C, Sanchez de Medina F (2014) Food derived bioactive peptides and intestinal barrier function. Int J Mol Sci 15: 22857-22873.
- 22. Bersudsky M, Luski L, Fishman D, White RM, Ziv-Sokolovskaya N, et al. (2014) Non-redundant properties of IL-1alpha and IL-1beta during acute colon inflammation in mice. Gut 63: 598-609.
- 23. Ernst M, Thiem S, Nguyen PM, Eissmann M, Putoczki TL. (2014) Epithelial gp130/Stat3 functions: An intestinal signaling node in health and disease. Semin Immunol 26: 29-37.
- Wang Y, Han G, Chen Y, Wang K, Liu G, et al. (2013) Protective role of tumor necrosis factor (TNF) receptors in chronic intestinal inflammation: TNFR1 ablation boosts systemic inflammatory response. Lab Invest 93: 1024-1035.
- 25. Becker K, Hartmann A, Ganzera M, Fuchs D, Gostner JM (2016) Immunomodulatory effects of the mycosporin-like amino acids Shinorine and Porphyra-334. Mar Drugs 14: 119.
- Sinha RP, Singh SP, Hader DP (2007) Database on mycosporins and mycosporin-like amino acids (MAAs) in fungi, cyanobacteria, macroalgae, phytoplankton and animals. J Photochem Photobiol B 89: 29-35.
- 27. Rastogi RP, Sinha RP (2009) Biotechnological and industrial significance of cyanobacterial secondary metabolites. Biotechnol Adv 27: 521-539.
- Rastogi RP, Sonani RR, Madamwar D, Incharoensakdi A (2016) Characterization and antioxidant functions of mycosporin-like amino acids in the *Cyanobacterium nostoc* sp. R76DM. Algal Res 16: 110-118.

- Favre-Bonvin J, Bernillon J, Salin N, Arpin N (1987) Biosynthesis of mycosporins: Mycosporin glutaminol in *Trichothecium roseum*. Phytochemistry 26: 2509-2514.
- Balskus EP, Walsh CT (2010) The genetic and molecular basis for sunscreen biosynthesis in cyanobacteria. Science 329: 1653-1656.
- 31. Miyamoto KT, Komatsu M, Ikeda H (2014) Discovery of gene cluster for mycosporin-like amino acid biosynthesis from Actinomycetales microorganisms and production of a novel mycosporine-like amino acid by heterologous expression. Appl Environ Microbiol 80: 5028-5036.
- Singh SP, Klisch M, Sinha RP, Hader DP (2010) Genome mining of mycosporine-like amino acid (MAA) synthesizing and non-synthesizing cyanobacteria: A bioinformatics study. Genomics 95: 120-128.
- 33. Zhu DQ, Liu F, Sun Y, Yang LM, Xin L, et al. (2015) Genome-wide identification of small RNAs in *Bifidobacterium animalis* subsp. *lactis* KLDS 2.0603 and their regulation role in the adaption to gastrointestinal environment. PLoS ONE 10: e0117373.
- 34. Ashwood P, Krakowiak P, Hertz-Picciotto I, Hansen R, Pessah I, et al (2011) Elevated plasma cytokines in autism spectrum disorders provide evidence of immune dysfunction and are associated with impaired behavioral outcome. Brain Behav Immun 25: 40-45.
- 35. Braun J (2017) Tightening the case for gut microbiota in autism-spectrum disorder. Cell Mol Gastroenterol Hepatol 3: 131-132.
- Ruddick JP, Evans AK, Nutt DJ, Lightman SL, Rook GA, et al. (2006) Tryptophan metabolism in the central nervous system: Medical implications. Expert Rev Mol Med 8: 1-27.
- 37. Wada N, Sakamoto T, Matsugo S (2015) Mycosporin-like amino acids and their derivatives as natural antioxidants. Antioxidants 4: 603-646.