

Review

Therapeutic approaches targeting the gut microbiota in ischemic stroke: current advances and future directions

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Ischemic stroke (IS) is the predominant form of stroke pathology, and its clinical management remains constrained by therapeutic time frame. The gut microbiota (GM), comprising a multitude of bacterial and archaeal cells, surpasses the human cell count by approximately tenfold and significantly contributes to the human organism's growth, development, and overall well-being. The microbiota-gut-brain axis (MGBA) in recent years has established a strong association between gut microbes and the brain, demonstrating their intricate involvement in the progression of IS. The regulation of IS by the GM, encompassing changes in composition, abundance, and distribution, is multifaceted, involving neurological, endocrine, immunological, and metabolic mechanisms. This comprehensive understanding offers novel insights into the therapeutic approaches for IS. The objective of this paper is to examine the mechanisms of interaction between the GM and IS in recent years, assess the therapeutic effects of the GM on IS through various interventions, such as dietary modifications, probiotics, fecal microbiota transplantation, and antibiotics, and offer insights into the potential clinical application of the GM in stroke treatment.

Key words: ischemic stroke, gut microbiota, microbiota-gut-brain axis (MGBA), neural pathways, endocrine pathways, immune pathways

INTRODUCTION

Ischemic stroke (IS), a cerebrovascular disorder resulting from inadequate blood flow to the brain caused by occlusion of cerebral blood vessels, notably blood clots, represents the predominant subtype of stroke [1]. With an annual incidence of approximately 30 million cases, IS constitutes a significant contributor to mortality and morbidity [2]. Presently, the existing therapeutic interventions for IS, such as thrombolysis, neuroprotection, anti-inflammation, and antioxidant therapies, fail to adequately address the clinical requirements [3, 4]. Moreover, the limited timeframe for early treatment and the subsequent harm caused by reperfusion can lead to enduring neuronal impairment or even fatality. The quality of life for IS patients is significantly compromised, and society bears a substantial burden. Therefore, the exploration of novel therapeutic strategies assumes paramount significance.

The gut microbiota (GM) is comprised predominantly of a diverse array of eukaryotic, bacterial, and archaeal organisms that colonize the gastrointestinal tract of animals. In recent times, the GM has garnered significant interest owing to its remarkably high microbial density. The advent of high-throughput sequencing technology has facilitated examination of the composition, distribution, and quantitative alterations of the GM, revealing its intricate associations with various organs, such as the gut-brain axis, gut-liver axis, and gut-cardiac axis [5]. Consequently, these interconnections have been linked to the development of numerous diseases, including neurodegenerative, metabolic, and autoimmune disorders [6]. The formation of human GM diversity remains relatively stable from the age of three, but can be influenced by various factors, such as disease, spirituality, age, dietary structure, and smoking. The microbiota-gut-brain axis (MGBA) serves as a pathway connecting the brain to the GM through neurological, endocrine, immune, and metabolic

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pathways [7]. In the presence of disease states, the development of IS can lead to changes in the GM, while disorders of the GM can worsen the progression of IS [8]. This paper examines the correlation between the GM and IS, with a particular focus on the potential of enhancing the GM to develop a cure for IS.

ISCHEMIC STROKE

IS is a condition characterized by high fatality and disability rates, predominantly affecting the elderly population. However, there has been a notable rise in the incidence of IS cases and related deaths. Recent trends indicate a shift towards younger age groups, particularly individuals aged 18-49, which has been attributed to risk factors like excessive sodium intake and smoking [9]. This demographic transition not only imposes a burden of disability on younger individuals but also contributes to a decline in national productivity. IS has been shown to result in a reduction in beneficial microorganism levels and an increase in opportunistic pathogenic bacteria [10]. This shift in microbial composition can also impact the ratio of the phylum Firmicutes to the phylum Bacteroides, potentially leading to gastrointestinal disorders [11]. The emerging recognition of the neuroprotective role of gut microbes and their metabolites in IS highlights the potential for modulating the GM as a novel therapeutic approach for treating IS.

GASTROINTESTINAL TRACT AND THE GUT MICROBIOTA

The gastrointestinal tract harbors a significant population of immune cells, constituting approximately 70% of the immune system and representing the most extensive aggregation of immune cells within the body [12]. The gut accommodates a vast number of microorganisms, approximately tenfold the total cell count of the body, which actively contribute to the regulation of the body's homeostasis. Perturbation in the ecological balance of the GM has been associated with the progression of various diseases. The GM is composed of various microorganisms, including bacteria, archaea, protozoa, viruses, and fungi. The phyla Firmicutes (Ruminococcus, Clostridium) and Bacteroides (Prevotella and Porphyromonas) are the predominant members, accounting for approximately 90% of the GM [13]. Additionally, the GM contains Actinobacteria (Bifidobacterium), Spirillaceae, Fusobacteria, Cyanophyta, Verrucomicrobium, and Proteus, among others. The GM is widely regarded as the "second brain" of human beings. It plays a crucial role in various physiological processes, including digestion, absorption of nutrients, and the production of metabolic substances. It also exerts influences on energy metabolism, physical development, and overall body functions. Additionally, the GM regulates tight junction proteins, thereby ensuring the integrity of the gastrointestinal barrier, defending against pathogens, and enhancing cellular immunity [14]. The main regulator of T cells is the GM, as evidenced by various studies [15]. In the presence of pathological conditions, GM disorders lead to a decline in immune function, resulting in local or systemic inflammation, translocation of microorganisms through the circulatory system, and the development of neurological, pulmonary, cardiovascular, and other diseases [16]. The GM serves as a producer of crucial bioactive metabolites, including but not limited to short-chain fatty acids (SCFAs,

such as isobutyric, isovaleric, butyric, acetic, propionic, and formic acids) [17], neurotransmitters (such as acetylcholine, 5-hydroxytryptophan, melatonin, norepinephrine, gammaaminobutyric acid, and dopamine), oxidized trimethylamine, and lipopolysaccharides (LPS). These metabolites not only participate in the synthesis of vitamin B and vitamin K, as well as the absorption and metabolism of essential substances like bile acids and sterols, but also actively contribute to maintenance of the body's immune function and energy metabolism [18]. The neurotransmitters generated by SCFAs activate the immune system and bind to G-protein coupled receptors on endocrine cells, thereby stimulating the secretion of gut hormones, such as cholecystokinin and glucagon-like peptide-1 (GLP-1). LPS have the ability to traverse damaged intestinal barrier cells and enter the bloodstream, resulting in a systemic immune response [19].

INTERACTIONS BETWEEN ISCHEMIC STROKE AND THE GUT MICROBIOTA

In recent years, the treatment of patients with IS has been notably impacted by alterations in the GM. Specifically, cerebral ischemic injury is associated with a significant decrease in GM diversity, primarily characterized by changes in the abundance of the phyla Firmicutes and Bacteroidetes [20]. Notably, immune cells in the gastrointestinal tract can be activated through various stimuli, such as injury-associated molecular patterns, exosomes, and antigens. This activation leads to the release of substantial quantities of cytokines, which, in conjunction with chemokines, traverse the blood-brain barrier, thereby intensifying the neuroinflammatory response in IS disease [21]. Additionally, this injury leads to compromised intestinal function, characterized by increased permeability of the intestinal wall and weakened intestinal motility. In contrast, a stable GM has the potential to ischemically safeguard injured brain tissue by modulating intestinal immune cells, specifically Treg cells. Consequently, it can be inferred that the brain, gut, and microbiota engage in intricate interactions following cerebral ischemia, which contribute to the progression of the disease. Therefore, comprehending the interplay among these three components assumes a crucial role in the effective management of IS.

The MGBA serves as a communication system facilitating the interaction between the gastrointestinal tract and the nervous system [22]. This interaction is primarily dependent on neural, endocrine, immune, and metabolic pathways. Through the MGBA, the brain's nervous system can regulate the gastrointestinal tract's functions, such as intestinal motility and secretion, to maintain homeostasis. Simultaneously, the gastrointestinal tract can also influence the neurological processes of the brain's nervous system by regulating neuroimmune and endocrine pathways via the MGBA. Therefore, the MGBA plays a crucial role in maintaining overall bodily homeostasis [23].

Neural pathways

The MGBA is bi-directionally regulated through the central nervous system (CNS), autonomic nervous system (ANS), spinal vagus nerve, and enteric nervous system (ENS). The vagus nerve is responsible for innervating around 70% of the parasympathetic nerves. This nerve pathway serves as the primary means by which gut microbes influence the brain. Research findings indicate that germ-free mice, which lack certain microorganisms, exhibited

significantly heightened functional brain networks and stronger connections compared with mice with standard microbiota. These enhanced networks were approximately twice as robust as those observed in mice with normal microbiota. However, the functional networks of germ-free mice displayed a greater density, less organization, and reduced efficiency in compensating for cerebral ischemic stress. The potential factors contributing to changes in dendritic spine density and immature microglia resulting from GM deletion have been identified [24]. The GM plays a significant regulatory role in modulating the functional connectivity and neuroinflammation in the hippocampal neurons, which are responsible for memory, learning, and cognition, the cingulate cortex, which is closely associated with working memory and executive attention [25], and the thalamus, a crucial brain region for regulating consciousness, alertness, and sleep, in mice with IS [26]. The study found a positive correlation between decreased strength of hippocampal neuronal functional connectivity and the presence of Corynebacterium diphtheriae, Pasteurella, and Proteus. Conversely, a negative correlation was observed between decreased strength of hippocampal neuronal functional connectivity and the relative abundance of Bartonella and Enterococcus faecalis [27].

Endocrine pathways

IS deteriorates intestinal barrier permeability and the GM composition through the levels of hormones, such as norepinephrine, epinephrine, and dopamine, released through endocrine pathways such as the HPA axis [28]. Conversely, it has also been noted that the release of endocrine hormones, such as norepinephrine, can be regulated by the GM, with Escherichia coli being able to stimulate the release of norepinephrine, leading to neuroprotective effects on the brain [29]. During cerebral ischemia, GLP-1 is secreted by intestinal secretory cells. This peptide activates the GLP-1 receptors located in the ischemic regions of the brain, thereby reducing the inflammation mediated by interleukin-6 (IL-6) and ultimately improving the condition of cerebral ischemic injury. Cerebral ischemia triggers a substantial release of IL-6 into the bloodstream, leading to stimulation of the hypothalamus, pituitary, and adrenal glands, resulting in secretion of the hormones [30]. Furthermore, estrogen plays a crucial role in regulating the initiation of immune responses through its promotion of the reestablishment of GM equilibrium, characterized by an increase in beneficial bacteria and a decrease in harmful bacteria. Estrogen also reduces the release of LPS, a metabolite associated with immune activation, while simultaneously enhancing the production of SCFAs [31]. Consequently, the incidence of stroke is significantly lower in women compared with men, and the administration of estrogen-treated fecal microbiota transplantation (FMT) yields noteworthy neurological enhancements in elderly female patients diagnosed with IS [32].

Immune pathways

The gastrointestinal system harbors a significant proportion of the body's immune cells, including regulatory T-cells, B-cells, dendritic cells, macrophages, and T-cells. IS-induced inflammatory responses can lead to alterations in the number of immune cells in the gut. Monocytes, lymphocytes, and neutrophils in the gut can also migrate into brain tissue, leading to secondary neurodegenerative diseases [33]. The GM, which plays a significant role in the immune system, serves as a crucial regulator of T-cell homeostasis [34]. Furthermore, it serves as a vital determinant in the progression of inflammation associated with IS. The involvement of the GM in the development of cerebral ischemic disease can be attributed to its impact on Treg cells, which release the anti-inflammatory factor IL-10 for neuroprotection, as well as yo T cells, known for their innate immune response and secretion of IL-17, which exacerbates neuroinflammation [35]. These findings suggest that the GM can potentially mitigate brain damage in cases of cerebral ischemic disease. Following the initiation of IS, intestinal T cells are instructed to migrate to the brain and release IL-17 in order to amplify the inflammatory response in the brain. Conversely, Treg cells were found to be elevated solely in the gut and facilitated the secretion of IL-10 and transforming growth factor- β (TGF- β) into the brain via the MGBA, thereby mitigating inflammation. Even after a span of 12 months post-IS in crab-eating monkeys, heightened levels of the pro-inflammatory factors interferon- γ (IFN- γ), tumor necrosis factor- α (TNF- α), and IL-6 were still observed in plasma. However, the persistence of systemic inflammation was diminished through regulation of the GM. Transplantation of GM into sterile animals experiencing cerebral ischemia resulted in the polarization of Th1 and Th17 cells, leading to increased levels of IFN- γ and IL-17. Additionally, more than 25% of the T-cell infiltrate in the ischemically injured brain tissue originated from the intestinal immune population. Conversely, FMT in a healthy state effectively regulates the peripheral immune system by reducing the number of pro-inflammatory cells, such as Th1 and Th17 cells, while increasing the number of Treg cells. This intervention also contributes to the reduction in cerebral infarction volume in ischemic mice. Moreover, the SCFA and LPS metabolites derived from the GM play a crucial role in regulating the inflammatory response. Specifically, the elevation of SCFAs influences G protein-coupled receptors and histone deacetylases, thereby facilitating the upregulation of anti-inflammatory factors (IL-10 and TGF- β) through immune cells. Additionally, SCFAs enhance the activation of Treg cells by augmenting the expression of forkhead box protein P3 (Foxp3), CD4, and CD25 proteins, ultimately mitigating cerebral ischemia/reperfusion injury in mice [36]. In contrast, LPS has the ability to traverse the intestinal barrier, activate B cells, and induce the release of proinflammatory factors such as IL-6, TNF- α , and chemokines from inflammatory cells via the Toll-like receptor 4 (TLR4)-myeloid differentiation factor 88 (MyD88) pathway. This activation can compromise the integrity of the blood-brain barrier, leading to cerebral edema and neurological damage [37]. Furthermore, following cerebral infarction, disruption of the GM can result in a sustained reduction in SCFAs and an enduring elevation in plasma LPS levels, thereby compromising the intestinal barrier and exacerbating the systemic immune response. The suppression of pathogenic bacterial abundance, such as Enterobacteriaceae, in the gastrointestinal tract following IS has been shown to effectively reduce inflammation and cerebral ischemic injury.

IMPROVEMENT OF THE GUT MICROBIOTA IN THE TREATMENT OF ISCHEMIC STROKE

Dietary interventions

Low-protein diets increase the abundance and diversity of GM (mainly in the phyla Firmicutes and Bacteroidetes), reverse pro-inflammatory T-cell polarisation, and exert anti-inflammatory effects [38]. Furthermore, calorie-restricted diets that involve a reduction of 30–40% in caloric intake may have a protective effect against IS through the regulation of GM, specifically the enrichment of *Bifidobacterium* [39]. Likewise, an elevated protein intake raises the likelihood of stroke, while a moderate diet with a reduced protein content (7% or 8% protein diet) effectively improves neurological harm, the blood-brain barrier, swelling, and inflammation in the brains of IS mice. However, it is crucial to avoid excessive reduction of dietary protein, as it can lead to extensive protein-energy malnutrition, worsen motor dysfunction in rats with cerebral ischemic injury, and intensify the inflammatory response in ischemic brain tissue. Additionally, the protective effect on ischemic brain tissue is not significantly influenced by the type of protein, as both soy protein and casein exhibit similar effects.

Probiotics

Probiotics, including species, such as Bifidobacterium spp., Bacillus spp., Streptococcus spp., Enterococcus spp., Lactobacillus, Schizosaccharomyces papyrifera, and Saccharomyces cerevisiae, have been found to positively impact body homeostasis when present in sufficient quantities. Multiple studies have demonstrated that the prolonged utilization of probiotics, specifically Lactobacillus and Clostridium butyricum, can impact the concentrations of brain-derived neurotrophic factor, serotonin, and neurotransmitters like γ -aminobutyric acid. Additionally, they can impede apoptosis, particularly through inhibition of the PI3K/Akt pathway, within ischemic brain regions. Furthermore, probiotics exhibit antioxidant enzyme activities that effectively mitigate neuronal damage and cognitive impairments, thereby ameliorating brain injuries. These findings have been supported by various sources (Table 1). The administration of probiotics to modulate the GM has demonstrated considerable therapeutic potential for cerebral ischemic diseases. Nevertheless,

given the intricate nature of the intestinal microecological environment, the selection and formulation of probiotics pose significant considerations.

Fecal microbiota transplantation

Following ischemic brain injury, GM disorders can elicit microglia-associated inflammatory responses and cognitive dysfunction, which may serve as a potential risk factor for cerebral ischemic diseases [40]. Both FMT and SCFAs demonstrated a reduction in nuclear factor- κ B (NF- κ B) and extracellular regulated protein kinase (ERK) protein activity in ischemic brain tissues by enhancing mitochondrial function, specifically the electron transport chain and oxidative phosphorylation. Additionally, FMT and SCFAs inhibited microglia polarization to M2 type and upregulated the level of tight junction proteins in the blood-brain barrier. Consequently, this pathway attenuated cerebral ischemic injury by suppressing NF- κ B and ERK protein activities in ischemic brain tissues and preventing the upregulation of downstream pro-inflammatory factors [41]. These findings have been supported by various sources (Table 2).

Antibiotics

The use of antibiotics causes changes in the GM, which then impact immune cells in the intestines. These changes include modifications in T cells, the movement of intestinal lymphocytes to ischemic brain tissue, and the regulation of Treg and Th17 cells. Additionally, antibiotics can affect the development of cerebral ischemic injury by influencing the peripheral immune system. The regulatory effects of antibiotics on cerebral ischemic diseases have also been investigated, as summarized in Table 3. The majority of these studies suggest that antibiotics may alleviate IS by exerting both anti-inflammatory and neuroprotective effects. However, there are also studies suggesting that antibiotics can

Table 1. Probiotics for the treatment of ischemic stroke (IS)

Year and researcher	Drugs and dosage	Subjects	Results	Reference
Han <i>et al.</i> , 2024	<i>Enterococcus faecalis</i> (10 ⁷ colony forming unit [CFU]), <i>Lactobacillus acidophilus</i> (10 ⁷ CFU), <i>Bifidobacterium</i> <i>longum</i> (10 ⁷ CFU); oral	Male C57BL/6L mice	Balances GM, improves brain tissue immune function, and promotes neurological recovery.	[46]
Cruz-Martínez et al., 2024	symbiotic mix of <i>Enterococcus faecium</i> (4×10^8 CFU) and agave inulin (860 mg/kg); oral	Male SD rats	Reduces TNF-α levels and increases brain- derived neurotrophic factor expression, promotes neurological recovery.	[47]
Rahman <i>et al.</i> , 2023	Limosilactobacillus reuteri UBLRu-87, Lactiplantibacillus plantarum UBLP-40, Lacticaseibacillus rhamnosus UBLR-58, Ligilactobacillus salivarius UBLS-22, and Bifidobacterium breve UBBr-01 (6.7–9.8×10 ⁹ CFU); oral	Male SD rats	Remodels GM, modulates pro-inflammatory mediators, and improves gut barrier and BBB permeability.	[48]
Daniele <i>et al.</i> , 2023	Bifidobacterium longum R0175, Lactobacillus helveticus R0052 (10^9 CFU); oral	Male C57BL/6L mice	Remodels GM, and promotes neurological recovery.	[49]
Rahmati <i>et al.</i> , 2019	Lacticaseibacillus casei ZT-Lca.106, Lactobacillus acidophilus ZT-Lac.51, Lacticaseibacillus rhamnosus ZT-Lrh.54, Lactobacillus bulgaricus ZT-Lbu.90, Bifidobacterium breve ZT-Bbr.22, Bifidobacteriumlongum ZT-Lca.106, and Streptococcus thermophilus ZT-Sth.20 (10 ⁹ CFU); oral	Male Swiss albino mice	Reduces apoptosis in brain tissue and improves spatial learning and memory deficits.	[50]
Sun <i>et al.</i> , 2016	<i>Clostridium butyricum</i> (10 ⁹ CFU); oral	Male ICR mice	Reduces oxidation levels and apoptosis in brain tissue, elevates butyric acid content and protects damaged nerve cells.	[51]

Year and researcher	Microbiological sources	Subjects	Results	Reference
Wei <i>et al.</i> , 2024	Normal male SD rats	Male SD rats	Reduces malondialdehyde levels in the brain, increases GSH and GPX4 expression, and reduces cerebral infarct volume.	[52]
Zeng <i>et al.</i> , 2023	Normal young or old male C57BL/6L mice	male C57BL/6L mice	GM from aged mice increases valproic acid levels in the brain, exacerbating inflammatory responses and neurological damage.	[53]
Guo <i>et al.</i> , 2023	Male SD rats administered dengzhan shengmai in the model	Male SD rats	Reduces the level of inflammation in the brain and improves neurological damage.	[54]
Li <i>et al.</i> , 2022	Male C57BL/6L mice administered Buzhong Yiqi decoction in the model	Male C57BL/6L mice	Reduces levels of apoptosis and neurological damage in brain tissue.	[55]
Wang <i>et al.</i> , 2022	Male SD rats administered Zhilong Huoxue Tongyu capsule in the model	Male SD rats	Reduces BBB permeability and neurological damage.	[56]
Wang <i>et al.</i> , 2022	Normal male or female C57BL/6L mice	Male or female C57BL/6L mice	GM from female mice reduces systemic inflammatory factors and promotes neurological recovery.	[57]
Li <i>et al</i> ., 2022	Male C57BL/6L mice pre-treated with electroacupuncture in the model	Male C57BL/6L mice	Produces indole-3-propionic acid, which exerts antioxidant effects, reduces inflammatory responses, and ameliorates neurological damage.	[42]
Feng <i>et al.</i> , 2022	Normal male C57BL/6L mice	Male C57BL/6L mice	Reduces systemic IL-17 expression and improves neurological function.	[58]
Zhang <i>et al.</i> , 2021	Male C57BL/6L mice treated with atorvastatin in the model	Male C57BL/6L mice	Regulates immune function, enhances anti-inflammatory effects, and ameliorates nerve damage.	[40]
Lee <i>et al.</i> , 2020	Normal young or aged male C57BL/6L mice	Aged male C57BL/6L mice	GM from young mice have higher levels of SCFAs, attenuating inflammatory responses and neurological damage in the brain.	[59]
Zhang <i>et al.</i> , 2020	Male SD rats administered Tong- Qiao-Huo-Xue Decoction	Male SD rats	Reduces nerve damage and cerebral infarct volume.	[60]
Chen <i>et al.</i> , 2019	Male SD rats administered Puerariae Lobatae Radix with Chuanxiong Rhizoma in the model	Male SD rats	Increases intestinal butyric acid levels, repairs the intestinal barrier, and alleviates cerebral oedema and neurological damage.	[61]
Xia <i>et al.</i> , 2019	Individuals of higher Stroke Dysbiosis Index	Male C57BL/6L mice	GM from patients with high Stroke Dysbiosis Index can exacerbate gut inflammation and exacerbate IS.	[62]
Spychala et al., 2018	Normal young or aged male C57BL/6L mice	Male C57BL/6L mice	GM from aged mice exacerbates inflammatory response and nerve damage.	[63]
Singh <i>et al.</i> , 2016	Normal male C57BL/6L and Rag1 ^{-/-} mice	Germ-free (GF) C57BL/6J and GF Rag1 ^{-/-} female mice	Initiates T_{helper} cells, reduces immune response, and improves neurological function.	[64]
Benakis <i>et</i> <i>al.</i> , 2016	Normal male C57BL/6L mice	Male C57BL/6L mice	Increases Treg cells, reduces inflammation levels and provides neuroprotection.	[65]

Table 2. Fecal microbiota transplantation (FMT) for the treatment of ischemic stroke (IS)

Table 3. Antibiotics for the treatment of ischemic stroke (IS) via the gut microbiota (GM)

Year and researcher	Drugs and dosage	Subjects	Results	Reference
Liu <i>et al</i> ., 2022	Ampicillin (1 g/L), vancomycin (500 mg/L), ciprofloxacin (200 mg/L), meropenem (250 mg/L) and metronidazole (1 g/L); delivered in the drinking water	Male Wistar rats	The simultaneous down-regulation of inflammatory in the cortex, effectively inhibits the initiation of inflammatory responses and facilitates the recovery process.	[66]
Benakis <i>et</i> <i>al.</i> , 2020	Ampicillin (1 g/L), vancomycin (0.5 g/L); delivered in the drinking water	Male C57BL/6L mice	Decrease the number of pro-inflammatory $\gamma\delta T$ cells and IL-17, to improves both the volume of cerebral infarction and neurological recovery following a stroke.	[67]
Chen <i>et al.</i> , 2019	Neomycin (450 mg/L), polymyxin B (150 mg/L); delivered in the drinking water	Male SD rats	Reduce nerve damage and cerebral infarct volume.	[<mark>61</mark>]
Benakis <i>et</i> <i>al.</i> , 2016	Vancomycin (0.5 g/L); delivered in the drinking water	Male C57BL/6L mice	By augmenting regulatory T cells, reducing $\gamma\delta$ T cells, and regulating immune response occurrence, the improvement of cerebral ischemic injury can be achieved.	[65]
Winek <i>et al.</i> , 2016	Ampicillin (1 g/L), vancomycin (500 mg/L), ciprofloxacin (200 mg/L), imipenem (250 mg/L), and metronidazole (1 g/L); delivered in the drinking water	Female C57BL/6L mice	By disrupting the protective effect of GM on the gut, thereby exacerbating the detrimental effects of ischemic injury.	[68]

worsen neurologic injury and the cerebral infarct volume by promoting the expansion of proinflammatory cells and reducing anti-inflammatory cells, thereby affecting Treg and Th17 cells. Hence, there is a need for further investigation into the use of antibiotics in the treatment of cerebral ischemic diseases. It is important to note that the effects of specific GM on cerebral ischemic diseases may vary. Additionally, GM are integral to a symbiotic relationship with the organism. Therefore, caution must be exercised when employing broad-spectrum antibiotics.

Other interventions

Changes in the GM can significantly influence the prognosis of IS. In addition to the aforementioned approaches, electroacupuncture preconditioning and treatment for cerebral ischemia/reperfusion injury have been found to restore microbial diversity and elevate the levels of the gut microbial metabolite isopropanol, which is known for its potent hydroxyl radical scavenging properties [42, 43], thereby exerting neuroprotective effects. Furthermore, exercise preconditioning has been shown to enhance the integrity of the intestinal barrier by augmenting the abundance of lactic acid bacteria and *Alistipes*, while reducing the presence of *Ruminococcus* [44]. This, in turn, leads to the inhibition of inflammatory factors such as NLR family pyrin domain-containing protein 3 (NLRP3), ultimately resulting in improved cognitive function within the brain [45].

SUMMARY AND PROSPECTS

This paper summarises the relationship between transgenics and IS, as well as the treatment of IS by modulating transgenics. As the largest immune organ in the human body, the gut is susceptible to systemic immune responses induced by changes in the GM. Therefore, modulating the GM provides new ideas for treating cerebral ischemic diseases. The following are worth noting: 1. A unified GM platform (containing detailed information on each genus of microorganisms and their roles, mechanisms, and interactions with other microbiota) needs to be established and improved. 2. Detailed molecular mechanisms of GM alteration of neurotransmitters and immune cells affecting ischemic brain tissues need to be further investigated. 3. Detailed studies are needed to investigate how drugs modulate the GM and their metabolites. 4. GM differences between rodents and humans should not be ignored in experimental studies. 5. Large differences in the GM exist between individuals, especially those residing in different regions, and personalised treatment of IS based on the GM is also a noteworthy topic. In conclusion, GM regulation plays an important role in the treatment of cerebral ischemic diseases, and elucidating its detailed mechanism of action will provide new ideas and options for the treatment of IS.

AUTHOR CONTRIBUTIONS

Zhiguo Mao and Jinying Zhang equally contributed to the conception and design of the research; Mingsan Miao contributed to the design of the research; Mingsan Miao, Lin Guo, Xiaoran Wang, and Zhengwang Zhu critically revised the manuscript; and Zhiguo Mao drafted the manuscript. All authors critically revised the manuscript, agreed to be fully accountable for ensuring the integrity and accuracy of the work, and read and approved the final manuscript.

CONFLICT OF INTEREST

No potential conflict of interest was reported by the authors.

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