

REVIEW ARTICLE

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The influence of the gut-brain axis on neurological and psychiatric well-being

Muhammad Sufyan¹, Mohammad Amjad Kamal²

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Abstract

The biological neurocentric view of brain health is being fundamentally organized by our understanding of the gut-brain axis (GBA). A complex, bidirectional communication network corelating the gastrointestinal tract with the central nervous system. This review represents current evidence on the profound influence of the GBA on neurological and psychiatric well-being, highlighting the gut microbiota as a central dogma. Herein, we present the key signaling pathways neural (vagus nerve), endocrine, immune, and metabolic through which the gut and brain continuously communicate. Additionally, the article implicates how disruptions in the gut-brain system contribute to the pathogenesis of disorders including Alzheimer's, Parkinson's, depression, anxiety, and autism disorder. The mechanisms involving neuroinflammation, altered neurotransmitter production, and dysregulated stress responses. Furthermore, critical considerations such as the enteric nervous system ("second brain"), early-life microbial colonization, the impact of common medications, and individual microbiome variability are highlighted. In this review, we also present the therapeutic potential of targeting the GBA through dietary interventions, psychobiotics, and personalized microbiome medicine, for promising approach to future research and clinical practice. This paradigm shift underscores that brain health is potentially linked to gut health, offering novel approaches for preventing and managing neuropsychiatric conditions.

Key words gut-brain axis, central nervous system, neuroinflammation, personalized microbiome medicine

1. Department of the Bioinformatics and Biotechnology, government College University Faisalabad, Pakistan.

2. Future Technologies Research Centre, King Faisal University, AI Ahsa, Saudi Arabia.

Correspondence: Mohammad Amjad Kamal (Future Technologies Research Centre, King Faisal University, AI Ahsa, PO Box 400, Post Code 31982, Saudi Arabia; E-mail: aakamal@kfu.edu.sa).

Introduction

The biological view of the brain as an isolated organ. Brain is impervious to peripheral influences, is fundamentally shifting. It has been recognized the profound interconnection between the central nervous system (CNS) and the gastrointestinal tract, a dynamic communication network known as the GBA [1]. This bidirectional system promotes constant crosstalk, integrating emotional and cognitive centers of the brain with peripheral intestinal functions [2]. There are numerous studies on it. Emerging research strongly implicates this axis as a crucial regulator in both psychiatric and neurological well-being. Studies suggested disruptions within this complex system are potentially linked to the pathogenesis of a diverse range of disorders, including Parkinson's and Alzheimer's disease to depression, anxiety, and autism spectrum disorder [3].

The gut-brain axis is best conceptualized as an extensive, bidirectional communication network. The gut-brain axis correlates the emotional and cognitive centers of the brain with the peripheral functional activities of the gastrointestinal tract [4]. This sophisticated system ensures the integration of gut homeostasis with brain function. The brain influences the gut primarily through the autonomic nervous system (ANS). It regulates primary functions like motility, secretion, and blood flow [5]. However, stress-mediated activation of the hypothalamic-pituitary-adrenal (HPA) axis can smoothly alter gut permeability and microbial composition [6]. On the other hand, the gut exerts a powerful influence on the brain by sending myriad signals that can influence mood, cognition, and behavior [7]. This bottom-up signaling is largely driven by the gut's intrinsic nervous system. The enteric nervous system (ENS), often called the "second brain," which contains over 100 million neurons [7]. The longest cranial nerve is vagus nerve, serves as a primary physical conduit for this regard, transmitting visceral sensory information directly to the brainstem [8, 9]. This continuous, two-way traffic ensures that our mental state can affect gut feelings, and conversely, our gut health can significantly influence our state of mind.

The action of the gut-brain axis is mediated through several parallel and interconnected signaling pathways, ensuring robust communication [10]. These can be classified into neural, endocrine, immune, and metabolic routes. The neural pathway is primarily executed by the vagus nerve. It transmits afferent signals regarding gut state such as distension, nutrient availability, and microbial activity directly to the brainstem [11, 12]. Efferent signals from the brain then modulate gut function in response. The endocrine (or hormonal) pathway linked to gut enteroendocrine cells, and promotes secretion of serotonin (5-HT) and peptide YY (PYY) in response to nutritional and microbial cues [13]. These hormones enter circulation to influence brain function or act locally on vagal terminals. The immune pathway is critical, the gut mucosa hosts approximately 70% of the body's immune cells. Dysbiosis can enhance pro-inflammatory cytokines release (e.g., IL-1 β , IL-6, TNF- α), which can pass the blood-brain barrier (BBB) or activate its endothelial cells to induce neuroinflammation [14]. Finally, metabolic pathways are associated with small molecules, particularly microbial metabolites, which serve as potent systemic messengers.

Importantly, at the core of the modern understanding of the gut-brain axis resides the gut microbiota: a vast, diverse ecosystem of trillions of bacteria, viruses, fungi, and archaea [15]. This microbial community is not a passive passenger but an active endocrine organ that fundamentally maintains axis communication. It influences brain physiology and functions via multiple mechanisms. Firstly, gut bacteria are vast biochemical factories, producing a broad array of neuroactive metabolites [16, 17]. These include short-chain fatty acids (SCFAs) like butyrate,

propionate, and acetate from dietary fiber fermentation. These metabolites exert anti-inflammatory characteristics and can strengthen the blood-brain barrier [18, 19]. Secondly, microbiota plays important roles producing key neurotransmitters; specific strains can synthesize gamma-aminobutyric acid (GABA), serotonin, dopamine, and acetylcholine, which influence host neurotransmission [20, 21]. Furthermore, the microbiota is indispensable for the proper development and function of host immune system. Microbiota play important roles on host immune system, educating immune cells and preventing inappropriate inflammation that could adversely affect the brain [22]. The gut microbiota establishes itself as a central regulator. It induces the complex symphony of the gut-brain axis by modulating the production of signaling molecules, regulating immune responses, and maintaining gut barrier integrity. This review article highlights the thorough mechanisms that underpin the GBA, focusing on the pivotal role of the gut microbiota. Herein, we present current evidence on how this gut-brain communication influences brain health and disease.

Mechanisms of communication: How the gut talks to the brain

A complex multi-channel network of signaling pathways facilitate the gut and the brain axis. These mechanisms ensure a constant flow of communication, allowing the brain to monitor gut activity. Also, the gut exerts a profound influence on brain function and behavior [23]. The primary routes of this communication can be classified into endocrine, neural, immune, and metabolic pathways. Importantly, the most direct line of communication is the neural pathway which is primarily mediated by the vagus nerve. This cranial nerve serves transmitting sensory information from the gut lumen (such as nutrient status and microbial activity) directly to the brainstem [24]. However, efferent signals from the brain modulate gut functions like motility and secretion. Besides, the endocrine pathway involves gut enteroendocrine cells that release neuroactive hormones, such as serotonin and peptide YY, into the bloodstream in response to nutritional and microbial cues [13, 25]. These hormones directly influence brain regions contributes in mood and appetite regulation. Thus, the most crucial pathway for pathology is the immune system [26, 27]. The gut mucosa houses a vast portion of the body's immune cells. Dysbiosis mediates intestinal integrity, leading to a "leaky gut" that allows bacterial fragments like lipopolysaccharide (LPS) to enter circulation [28, 29]. This facilitates a systemic inflammatory response, producing cytokines that can cross the blood-brain barrier or activate its cells, leading to neuroinflammation a key contributor to numerous neurological and psychiatric disorders [30, 31]. The gut microbiota produces small molecules involves in the metabolic pathway. Key small molecules are SCFAs like butyrate, which possess anti-inflammatory properties and can strengthen the blood-brain barrier, directly influencing brain health and function (**Figure 1**).

The gut-brain axis in neurological disorders

Accumulating evidence suggested the critical role of GBA dysfunction in the pathogenesis and progression of major neurological disorders [32]. The neural, immune, and metabolic communication channels between the gut and brain become conduits for both detrimental and protective influences. This mechanism manifests new perspectives on disease origins and potential therapeutic avenues. In Parkinson's disease (PD), the GBA is central player to the "dual-hit" hypothesis, which proposes that an unknown pathogen may trigger the misfolding of alpha-synuclein protein first in the gut's enteric nervous system [33, 34]. This pathology is then thought to propagate via the vagus nerve to the brainstem and ultimately the substantia nigra [35].

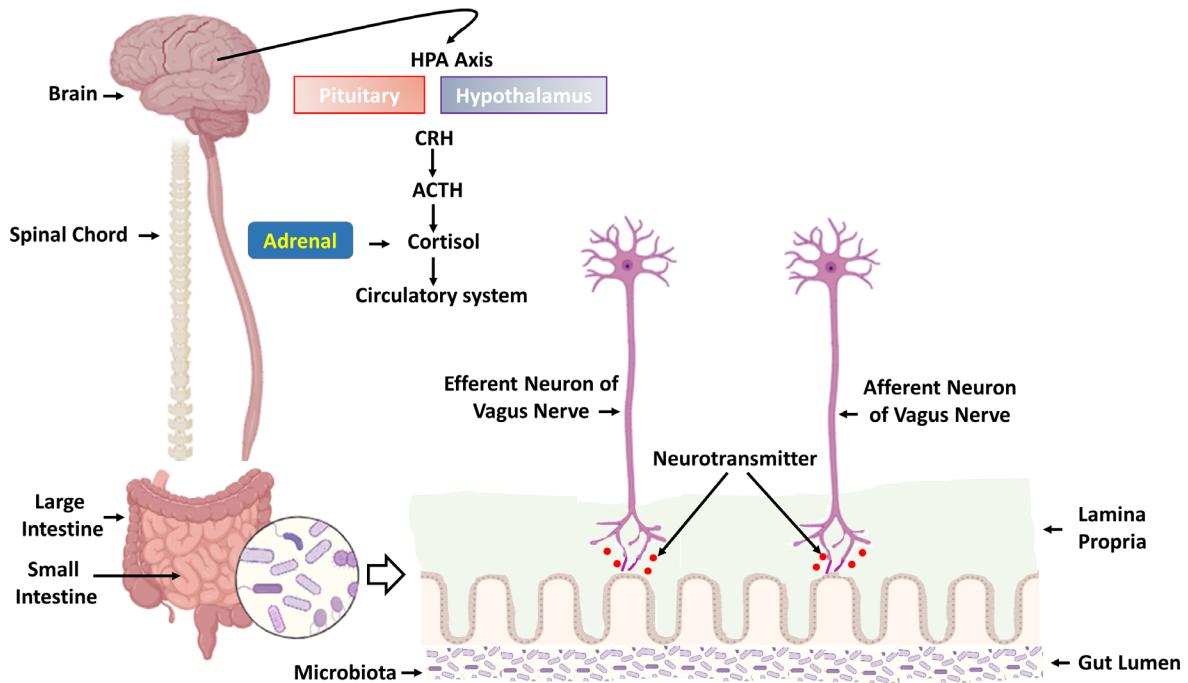


Figure 1. The integrated communication pathways of the gut-brain axis. This schematic illustrates the bidirectional signaling network connecting the Central Nervous System (CNS), comprising the Brain and Spinal Cord, with the gastrointestinal tract. The Hypothalamus, Pituitary, and Adrenal glands form the HPA axis, the body's central stress response system, which is modulated by gut-derived signals. Key communication occurs via the Vagus Nerve, where Afferent neurons relay visceral information from the gut to the CNS, and Efferent neurons carry regulatory commands back to the gut. Within the intestinal wall, the Lamina Propria houses immune cells and nerve fibers that interact with the Gut Lumen, where the Microbiota reside. These microbes produce Neurotransmitters and metabolites that influence local and central nervous system function, creating a continuous feedback loop between the Small Intestine and Large Intestine and the brain, fundamental to both neurological and psychiatric well-being.

Additionally, individuals with vagotomies show a reduced risk of PD, and specific gut microbial profiles. It is characterized by increased pro-inflammatory species and decreased SCFA-producers, are consistently identified in patients [36, 37]. These microbial shifts promote inflammation and exacerbate alpha-synuclein aggregation.

Similarly, in Alzheimer's disease (AD), gut dysbiosis is linked to core disease mechanisms [38]. Certain microbiota influence the production and clearance of amyloid-beta peptides through the modulation of the blood-brain barrier and systemic inflammation [39]. Moreover, microbial metabolites including pro-inflammatory molecules from detrimental bacteria and a deficiency of the anti-inflammatory short-chain fatty acid butyrate, are implicated in driving neuroinflammation and tau phosphorylation [40, 41]. The GBA also plays a role in stroke outcomes, conversely, pre- and post-stroke microbiota composition influences the severity of brain injury and the effectiveness of recovery through immune modulation [42]. Furthermore, alterations in the gut microbiome have been observed in autism spectrum disorder (ASD) and multiple sclerosis (MS), whereas microbial metabolites appear to influence neurodevelopment, immune activation, and demyelination [43, 44]. This collective evidence demonstrates the GBA not as a peripheral player, but as a fundamental component in understanding the complex etiology and progression of a wide spectrum of neurological conditions.

The gut-brain axis in psychiatric well-being and disorders

The potential influence of the GBA extends decisively into the

realm of psychiatric well-being, with dysregulation of gut-mediated biological pathways. It was now recognized as a significant factor in the etiology and symptomology of major mental health disorders. Microbiota possesses vital role in producing neuroactive compounds, regulation of inflammation, and influencing the stress response system. It provides a strong biological correlation between intestinal health and emotional states, manifesting a paradigm shift in how psychiatric conditions are conceptualized and potentially treated [45, 46]. Additionally, distinct gut microbial population have been identified, with a decreased overall diversity and reduction of anti-inflammatory, SCFA-producing bacteria in Major Depressive Disorder (MDD) and anxiety disorders [47, 48]. The dysbiosis promotes a leaky gut, allowing bacterial LPS to enter circulation and trigger a chronic, low-grade inflammatory state [49]. The Pro-inflammatory cytokines can cross the blood-brain barrier, disrupting neuroplasticity, reducing the production of critical neurotrophins like BDNF, and shifting neurotransmitter metabolism [50]. Also, inflammation can shunt tryptophan away from serotonin synthesis and towards the neurotoxic kynurene pathway, directly impacting mood regulation (Figure 2).

The GBA also critically modulates the body's central stress response, the HPA axis [51]. The gut microbiota is essential for its healthy development in infancy and its regulation in adulthood [52]. Dysbiosis promotes HPA axis hyperactivity, resulting in exaggerated cortisol release and a heightened physiological response to stress, which is a well-established risk factor for anxiety and depression [53, 54]. This is specifically relevant in the context of early-life adversity. Whereas, stress-induced alterations to the microbiome can create a lifelong vulnerability to psychiatric

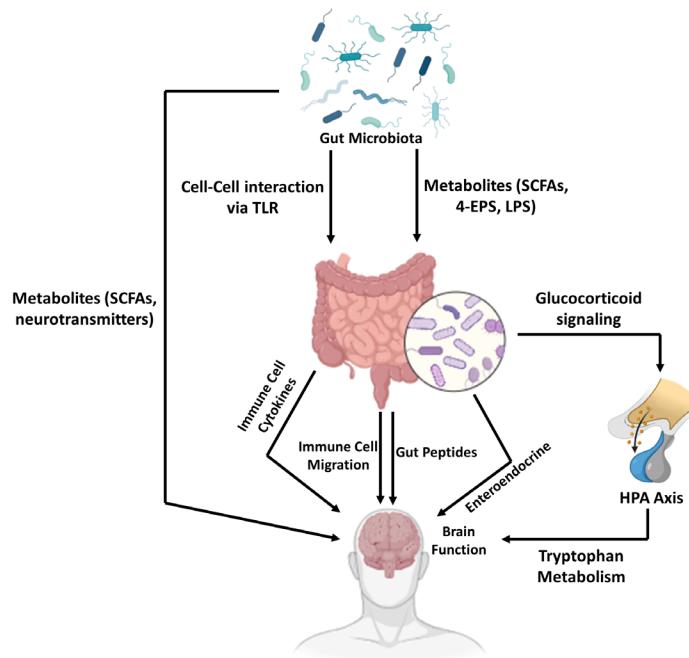


Figure 2. Molecular and Cellular Pathways of Gut-Brain Axis Communication. This schematic detail the key mechanisms by which the gut microbiota signals the brain. Commensal bacteria produce diverse Metabolites, including beneficial SCFA, neuroactive compounds (Neurotransmitter), and the pro-inflammatory LPS. These molecules mediate Cell-Cell interaction via TLR on host cells, triggering Immune cell cytokines and systemic Immune cell migration, potentially driving neuroinflammation. Enterendoocrine cells sense these signals and release Gut Peptides that act on local nerves or enter circulation. The microbiota also critically regulates Tryptophan Metabolism, influencing the production of neuroactive kynurene pathway metabolites. Collectively, these pathways modulate Brain Function and the HPA Axis, leading to Glucocorticoid release, which provides feedback to the gut, completing a bidirectional loop central to neurological and psychiatric health.

illness. Furthermore, emerging research in ASD suggests a strong gut-brain connection. Conversely, microbial imbalances contribute to behavioral symptoms through the generation of metabolites that alter neuronal function and immune communication [55]. Therefore, this collective evidence strongly suggests the gut microbiome is an active contributor to psychiatric well-being. Besides, gut microbiome manifests promising avenues for microbiome-targeted interventions like psychobiotics and dietary strategies to augment mental health (Table 1) [56-65].

The role of the enteric nervous system (ENS): The "Second Brain"

The "second brain", the ENS is an intricate network of over 100 million neurons embedded within the walls of the gastrointestinal tract [66]. This vast and autonomous system is a fundamental, yet often overlooked. Component of the gut-brain axis, acting not as a passive conduit but as a primary regulator and translator of signals between the gut lumen and the CNS [67]. Its sophisticated integration of local microbial, hormonal, and mechanical cues positions is a critical regulator of both digestive and neurological health. The ENS operates with a significant degree of independence from the CNS. It has been revealed that coordinating complex digestive processes such as motility, secretion, and blood flow on its own [68]. However, it maintains constant bidirectional communication with the brain, primarily through the vagus nerve. This connection allows the ENS to get the crucial information about the gut's state, including nutrient status, the presence of pathogens, and microbial metabolic activity [69]. Similarly, the ENS functions as a sensory interface, sampling

the gut's environment and sending integrated signals that can influence mood, stress levels, and overall well-being [69, 70].

The ENS plays important role in neurological and psychiatric disorders. The ENS is a major site of neurotransmitter production. The ENS synthesizes approximately 90% of the body's serotonin, a key regulator of mood, appetite, and sleep [71]. Pathologically, it is implicated as a potential starting point for diseases like Parkinson's. In Parkinson's misfolded alpha-synuclein protein is believed to first aggregate in ENS neurons before propagating to the brain via the vagus nerve [72, 73]. Furthermore, ENS integrity is crucial for maintaining the gut barrier. Dysfunction of gut contributes to "leaky gut," inflammation, and the subsequent systemic immune responses leading to several neuropsychiatric conditions [49, 74]. Importantly, the ENS is a central player in translating gut microbiome activity into signals that profoundly impact the brain [75].

Critical windows of development: Early-life microbiome and neurodevelopment

The gut microbiome is established during early life represents a critical developmental window. The gut microbiome exerts a long-lasting influence on neurological and psychiatric well-being. The prenatal, perinatal, and early postnatal periods are characterized by rapid brain development and concurrent microbial colonization. It creates a vulnerable yet programmable interface where gut-brain communication is first established [10, 76]. Disruptions during this sensitive period can have profound and enduring consequences on brain structure, function, and an individual's vulnerability to disorders later in life [77, 78]. The initial microbial

Table 1. Neurodegenerative & psychiatric diseases and the microbiome.

Disease	Associated microbiome (Dysbiosis)	Hypothesized role of the microbiome	References
Alzheimer's Disease (AD)	<ul style="list-style-type: none"> Reduced diversity. Decreased SCFA-producers (e.g., <i>Faecalibacterium</i>). Increased pro-inflammatory taxa & LPS-producers. 	<ul style="list-style-type: none"> LPS / Inflammation: Gut LPS → systemic inflammation → microglial activation → neuroinflammation. Amyloid Seeding: Bacterial amyloids → cross-seeding of Aβ plaques. BBB Breakdown: SCFA deficiency → impaired tight junctions → compromised BBB. 	[56, 57]
Parkinson's Disease (PD)	<ul style="list-style-type: none"> Reduced diversity. Decreased <i>Prevotellaceae</i>. Increased <i>Enterobacteriaceae</i>. 	<ul style="list-style-type: none"> Pathogen Propagation: Gut inflammation → α-syn misfolding in gut → vagus nerve propagation to brain. Metabolic Toxicity: Microbial metabolites → mitochondrial dysfunction → oxidative stress. Drug Metabolism: Gut bacteria metabolize levodopa → reduced drug efficacy. 	[58, 59]
Autism Spectrum Disorder (ASD)	<ul style="list-style-type: none"> Reduced diversity. Increased <i>Clostridium</i>, <i>Desulfovibrio</i>. Decreased <i>Bifidobacterium</i>. 	<ul style="list-style-type: none"> Toxic Metabolites: Bacterial propionic acid → BBB crossing → neuroinflammation & mitochondrial dysfunction. Leaky Gut: Intestinal permeability → neuroactive peptides enter blood → immune activation. Neurotransmitter Imbalance: Dysbiosis → altered GABA/glutamate precursor synthesis. 	[60, 61]
Major Depressive Disorder (MDD)	<ul style="list-style-type: none"> Reduced diversity. Decreased <i>Faecalibacterium</i>, <i>Coprococcus</i>. Increased pro-inflammatory genera. 	<ul style="list-style-type: none"> Cytokine Release: Pro-inflammatory microbes → IL-6, TNF-α release → HPA axis overactivation & reduced neurogenesis. Tryptophan Diversion: Inflammation → shunts tryptophan to kynurenone → away from serotonin. SCFA Deficiency: Low butyrate → impaired microglial function → poor neuronal support. 	[62, 63]
Anxiety Disorders	<ul style="list-style-type: none"> Reduced diversity. Decreased <i>Lactobacillus</i>, <i>Bifidobacterium</i>. Increased <i>Enterobacteriaceae</i>. 	<ul style="list-style-type: none"> Vagal GABA Modulation: Specific microbes → vagus nerve signaling → altered GABA receptor expression in amygdala. HPA Axis Dysregulation: Dysbiosis → exaggerated corticosterone response to stress. Direct Synthesis: Gut bacteria produce GABA → influences central nervous tone. 	
Multiple Sclerosis (MS)	<ul style="list-style-type: none"> Reduction in SCFA-producing <i>Clostridia</i>. Increase in <i>Akkermansia</i>, <i>Methanobrevibacter</i>. 	<ul style="list-style-type: none"> Immune Cell Deficit: Low SCFAs → impaired Treg cell generation → loss of autoimmunity suppression. Molecular Mimicry: Microbial peptides → mimic myelin antigen → trigger cross-reactive T cell attack. Bile Acid Alteration: Dysbiosis → altered bile acid metabolism → favors pro-inflammatory Th17 cells. 	[64, 65]

inoculum is shaped by several factors such as mode of delivery (vaginal birth versus Caesarean section) and feeding practices of infants (breastfeeding versus formula). It dictates the pioneer species that colonize the sterile gut. These early colonists are not passive inhabitants; they play an instructive role in educating the developing immune system and are essential for the normal maturation of key neural systems [79, 80]. Crucially, the microbiota guides the development of the blood-brain barrier, ensuring its integrity, and regulates the maturation and function of microglia the brain's resident immune cells [14, 81]. Properly functioning microglia are vital for synaptic pruning, the process of refining neural connections that is fundamental for learning and cognitive function [82, 83].

Furthermore, the early-life microbiome is integral to the action of the HPA axis, the body's central stress response system [84]. Importantly, a healthy, diverse gut microbiota helps to establish appropriate stress reactivity and resilience [85, 86]. Conversely, early-life adversity, antibiotic use, or pathogenic infections can cause dysbiosis, leading to microglial dysfunction, an exaggerated HPA axis response, and altered neurodevelopment [87, 88]. The etiology of neurodevelopmental disorders such as ASD and ADHD were initiated due to the dysregulation of microbiota. It highlights that the seeds of future neurological and psychiatric health are sown within the first years of life [89, 90].

The impact of common medications on the GBA

The profound influence of commonly prescribed medications on the GBA is an emerging area. It was revealed that their therapeutic and side effects may be partially mediated through unintended alterations to the gut microbiome [32, 91]. Treating specific conditions, many widely used drugs exerts significant off-target effects that disrupt microbial communities, compromise intestinal barrier integrity. Subsequently, modulate neuroinflammatory pathways, thereby influencing neurological and psychiatric health [92, 93]. Antibiotics designed to kill or inhibit bacteria, [94] while life-saving, their broad-spectrum action can severely reduce microbial diversity for long time, killing beneficial commensal species [95]. Additionally, this dysbiosis induces reduction of the production of crucial neuroactive metabolites like SCFAs, potentially increasing vulnerability to stress, anxiety, and cognitive disturbances, particularly when used during critical developmental windows [19, 96, 97].

In addition, non-antibiotic drugs also exhibit significant antimicrobial actions [98]. It was found to gut microbial populations were altered dramatically due to the action of Proton-pump inhibitors (PPIs), metformin, and non-steroidal anti-inflammatory medications (NSAIDs) [99]. PPIs alter pH of the gut, which can lead to the overgrowth of orally-derived bacteria and potentially increasing infection risk [100, 101]. Importantly, many psychoactive medications, including antidepressants and antipsychotics, demonstrate antimicrobial properties *in vitro* [102, 103]. A compelling hypothesis suggests that a part of their potent efficacy may stem not from direct action on neurons. It indirectly through modifying the gut microbiome and its production of neurotransmitters like serotonin or GABA, thereby influencing the GBA [1, 32, 104]. Herein, we highlights a paradigm shift: the gut microbiome must be considered a primary mediator of both the therapeutic benefits and adverse effects of pharmacological interventions [105, 106]. Therefore, understanding these interactions is essential for predicting side effects, elucidating individual variability in drug response, and developing novel strategies that combine traditional therapeutics with microbiome-enhancing co-treatments to enhance patient outcomes in neurological and psychiatric care (**Table 2**) [107-120].

Individual variability and the concept of "microbiome fingerprinting"

The factors, such as a complex interplay of genetics, geography, diet, early-life exposures, and medication history potentially influence the gut microbial composition. It indicates that each individual exhibits a unique microbial profile which is fully different from other individuals, called "microbiome fingerprint" [121]. Importantly, this concept fundamentally shifts the therapeutic paradigm from a one-size-fits-all approach towards a future of personalized, precision medicine targeting the GBA for neurological and psychiatric well-being [122]. The generalized dietary or probiotic interventions are the key reason for this individual variability leading to inconsistent results across populations. Depending on the unique microbial community structure and host genetics, a microbial species or metabolite that is beneficial for one person may be neutral or even detrimental to another. For instance, the efficacy of a probiotic strain in alleviating anxiety symptoms may depend entirely on the recipient's baseline microbiota's ability to support its colonization or on their specific immune response to its introduction [123, 124]. This explains the high degree of heterogeneity observed in clinical trials of prebiotics and psychobiotics.

Consequently, advanced sequencing and machine learning algorithms decode microbiome fingerprint of each individual which is the future of GBA-based therapeutics [125]. The goal is to move beyond simple taxonomic analysis to a functional understanding of microbial metabolic potential of each individual. This could enable clinicians to predict disease susceptibility, tailor nutritional plans to boost specific neuroprotective metabolites, and select microbial consortia (next-generation probiotics) designed to correct a patient's specific functional deficits [126]. By acknowledging and embracing individual variability through microbiome fingerprinting, the field can progress towards developing truly effective, personalized interventions that modulate the GBA to enhance neurological resilience and psychiatric health.

Beyond bacteria: the virome and mycobiome

The traditional focus of GBA research has focused on bacteria. However, the gut microbiome is a complex ecosystem comprising other microbial diversity, notably viruses and fungi, which are now recognized as essential contributors to neurological and psychiatric health. Thus, the collective genomes of viruses the virome and fungi the mycobiome interact with bacterial communities and the host, adding profound layers of complexity to gut-brain signaling and opening new frontiers for understanding disease mechanisms [127, 128]. Gut bacteria are infected by the human gut virome is primarily comprised of bacteriophages, viruses [129, 130]. Furthermore, pathogens, phages are key regulators of bacterial population dynamics, diversity, and function through a predator-prey relationship [131]. The gut ecosystem's structure is rapidly altered by phages through lysing specific bacterial hosts or transferring genetic material through transduction, leading to influencing the production of bacterial metabolites like SCFAs [132]. Therefore, bacterial dysbiosis are mediated by dysbiosis in the phage community. It potentially contributes to the neuroinflammatory processes leading to disorders like AD and major depression [133].

Similarly, the mycobiome which is comprised of commensal fungi like *Candida* and *Saccharomyces* species, maintain balance with bacteria and the host immune system [134]. Fungi are less abundant than bacteria but potent regulators of immune responses. An overgrowth of certain fungal species can disrupt intestinal barrier integrity and amplify systemic immune activation,

Table 2. Mechanisms, messengers, and effects in the gut-brain axis.

Component	Key elements	Primary function in GBA	Effect on neurological & psychiatric health	References
Primary Communication Pathways	Neural (Vagus Nerve): Main physical conduit for bidirectional signals. Endocrine: Gut hormones (serotonin, PYY). Immune: Cytokines (IL-1 β , IL-6, TNF- α). Metabolic: Microbial metabolites (SCFAs).	Transmits information between the gut and brain. Neural is direct and fast; endocrine/immune are slower, systemic; metabolic is a foundational biochemical influence.	Ensures integration of gut homeostasis with brain function. Dysregulation in any pathway is a mechanism for disease.	[107, 108]
Key Microbial Metabolites	Short-Chain Fatty Acids (SCFAs): Butyrate, Propionate, Acetate. Neurotransmitters: GABA, Serotonin, Dopamine. Tryptophan Metabolites: Kynurenone, Quinolinic acid. Bile Acids: Secondary bile acids.	SCFAs: Anti-inflammatory, energy for colonocytes, strengthen BBB. Neurotransmitters: Directly influence host neurotransmission. Tryptophan Metabolism: Balances neuroprotective vs. neurotoxic compounds.	Protective (Eubiosis): SCFAs and neuroprotective kynurenic acid support brain health. Detrimental (Dysbiosis): Neurotoxic quinolinic acid and neurotransmitter imbalance promote disease.	[109, 110]
Gut Barrier Integrity	Intestinal Epithelium: Single layer of cells with tight junctions. Mucus Layer: Physical barrier. Immune Cells: Monitor barrier status.	Prevents translocation of bacteria and pro-inflammatory molecules (e.g., LPS) into systemic circulation.	Intact Barrier ("Tight Gut"): Prevents systemic inflammation and neuroinflammation. Compromised Barrier ("Leaky Gut"): Allows LPS & cytokines to trigger neuroinflammation, linked to depression, AD, and anxiety.	[111, 112]
Central Nervous System Barriers	Blood-Brain Barrier (BBB): Semi-permeable lining of brain's capillaries. Glymphatic System: Clears waste from brain.	Protects the brain from toxins and pathogens in the blood while allowing nutrient passage.	Strong BBB: Maintains a stable environment for neuronal function. Weakened BBB: Permits influx of inflammatory cytokines and immune cells, driving neuroinflammation and neurodegeneration.	[113, 114]
Immune System Regulation	Gut-Associated Lymphoid Tissue (GALT): Body's largest immune organ. Microglia: Brain's resident immune cells.	The microbiome educates the immune system. Gut immune responses can activate microglia in the brain.	Healthy Regulation: Prevents excessive inflammation; supports synaptic pruning via microglia. Dysregulation: Chronic systemic inflammation leads to microglial activation, neuroinflammation, and impaired neuroplasticity.	[115, 116]

Component	Key elements	Primary function in GBA	Effect on neurological & psychiatric health	References
Stress Response System	Hypothalamic-Pituitary-Adrenal (HPA) Axis	The body's central stress response system. Releases cortisol.	Well-Regulated: The microbiome helps calibrate the HPA axis for a balanced stress response. Dysregulated (Dysbiosis): Leads to HPA axis hyperactivity, exaggerated cortisol release, and increased vulnerability to anxiety and depression.	[117, 118]
State of the Gut Ecosystem	Eubiosis: High diversity, balanced community, beneficial microbes dominate. Dysbiosis: Low diversity, loss of beneficials, overgrowth of pathogens.	Eubiosis supports all healthy GBA functions. Dysbiosis disrupts signaling, barrier function, and immune regulation.	Eubiosis: Foundation of neurological homeostasis and resilience. Dysbiosis: A key disruptor of brain homeostasis and a driver of neuropsychiatric disorders.	[119, 120]

resulting the risk factor for neuroinflammation [14]. Furthermore, bioactive metabolites and mycotoxins are produced by some fungi that may directly influence neuronal function [135]. Focusing these interactions is crucial for a holistic understanding of how gut microbial dysbiosis impacts the brain.

Circulating metabolites as key messengers: A molecular perspective

By production and modulation of a vast array of circulating metabolites, the gut microbiome significantly influences the brain functions [136]. These small molecules produced by microbiome, taken into the circulation from the intestines are crucial messengers for the endocrine system. It makes a direct molecular connection between brain function and microbial metabolic activity. This biochemical interaction is a fundamental mechanism of the gut-brain axis acts to maintain neurological and psychiatric well-being. It is a tangible pathway through which diet and microbiota can shape mental health. These microbial metabolites are SCFAs, such as butyrate, propionate, and acetate, produced from the fermentation of dietary fiber [137]. Butyrate, in particular, demonstrates potent neuroprotective properties. It passes the blood-brain barrier, lowers neuroinflammation by blocking histone deacetylases (HDACs), and serves as a main energy source for colonocytes, supporting overall gut health [138]. Beyond SCFAs, gut bacteria play important role in bile acid metabolism. Gut microbiome alters primary bile acids from the liver into secondary bile acids. It acts as signaling molecules through receptors like FXR and TGR5, reshaping glial cell activity and neuroinflammation [139].

Moreover, the essential amino acid tryptophan is associated with the most critical metabolic pathway. Gut microbes directly consume dietary tryptophan, but also critically regulate its host metabolism. It is very important to maintain equilibrium between the serotonin and kynurenine pathways. Microbial dysbiosis can move tryptophan away from producing serotonin, a key neurotransmitter, towards the kynurenine pathway, leading to formation of neuroactive metabolites that can be either neuroprotective (kynurenic acid) or neurotoxic (quinolinic acid) [140]. A shift towards neurotoxicity is strongly implicated in the pathogenesis of depression, anxiety, and neurodegenerative diseases. Thus, circulating metabolites provide a molecular basis for how gut microbial composition directly influences brain chemistry and vulnerability to disorder.

Eubiosis: Foundation of neurological homeostasis

A state of eubiosis is characterized by a diverse and functionally balanced gut microbiota [141]. Eubiosis has major contribution on brain homeostasis and supporting neurological and psychiatric well-being. Thus, a resilient and stable microbial community directly and indirectly promote central nervous system functions. Beneficial commensal gut microbiota crucially performs fermenting dietary fiber to generate key neuroactive essential metabolites. These metabolites possess systemic anti-inflammatory effects, reinforce the integrity of both the gut and blood-brain barriers, and provide an energy source for colonocytes. Thereby, these metabolites support overall gastrointestinal health. Furthermore, a eubiotic microbiome contributes to the regulation of the extensive endocrine system of gut. It affects the production and secretion of neurotransmitters and gut hormones [142]. It also plays an important role in educating and modulating the host immune system, ensuring appropriate inflammatory responses and preventing systemic immune activation. This harmonious relationship between the host and its microbial inhabitants establishes a foundation for optimal gut-brain communication,

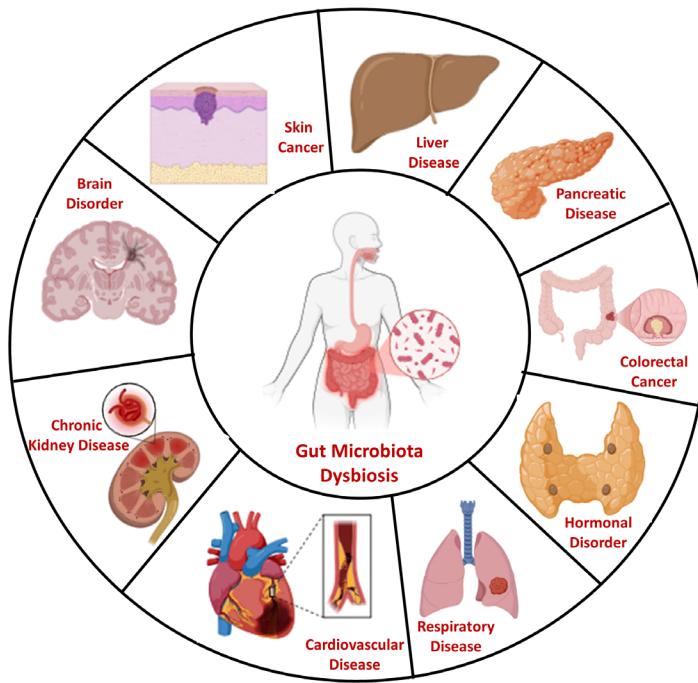


Figure 3. The Systemic reach of the gut-brain axis and connections to peripheral organ systems. This schematic illustrates how gut microbiota dysbiosis and a compromised gut-brain axis contribute to systemic pathophysiology beyond the brain. Through mechanisms of chronic systemic inflammation, immune dysregulation, and altered microbial metabolite production, gut-derived signals can disrupt the function of peripheral organs. This establishes mechanistic links between an imbalanced gut ecosystem and the development or progression of a spectrum of conditions, including cardiovascular disease, chronic kidney disease, respiratory disease, liver disease, pancreatic disease, colorectal cancer, skin cancer, hormonal disorder, and brain disorder. This holistic view underscores the gut's role as a central modulator of whole-body health.

facilitating healthy neurodevelopment, synaptic plasticity, stress resilience, and cognitive function.

The healthy gut-brain ecosystem

Eubiosis serves as the essential foundation for normal brain function and psychological well-being. Therefore, in the symbiotic equilibrium state, the complex community of microorganisms performs essential functions that actively support central nervous system functions through multiple interconnected mechanisms. Furthermore, a healthy gut microbiome promotes production of neurotransmitters including serotonin, GABA, and dopamine. Besides, gut microbiome simultaneously supports the integrity of the intestinal epithelial barrier to prevent the translocation of pro-inflammatory substances [143]. This host-microbe symbiotic relationship facilitates appropriate immune system regulation and balanced stress response through HPA axis alteration [144]. Collectively, Efficient nutrient absorption maintains the delicate biochemical environment necessary for neural plasticity, cognitive function, and emotional regulation. Therefore, the preservation of this intricate ecological balance is required for neurological homeostasis, and its disruption represents a critical pathway through which brain well-being becomes compromised.

Moreover, the systemic influence of the gut-brain axis, demonstrating how gut microbiota dysbiosis manifests pathophysiology to peripheral organs. Driven by chronic inflammation, immune dysregulation, and aberrant microbial metabolite signaling, gut-derived alteration is mechanistically associated with a spectrum of disorders, such as, cardiovascular, renal, hepatic, pancreatic, and respiratory diseases [145]. This perspective positions the gut as a pivotal regulator of holistic

organismal health (Figure 3).

Dysbiosis of human intestinal microbiota and brain homeostasis

Dysbiosis severely disrupts brain homeostasis, resulting onset of neurological and psychiatric diseases. This pathological alteration in the composition and functionality of the gut microbiota characterized by reduced biodiversity, loss of beneficial gut microbiota, and overgrowth of opportunistic pathogens [146]. As a result, signaling pathways are disrupted and this disruption potentially destabilizes microbial population, resulting impairment of brain well-being and neural function.

The potential impact of dysbiosis on brain homeostasis occurs through several interconnected mechanism. Firstly, it diminishes the intestinal epithelial barrier integrity, leading to enhancement of gut permeability. Subsequently, bacterial endotoxins, LPS enters into blood stream through gut leakage which triggers inflammation [147]. Then, onset of neuroinflammation, AD, and anxiety disorders caused by pro-inflammatory cytokines which can pass through the blood-brain barrier. Secondly, alteration of essential microbial metabolites production due to dysbiosis. Reduced production of beneficial SCFAs such as butyrate attenuates their neuroprotective, anti-inflammatory, and blood-brain barrier enhancing effects [148]. Concurrently, dysbiosis can shift tryptophan metabolism away from serotonin synthesis towards the neurotoxic kynurene pathway, further exacerbating neurological dysfunction [149]. Furthermore, dysbiosis disrupts the regulation of the HPA axis, leading to aberrant stress responses and elevated cortisol levels that negatively influence brain regions like the hippocampus and prefrontal cortex [150]. By disruption of

these vital communication networks, intestinal dysbiosis mediates synaptic plasticity, neurotransmitter balance, and glial cell function (Figure 4).

Gut microbiome on insomnia and schizophrenia

Emerging evidence reveals critical connections between gut microbiome composition and two disorders: insomnia and schizophrenia [151]. In both disorders, specific microbial alterations contribute to pathophysiology through gut-brain axis mechanisms. Subsequently, onset of immune activation, neurotransmitter production, and sleep-wake cycle regulation. Sleep pattern and quality of the sleep are altered through multiple pathways by gut microbiome diversity. Certain *Lactobacillus* and *Bifidobacterium* strains enhance GABA production, promoting relaxation and sleep initiation [152]. Conversely, microbial dysbiosis impairs normal sleep patterns and reduce sleep quality through increased formation of pro-inflammatory cytokines. In addition, the microbiome controls circadian rhythms through metabolic products that influence central clock gene expression in the hypothalamus leading alteration of sleep pattern.

In schizophrenia, microbiome alterations have been characterized by reduced microbial diversity and higher expression of pathogenic species. Alteration of microbial population is associated with onset of disease by multiple mechanisms [153]. Firstly, triggering neuroinflammation by elevated intestinal permeability allowing bacterial metabolites. Secondly, altered tryptophan metabolism and impaired dopamine and glutamate signaling. Importantly, microbial composition is affected by antipsychotic medications, that may influence therapeutic efficacy and side effects [154]. Therefore, probiotic supplementation

may improve both gastrointestinal symptoms and psychological wellbeing in schizophrenia patients, possibly by reducing inflammatory markers and oxidative stress (Figure 4).

Probiotics on CNS and neurological disorders

It has been revealed that specific probiotic strains, often termed "psychobiotics", can significantly improve CNS function and potentially alleviate various neurological disorders [155]. Thus, this beneficial gut microbiota modulates neurochemical, inflammatory, and endocrine signaling. Additionally, the production of GABA, serotonin, and brain-derived neurotrophic factor (BDNF) are raised by the action probiotics which are crucial for neuronal health, synaptic plasticity, and mood regulation [156]. It was unveiled that certain *Lactobacillus* and *Bifidobacterium* strains in AD potentially reduced amyloid-beta aggregation and tau phosphorylation through anti-inflammatory mechanisms and enhanced production of neuroprotective butyrate [157]. Moreover, probiotics may improve gastrointestinal disorders by reducing systemic inflammation and alpha-synuclein aggregation in Parkinson's disease. Specific probiotic formulations have demonstrated remarkable immunomodulatory effects in multiple sclerosis.

The meta-analyses report indicate that depression and anxiety significantly reduced by the action of particular probiotic combinations with use of anti-depressant medications [158]. Probiotics mediate effect through reduced inflammatory cytokines, normalized HPA axis activity, and elevated tryptophan availability for serotonin synthesis. Further research is needed to establish optimal strains, dosages, and treatment durations for specific neurological conditions, paving the way for more targeted

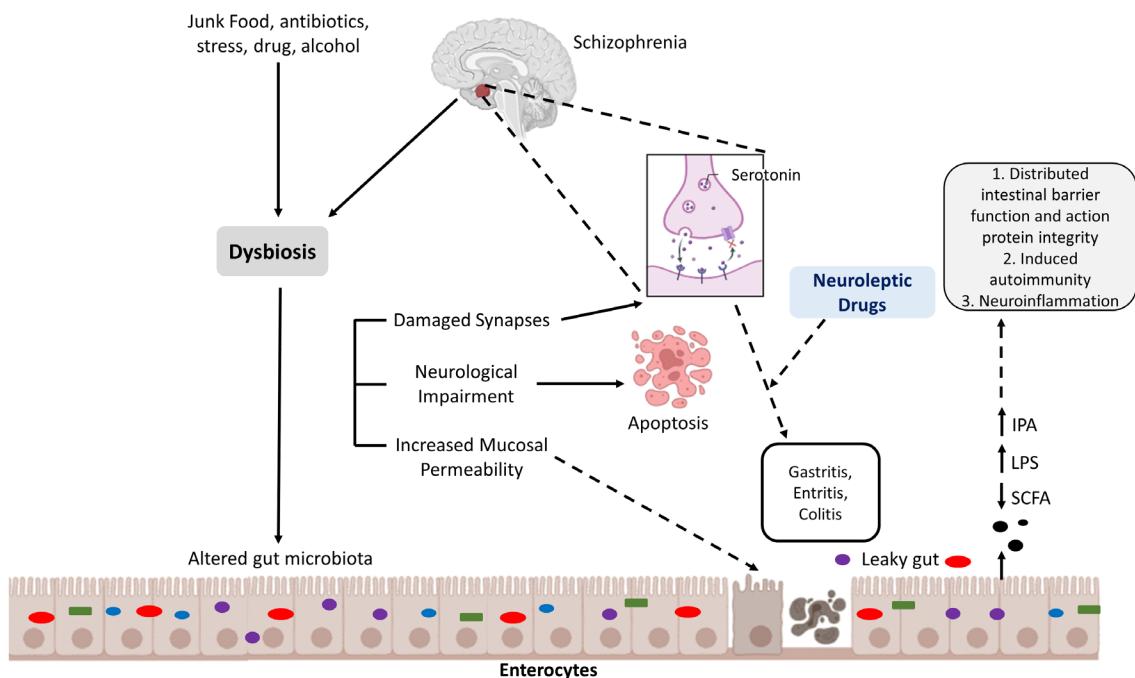


Figure 4. The vicious cycle of gut-brain dysregulation in schizophrenia pathophysiology. This schematic illustrates a proposed pathological loop linking environmental triggers, gut integrity, and brain function in Schizophrenia. Exogenous factors like junk food, stress, alcohol, antibiotics, and drug use initiate altered gut microbiota and dysbiosis. This disrupts the gut ecosystem, reducing beneficial metabolites like SCFA and IPA, while increasing harmful LPS. Consequently, increased mucosal permeability and a leaky gut cause inflammation (gastritis, enteritis, colitis) and enterocyte damage, leading to distributed intestinal barrier function and action protein integrity. Bacterial products translocate into circulation, driving systemic inflammation and neuroinflammation, which contributes to apoptosis, damaged synapses, and neurological impairment. This cycle may be further exacerbated by, or influence the efficacy of, neuroleptic drugs.

microbial-based interventions in neurology and psychiatry.

Therapeutic interventions and future directions

The growing interest of the GBA has increased the exploration of novel therapeutic strategies aimed to improve neurological and psychiatric well-being. These interventions target the gut microbiome and its outputs, offer a promising neurological and psychiatric treatments for the patient care. Current research focuses primarily on dietary modifications, prebiotics, and probiotics. Specific diets, particularly those high in fiber and polyphenols like the Mediterranean diet, are shown to increase microbial diversity and the production of beneficial metabolites, such as SCFAs, which confer neuroprotective and anti-inflammatory effects [159]. Targeted psychobiotics live organisms that produce health benefits in patients with psychiatric illnesses are being investigated for their potential to alleviate symptoms of depression and anxiety [160].

Importantly, future studies should be done to prioritize large-scale, longitudinal human trials to establish causal relationships between microbial changes and clinical outcomes. By integrating an individual's unique microbial fingerprint, genetic background, and lifestyle factors, therapies can be tailored for maximum efficacy. This may involve designing specific synbiotic (combined prebiotic and probiotic) formulations or using phage therapy to precisely modulate bacterial populations. Collectively, harnessing the GBA therapeutically holds immense potential to revolutionize treatment paradigms, offering new hope for preventing and managing a spectrum of neurological and psychiatric disorders.

Conclusions

In conclusion, this review establishes the GBA as fundamental to neurological and psychiatric well-being. The complex bidirectional communication between the gastrointestinal tract and central nervous system through interconnected multiple pathways reveals that brain health cannot be completely explored separately from peripheral influences. To understand fully the brain health, gut microbiome also needs to be focused equally. This paradigm challenges traditional neurocentric views and underscores the need for a comprehensive, integrative approach to patient care. Future therapeutic approaches should extend beyond targeting the brain alone to include gut ecosystem modulation. Nutritional psychiatry, personalized microbial regimens, and medication mindfulness emerge as promising adjunctive treatments. This perspective emphasizes early-life interventions and lifestyle factors in establishing neurological resilience. Collectively, advancing GBA understanding requires greater interdisciplinary collaboration integrating neurology, psychiatry, gastroenterology, and nutrition science.

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No applicable.

Data availability

This narrative review is based on previously published studies and publicly available data. No new datasets were generated or analyzed for the current review.

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MS contributed to the design, writing, collected data and drew figures for the manuscript. MAK revised the manuscript and approved the submission.

Competing interests

The authors declare no competing interests.

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