

# Fecal microbiota transplantation and next-generation therapies: A review on targeting dysbiosis in metabolic disorders and beyond

SAGE Open Medicine

Volume 12: 1–12

© The Author(s) 2024

Article reuse guidelines:

sagepub.com/journals-permissions

DOI: 10.1177/20503121241257486

journals.sagepub.com/home/smo



Zenawork Sahle<sup>1</sup> , Getabalew Engidaye<sup>1</sup>, Demissew Shenkute Gebreyes<sup>1</sup> , Behailu Adenew<sup>2</sup> and Tsegahun Asfaw Abebe<sup>1</sup> 

## Abstract

The human microbiome, particularly the gut microbiome, has emerged as a central determinant of health and disease. Dysbiosis, an imbalance in the microbial composition of the gut, is associated with a variety of metabolic and other diseases, highlighting the potential for microbiota-targeted treatments. Fecal microbiota transplantation has received considerable attention as a promising therapy to modulate the gut microbiome and restore microbial homeostasis. However, challenges remain, including standardization, safety, and long-term efficacy. This review summarizes current knowledge on fecal microbiota transplantation and describes the next generation therapies targeting microbiome. This review looked at the mechanistic understanding of fecal microbiota transplantation and alternative strategies, elucidating their potential role in improving dysbiosis-associated metabolic disorders, such as obesity, and type 2 diabetes and others. Additionally, this review discussed the growing application of therapies targeting the gut microbiome. Insights from clinical trials, preclinical studies, and emerging technologies provide a comprehensive overview of the evolving landscape of microbiome-based interventions. Through a critical assessment of current advances and prospects, this review aims to highlight the therapeutic potential of targeting gut microbiome and pave the way for innovative approaches in precision medicine and personalized treatments.

## Keywords

Fecal microbiota transplantation, gut microbiota, metabolic disorders, microbiota-targeted therapies, dysbiosis

Date received: 15 September 2023; accepted: 9 May 2024

## Introduction

The human gut microbiota, a diverse ecosystem of microorganisms, influences host physiology, metabolism, and immune function.<sup>1,2</sup> Dysbiosis, or disruptions in microbial composition and function, is associated with various diseases.<sup>2</sup> Metabolic disorders like obesity, diabetes, and metabolic syndrome are a major global health issue, causing illness, death, and high healthcare expenses.<sup>3</sup> Emerging evidence indicates that imbalanced gut microbiota may also play a role in causing these conditions.<sup>4</sup> Therefore, there is a growing interest in microbiota-targeted therapies to modulate microbial communities and restore homeostasis in metabolic disorders.<sup>5–9</sup> The microbes in the digestive tract are called the human gastrointestinal microbiota. They outnumber human cells by 10 times, with an average ratio of 1:3 microbiota to human cells.<sup>10</sup> FMT is a potential intervention to reshape the gut microbiota and treat metabolic dysfunction and autoimmune disease.<sup>11,12</sup> It involves transferring

fecal microbial communities from a healthy donor to restore diversity and function. Initially used mainly for recurrent *Clostridiodes difficile* infection, FMT has now also been applied to other conditions, including metabolic disorders.<sup>13</sup>

Despite promising results several studies,<sup>13</sup> challenges remain, including standardization of protocols, safety concerns, and variability in long-term effectiveness.<sup>14–17</sup> Additionally, the mechanisms underlying the therapeutic effects of FMT and the optimal selection of donors and

<sup>1</sup>Department of Medical Laboratory Science, Asrat Weldeyes Health Science Campus, Debre Berhan University, Debre Berhan, Ethiopia

<sup>2</sup>Department of Medical Laboratory Science, Debre Berhan Compressive Specialized Hospital, Debre Berhan, Ethiopia

### Corresponding author:

Zenawork Sahle, Department of Medical Laboratory Science, Asrat Weldeyes Health Science Campus, Debre Berhan University, Debre Berhan 445, Ethiopia.

Email: zenasahle@gmail.com



Creative Commons Non Commercial CC BY-NC: This article is distributed under the terms of the Creative Commons

Attribution-NonCommercial 4.0 License (<https://creativecommons.org/licenses/by-nc/4.0/>) which permits non-commercial use, reproduction and distribution of the work without further permission provided the original work is attributed as specified on the SAGE and Open Access pages (<https://us.sagepub.com/en-us/nam/open-access-at-sage>).

recipient are areas of ongoing investigation.<sup>15,18</sup> In addition to FMT, a variety of next-generation therapies targeting microbiome has emerged, including prebiotics, probiotics, postbiotics, antibiotics, and microbial therapies.<sup>19</sup> These approaches aim to modulate the gut microbiota through various mechanisms, including promoting the growth of beneficial bacteria, inhibiting pathogens, and modulating host-microbiota interactions. This review aims to provide a comprehensive overview of the current landscape of FMT and next-generation microbiota-targeted therapies to target physiological dysregulation in metabolic disorders and beyond. Through a critical appraisal of the existing evidence, mechanistic insights, and clinical applications, we seek to elucidate the therapeutic potential of these interventions and identify future directions for research and future clinical application in the growing field of microbiome-based therapies.

## Pathophysiological justification for fecal microbiota transplantation

Gut microbiota encompasses the set of microorganisms that colonize the gastrointestinal tract with mutual relationships that are key for host homeostasis.<sup>20</sup> In the first year of life, the gut microbiome is developed and is impacted by a variety of internal and environmental variables, including nutrition and antibiotic use.<sup>21</sup> Low diversity and relative dominance of *Proteobacteria* and *Actinobacteria* define the gut microbiota of neonates.<sup>22</sup> The adult microbiota, which is characterized by *Firmicutes* and *Bacteroidetes* dominance, then causes the microbiota to become more varied.<sup>23</sup> The host may benefit from the metabolites that the microbiota creates, including those that have anti-inflammatory and antioxidant activity, regulate intestinal barrier function, and provide vitamins and energy.<sup>24</sup> Human intestine commensal bacteria play a role in the growth and maintenance of gut sensory, motor, and immunologic functions.<sup>25</sup>

Fecal microbiota transplantation has been used in several clinical investigations where metabolic syndrome like obesity,<sup>26</sup> and non-alcoholic fatty liver disease<sup>27</sup> have been present. According to previous study, several clinical investigations using FMT have been carried out in the context of metabolic syndrome, obesity, and non-alcoholic fatty liver disease in both rats and humans. Patients in those studies had lower levels of gut microbial diversity, and after receiving FMT from lean, healthy donors, the gut microbial diversity levels were increased.<sup>28</sup>

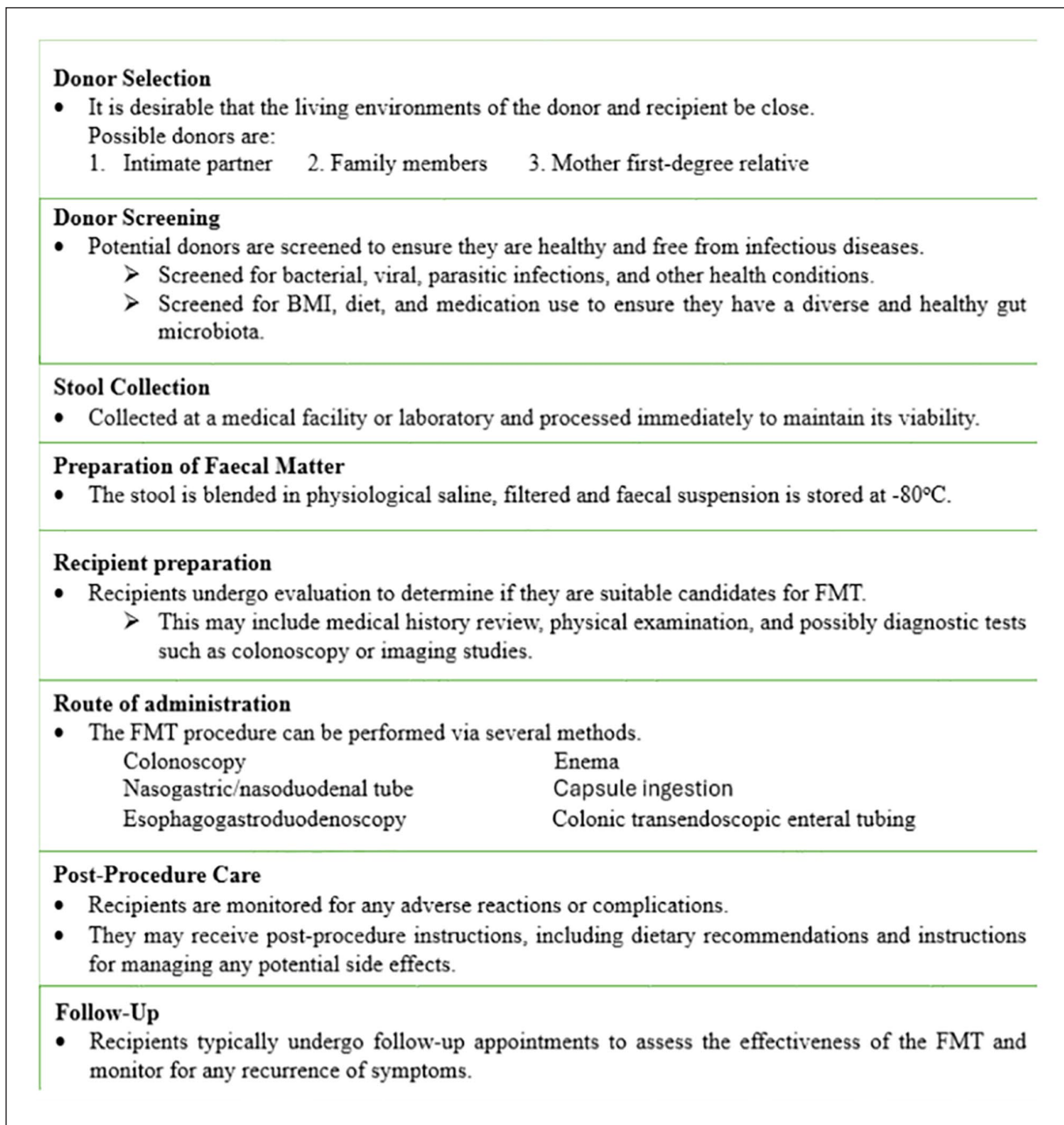
There's a growing recognition of the complex relationship between a host and its microbiota across different species.<sup>29</sup> Recent progress in identifying and isolating specific members of the gut microbiota, advancements in gnotobiology and a deeper understanding of host genetics have collectively facilitated research in this field.<sup>30</sup> This has enabled researchers to delve deeper into the dynamics of this relationship and its implications for health and disease. The gut

microbiota is indispensable for the host's overall health, influencing various aspects of host biology. It actively participates in the development and differentiation of the intestinal epithelium and the immune system,<sup>31</sup> crucial for maintaining tissue homeostasis and defending against pathogen invasion.<sup>32,33</sup> Additionally, it plays a pivotal role in metabolizing indigestible polysaccharides, providing essential nutrients and energy sources to the host.<sup>34–36</sup> Furthermore, certain members of the gut microbiota contribute to the production of vital vitamins necessary for various physiological processes.<sup>37,38</sup> This intricate symbiotic relationship underscores the importance of a balanced and diverse gut microbiota in promoting optimal health and wellbeing.

The gut microbiota actively regulates numerous metabolic pathways in the host, fostering intricate metabolic, signaling, and immune-inflammatory axes that connect the gut, liver, muscle, and brain.<sup>39</sup> A thorough comprehension of these axes is essential for refining therapeutic approaches aimed at modulating the gut microbiota to address diseases and enhance overall health. Organ morphogenesis, intestinal vascularization, tissue homeostasis, carcinogenesis, bone mass, and behavior are just a few of the physiological aspects of the host that the gut microbiota influences.<sup>20</sup> It also expands nutrient sources, produces essential vitamins, and carries out xenobiotic metabolism.<sup>37</sup>

## Fecal transplantation procedure and selection

fecal transplantation has emerged as a promising therapeutic option for restoring gut microbiota balance and treating certain gastrointestinal conditions, with ongoing research focused on optimizing the procedure and selecting appropriate candidates for treatment. Even though the preparation of stool sample was varied, it is feasible to reestablish a healthy intestinal microbiome by transferring fecal material from a healthy donor to the patient to increase the intestinal microbial variety.<sup>40</sup> Fecal transplantation was first carried out on people in 1958, and it has been done on animals for more than a century.<sup>41</sup> The selection of donors for FMT is crucial and typically involves screening for infectious diseases and medical conditions to ensure the safety of the procedure.<sup>42</sup> Donors are often selected based on strict criteria to minimize the risk of transmitting pathogens or other adverse effects to the recipient.<sup>43</sup> Whether the donor is a close friend or family member, a first-degree relative, or even a stranger, it makes no difference. Clinicians must choose a donor for fecal transplantation who is free of infectious pathogens that could be transferred to the recipient. If a potential donor has been exposed to hepatitis B, hepatitis C, or HIV viruses within the past year or has a confirmed diagnosis of one of these conditions, they are disqualified from participating.<sup>44</sup> The donation criteria also disallow anyone with a tattoo or body piercing, as well as those who engage in risky sexual activity or use illegal substances<sup>44</sup> (Figure 1).



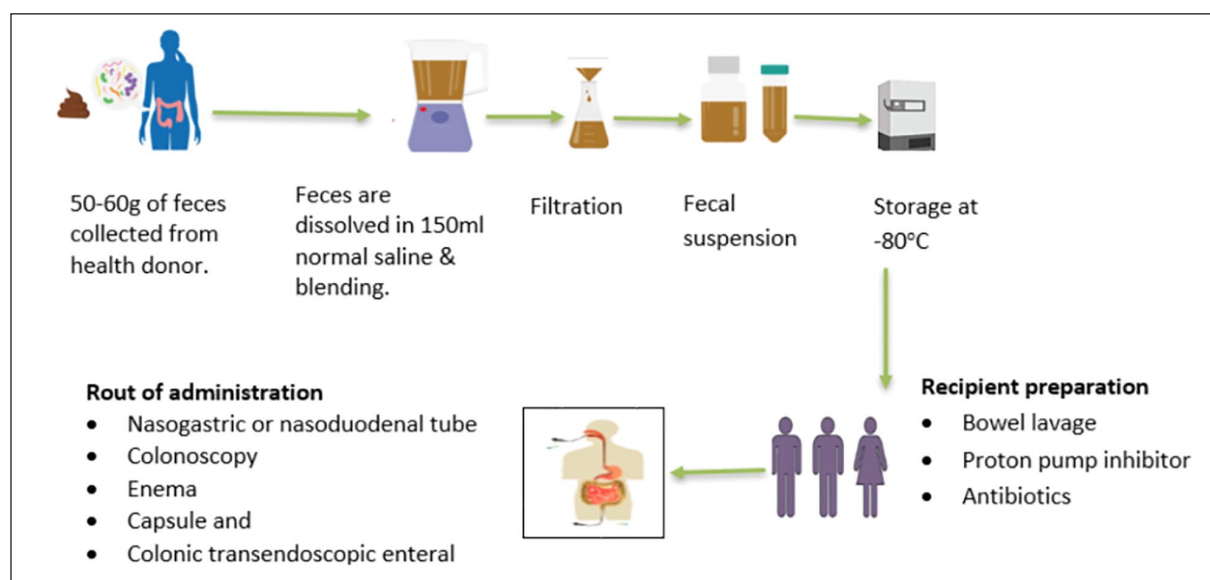
**Figure 1.** The flow of fecal microbiota transplantation.

Following donor selection, a fecal sample from the donor is suspended in milk, tap or bottled water, or no bacteriostatic saline solution, however the latter is thought to be less likely to impact the microbiota of donor stool. Then, the donor feces are homogenized, either by hand agitation and shaking or by means of a machine blender. The mixture is strained through a steel strainer or gauze after being suspended with the diluent to get rid of the bigger particles<sup>41,45</sup> (Figure 2). A fecal suspension can be delivered via nasogastric or nasoduodenal tube, colonoscopy, enema, or capsule and colonic transendoscopic enteral tubing.<sup>46</sup> After FMT

procedure the receipt should be followed for any adverse effect, complications, and for the effectiveness of the procedure (Figure 1).

### Composition of fecal microbiota

There hasn't been much research done on the makeup of human feces. However, studies that have looked at composition have revealed variable findings, possibly due to various factors including genetics, diet, age, environment, and medication use.<sup>24</sup> Adult feces typically contain 25% solid



**Figure 2.** The schematic diagram of the stool collection and fecal microbiota preparation.

substance and 75% water.<sup>47</sup> Organic material, of which 25%–54% are microbial cells, makes up the great majority of solid matter.<sup>48</sup> Fecal microbiota primarily composed of bacteria, but also including viruses, fungi, and archaea.<sup>49</sup> The fecal microbiota plays crucial roles in digestion, immune function, metabolism, and even neurological processes.<sup>50</sup> Bacterial species within the fecal microbiota belong to various phyla, with *Firmicutes* and *Bacteroidetes* typically dominating in healthy individuals. *Firmicutes* are adept at breaking down complex carbohydrates, while *Bacteroidetes* contribute to the degradation of dietary fiber. Other phyla such as *Actinobacteria* and *Proteobacteria* are also present, even though in smaller proportions.<sup>50</sup> Within these phyla, there exists a vast array of genera and species, each with its own unique functions and interactions. For instance, members of the genera *Bifidobacterium* and *Lactobacillus* are renowned for their probiotic properties, promoting gut health and aiding in the synthesis of vitamins.<sup>51</sup> Conversely, certain taxa like *Clostridioides difficile* can become pathogenic under dysbiotic conditions, causing infections and gastrointestinal disorders.<sup>52</sup> Apart from bacteria, the fecal microbiota also harbors viruses (mainly bacteriophages), fungi (such as *Candida* and *Saccharomyces*), and archaea (including *Methanobrevibacter smithii*).<sup>49,50</sup> These microbial constituents contribute to the overall balance and functionality of the gut ecosystem.

### Fecal microbiota transplantation and its clinical application

Fecal bacteriotherapy, also known as fecal microbiota transplantation (FMT), is a revolutionary therapeutic approach that involves the transfer of fecal microbiota from a healthy

donor to a recipient with a dysbiotic gut microbiome. This procedure aims to restore microbial balance and functionality in the gastrointestinal tract, offering a promising treatment for various gastrointestinal and systemic disorders.<sup>46</sup> FMT has shown remarkable efficacy in the management of recurrent *Clostridioides difficile* infection, with success rates exceeding 90%, surpassing those of conventional antibiotic therapy.<sup>53</sup> Now a day, the most frequent indication for fecal microbiota transplantation is recurrent *Clostridioides difficile* infection; however, fecal transplantation is also being tested as a treatment for other gastrointestinal diseases as well as some non-gastrointestinal conditions, such as Parkinson's disease, fibromyalgia, chronic fatigue syndrome, multiple sclerosis, obesity, insulin resistance, metabolic syndrome, autism, and more, though these should be further evaluated in clinical trials.<sup>54</sup>

### *Clostridium difficile* infection

Even though *Clostridioides difficile* is a normal component of the intestinal microbial environment, pre-exposure to broad-spectrum antibiotics can increase the risk of infection because they upset the gut flora's balance, which *Clostridioides difficile* needs to survive.<sup>55</sup> The speed and accuracy of the prescribed therapy will determine how quickly and effectively the infection progresses, but even in the presence of an effective treatment, there is a chance that the infection will return.<sup>56</sup> In the past few decades, FMT has received considerable attention because of a convincing clinical trial of treatment of recurrent *Clostridioides difficile* infection. The first randomized controlled trial of FMT for 43 patients with recurrent *Clostridioides difficile* infection compared FMT administered via nasoduodenal tube after



4–5 days of oral vancomycin with 14 days of continued vancomycin alone and with 14 days of vancomycin plus bowel lavage.<sup>57</sup>

Numerous research has examined the fecal microbiota of donors and patients with *Clostridioides difficile* infection recurrence, and they have demonstrated that the *Clostridioides difficile* infection is linked to less changes in the diversity and composition of the fecal microbiota. When compared to samples from post-FMT patients with recurrent *Clostridioides difficile* infection and healthy donor samples, the number of members of the *Streptococcaceae*, *Enterococcaceae*, or *Enterobacteriaceae* was raised, while the number of butyrate-producing (*Lachnospiraceae* and *Ruminococcaceae*) species was decreased.<sup>58</sup> According to the review article,<sup>59</sup> 85% of recurrent *Clostridium difficile* infection and 55% of new *Clostridioides difficile* infection were successfully treated with FMT, whereas medical therapy had success rates ranging from 30% to 80%. Recently, for individuals with recurring *Clostridioides difficile* infection, RBX2660, a live biotherapeutic agent, presents a very promising therapy alternative. RBX2660 helps patients achieve clinically significant improvements by reestablishing a healthy gut microbiota.<sup>60</sup> In parallel, researchers are investigating the potential of adjunctive therapies, including bacteriophages,<sup>61</sup> Bacteriocin,<sup>62</sup> Probiotics,<sup>62</sup> antimicrobial peptides,<sup>63–65</sup> and immunomodulators,<sup>66</sup> to enhance the efficacy of FMT or next-generation microbial therapies for *Clostridioides difficile* infection. These combination therapies target various aspects of *Clostridioides difficile* infection pathogenesis, such as bacterial virulence, host immune response, and microbiota restoration, to provide more comprehensive and durable treatment outcomes.

### Irritable bowel syndrome

Fecal microbiota transplantation has emerged as a promising avenue for the treatment of irritable bowel syndrome (IBS), a common gastrointestinal disorder characterized by abdominal pain, bloating, and changes in bowel habits.<sup>67</sup> While the exact etiology of IBS remains unclear, growing evidence suggests that dysbiosis, an imbalance in the gut microbiota composition, plays a significant role in its pathogenesis.<sup>68</sup> It is thought that the intestinal microbiota, immune system, and brain-gut axis interact intricately in the etiology of irritable bowel syndrome, which is complicated and poorly understood.<sup>69</sup> FMT involves the transfer of fecal microbiota from a healthy donor to an IBS patient, with the aim of restoring microbial balance. Initial clinical studies have shown encouraging results, with some patients experiencing significant improvements in their gastrointestinal symptoms and overall quality of life following FMT.<sup>67,68</sup>

Furthermore, ongoing research is exploring next-generation therapies that build upon the principles of FMT to enhance efficacy and safety. These include targeted microbial interventions, such as the administration of specific microbial strains or microbial consortia tailored to address

the dysbiosis observed in IBS patients.<sup>70,71</sup> Additionally, advances in microbiome science have led to the development of personalized approaches, where fecal microbiota from a healthy donor is selected based on the recipient's unique microbial profile, maximizing compatibility and therapeutic benefits.<sup>72</sup> Moreover, innovative delivery methods, such as oral capsules or microbial biofilms, are being investigated to optimize the delivery and retention of microbial therapeutics in the gastrointestinal tract.<sup>73,74</sup> These next-generation therapies hold great promise in reshaping the treatment landscape for IBS, offering patients more effective and personalized interventions that target the underlying mechanisms driving their symptoms.<sup>75</sup> As research in this field continues to advance, the integration of FMT and next-generation therapies into clinical practice has the potential to revolutionize the management of IBS and improve outcomes for millions of individuals worldwide.

### Carcinoma of the colon

It is now thought that dysbiosis and the pro-carcinogenic qualities of bacteria (genotoxicity, inflammation, and oxidative stress) may be connected to the development of colorectal cancer. Some bacterial species, including *Bacteroides fragilis*, *Streptococcus bovis*, *Clostridium septicum*, *Helicobacter pylori*, *Enterococcus faecalis*, *Escherichia coli*, and *Fusobacterium spp.*, have been found and are suspected of contributing to the development of colorectal cancer.<sup>76</sup> The complicated interaction between tumor cells, non-neoplastic cells (stromal cells), and a great deal of microbes results in colorectal cancer. In recent decades, more focus has been placed on the role of microbial infection in carcinogenesis in addition to known risk factors (fat-rich diets, obesity, population, and living in a developed country) and uncontrolled cellular proliferation. Microbes are suspected to be responsible for 20% of cancers, particularly colorectal cancer.<sup>77</sup> Fecal transplantation procedures could take the place of colorectal cancer associated dysbiosis and restore eubiosis in chronic disease, assisting in lowering the activation of inflammatory, proliferative, and pro-carcinogenic pathways as well as microbiota-induced genotoxicity. Future transplantation studies will be a crucial next step in this line of study, even if fecal transplantation has not been well investigated in colorectal cancer.<sup>78</sup>

### Crohn's disease and ulcerative colitis

Crohn's disease and ulcerative colitis, collectively known as inflammatory bowel diseases (IBD), chronic inflammatory conditions of the intestines, and the etiology of these conditions involves dysbiosis, which causes the mucosal immune system to become activated, causing chronic inflammation and the emergence of mucosal lesions. While the exact causes of IBD remain elusive, dysbiosis of the gut microbiota is believed to play a significant role in disease

pathogenesis.<sup>10</sup> Fecal microbiota transplantation (FMT) has emerged as a promising therapeutic option for IBD, particularly for patients who fail to respond to conventional treatments.

Initial clinical studies have shown that FMT can induce remission and improve symptoms in a subset of patients with IBD.<sup>79–81</sup> However, the efficacy of FMT varies widely among individuals, highlighting the need for optimized protocols and personalized approaches. Next-generation therapies for IBD are now being explored to enhance the therapeutic potential of FMT and address the limitations associated with donor variability, safety concerns, and unpredictable outcomes. One approach involves the development of microbial-based therapeutics, such as defined microbial consortia or engineered probiotics, tailored to target specific dysbiotic patterns associated with Crohn's disease and ulcerative colitis.<sup>82,83</sup> Furthermore, advancements in delivery methods, such as encapsulation or targeted delivery systems, are being investigated to improve the efficacy and safety of microbial-based interventions for IBD.<sup>84,85</sup> By enhancing the delivery and retention of therapeutic microbes in the gut, these innovative approaches may improve treatment outcomes and reduce the need for repeated administrations.

In addition to microbial-based therapies, other next-generation treatments for IBD include microbiome-modulating agents,<sup>86</sup> immunomodulators,<sup>87</sup> and targeted biologics<sup>88</sup> that aim to modulate the host immune response and restore intestinal homeostasis. These novel therapies offer promising avenues for precision medicine approaches in the management of Crohn's disease and ulcerative colitis, with the potential to improve patient outcomes and quality of life.

### Neuropsychiatric diseases

Today, it's thought that changes to the intestinal microbiome during the first few months of life may be the cause of serious neuropsychiatric conditions that manifest as adulthood approaches, including schizophrenia, autism, behavioral or cognitive disorders, depression, anxiety, Alzheimer's disease, multiple sclerosis, chronic fatigue syndrome, and schizophrenia.<sup>89</sup> There is an increasing focus on the impact of intestinal bacteria on human health, and recent data points to a potential function for the microbiota, gut, and brain axis in neuropsychiatric diseases. There has been a lot of research in this area during the past few years. Although the paths for this relationship are not completely understood, they include metabolic, humoral, immunological, and neurological pathways.<sup>90</sup> It was established that the endotoxin lipopolysaccharide would have an impact on how the central nervous system was modulated. Additionally, it generates inflammatory cytokines, which alter the physiological activity of the brain.<sup>91</sup>

Recently, a child with chronic illness and epilepsy who had been receiving sodium valproate medication up to the transplant was described as the first patient to use fecal

transplantation to achieve remission of digestive symptoms. After 20 weeks, effectiveness in preventing seizures was seen without the use of anti-epileptic medications.<sup>89</sup>

### Obesity and metabolic syndrome

Obesity and metabolic syndrome are multifactorial conditions influenced by genetic, environmental, and lifestyle factors. Emerging research suggests that alterations in the gut microbiota composition and function may contribute to the development of these conditions. Intestinal microbiota is one the primary cause of obesity,<sup>92–94</sup> modifying nutrient absorption and energy management in addition to food consumption,<sup>95,96</sup> which is crucial to the physiology of obesity. Obesity is a significant contributor to the risk of developing diabetes, hypertension, and the metabolic syndrome.<sup>97,98</sup> The etiology of obesity and related disorders is therefore greatly influenced by the gut microbiota.

Fecal microbiota transplantation (FMT) has been proposed as a new way to change the gut microbiota that may result in beneficial metabolic changes, though the evidence for FMT effectiveness in the treatment of obesity.<sup>98–102</sup> Now, bariatric surgery is the only treatment option for morbid obesity that sustains significant weight loss.<sup>103</sup> Next-generation therapies for obesity and metabolic syndrome may involve more targeted approaches, such as precision microbiome modulation.<sup>104</sup> This could include the development of microbial-based interventions, such as engineered probiotics or prebiotics designed to specifically target metabolic pathways or modulate the gut microbiota composition in a more controlled manner.<sup>105</sup> Next-generation therapies that leverage advances in microbiome science may offer more targeted and effective approaches for managing these conditions in the future.

### Glucose intolerance

Chronic hyperglycemia and changes to the metabolisms of carbohydrates, lipids, and proteins are the hallmarks of the metabolic illness known as diabetes mellitus (DM), which is on the rise throughout the world. Polymorphonuclear leukocytes, T lymphocytes, and the immune system's reaction to antigen exposure are all altered in diabetic patients, along with bactericidal activity, bactericidal response, and polymorphonuclear leukocyte function.<sup>106</sup> Numerous potential long-term consequences are associated with both type 1 and type 2 diabetes mellitus, and they are often inversely correlated with the degree and persistence of hyperglycemia.<sup>107</sup>

Studies have shown that individuals with glucose intolerance often exhibit dysbiosis, characterized by alterations in the gut microbiota composition and decreased microbial diversity.<sup>108</sup> It has been found that there is a correlation between the ratio of *Bacteroidetes* to *Firmicutes* and plasma glucose levels in type 2 diabetic and obese patients, suggesting that altering the microbial composition may offer a fresh

method for preventing and treating obesity and type 2 diabetes.<sup>109</sup> Preclinical studies in animal models and some small-scale human trials have demonstrated improvements in glucose metabolism following FMT.<sup>110–112</sup> These improvements are thought to be mediated by changes in the gut microbiota that positively influence metabolic function.

Precision microbiome modulation represents a cutting-edge approach to targeting specific microbial taxa or metabolic pathways associated with glucose intolerance. Engineered probiotics or prebiotics can be designed to selectively alter the gut microbiota, aiming to enhance glucose metabolism.<sup>113</sup> Additionally, the development of microbiome-targeted drugs holds promise for directly targeting microbial metabolites or signaling pathways involved in glucose regulation.<sup>114</sup> By leveraging advances in microbiome research, including metagenomics and metabolomics, personalized interventions tailored to an individual's unique gut microbiota profile could revolutionize the management of glucose intolerance. These innovative strategies offer the potential for more targeted and effective interventions.

### *Illnesses resulting from allergies*

Allergy-related conditions such as food allergies, asthma, and eczema have become more common in recent years. According to recent research, gut microbial alterations have a significant impact on the immunological processes that may result in the emergence of allergy illnesses. These are susceptible to modulation by a variety of environmental factors, including nutrition, antibiotic use, and early-life microbial exposures.<sup>115</sup> Immunological processes that might result in the emergence of allergy disorders are significantly influenced by changes in gut microbiota. The potential of the microflora to affect the immune response has led to new therapeutic modalities that utilize these variations in microbiota for the treatment and prophylaxis of allergy. By transplanting a complex population of bacteria that is more stable and has a larger capacity to colonize, FMT appears to hold promise for restoring immunological homeostasis.<sup>116,117</sup>

### **Adverse effect of fecal microbiome transplantation**

Fecal microbiota transplantation (FMT) is generally considered a safe and effective treatment for certain conditions, but there are potential adverse effects associated with the procedure.<sup>118</sup> One significant risk is the transmission of infections from the donor to the recipient, despite strict donor screening protocols. These infections can include bacterial, viral, and parasitic agents.<sup>118,119</sup> Additionally, recipients may experience gastrointestinal symptoms such as abdominal pain, bloating, diarrhea, or constipation following FMT, although these symptoms are typically mild and transient.<sup>120,121</sup> In rare cases, allergic reactions to components of the fecal material or medications used during the procedure may occur,

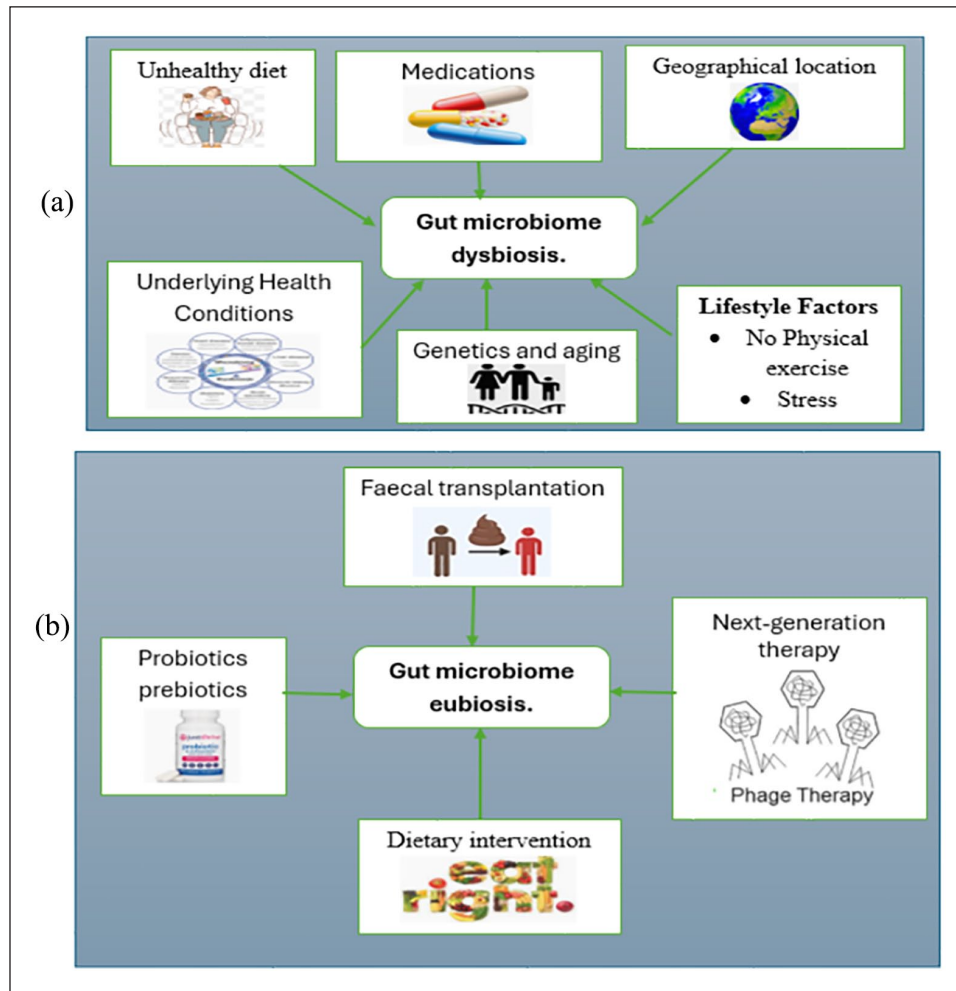
ranging from mild itching to severe anaphylaxis.<sup>119,122</sup> There is also a risk of unintended changes or imbalances in the recipient's gut microbiota, potentially leading to new or exacerbated gastrointestinal symptoms.<sup>119,123</sup> Furthermore, since FMT is still a relatively novel procedure, there may be unknown long-term risks associated with it. These could include the transmission of chronic diseases from the donor to the recipient or triggering immunological reactions in the recipient. Thus, careful consideration of the potential risks and benefits, along with thorough donor screening and close monitoring of recipients, is essential in the use of FMT as a therapeutic intervention.

### **Conclusions and recommendation**

Dysbiosis, the imbalance of gut flora, is influenced by various factors including diet, lifestyle, medications, and underlying health conditions (Figure 3(a)). To counter dysbiosis and restore a healthy gut composition (eubiosis), microbiota modulation techniques are being explored (Figure 3(b)). By targeting specific microbial communities or metabolic pathways, these techniques hold promise for precisely and effectively restoring gut microbiota balance, offering potential advancements in therapeutic. In conclusion, fecal microbiota transplantation (FMT) and next-generation therapies represent exciting avenues for targeting dysbiosis in metabolic disorders and other health conditions. As our understanding of the gut microbiome continues to evolve, so does the potential for innovative interventions aimed at restoring microbial balance and promoting health. FMT, with its demonstrated efficacy in treating conditions like recurrent *Clostridioides difficile* infection, serves as a cornerstone in this field, while emerging therapies such as personalized microbiome modulation and synthetic microbial consortia offer promise for more tailored and precise interventions. Regulatory considerations, ongoing research efforts, and public awareness initiatives will be instrumental in realizing the full therapeutic potential of these approaches. With continued advancements in microbiome science and therapeutic innovation, the future holds great promise for harnessing the power of the gut microbiota to improve health and well-being across diverse patient populations.

### **Future perspective of fecal microbiota transplantation**

The future of fecal microbiota transplantation is promising, with ongoing research to further elucidate its therapeutic potential and expand its clinical applications across a wide range of health conditions. By altering the human gut flora, fecal microbiota transplantation could benefit HIV-infected patients.<sup>124</sup> Phage therapy may be able to eradicate aggressive bacteria from a diseased gut and promote the growth of commensal bacteria. Future research in the field of gut microbiota modulation should focus on



**Figure 3.** (a) Factors influencing dysbiosis, that is, alteration of gut flora and (b) microbiota modulation techniques inducing eubiosis, that is, restoration of altered gut composition.

several key areas to advance our understanding and enhance therapeutic strategies.

**Refinement of Treatment Protocols:** Continued research and clinical trials will likely lead to the refinement of FMT protocols, including optimal donor selection criteria, preparation methods, and delivery techniques. This could improve the safety, efficacy, and reproducibility of FMT procedures.

**Personalized Medicine Approaches:** Advances in microbiome sequencing technologies and computational analysis techniques may enable personalized FMT treatments designed to the unique microbial profiles of individual patients. This could involve matching donors and recipients based on specific microbial signatures or using synthetic microbiota formulations designed to address specific dysbiosis patterns.

**Microbiome Modulation Therapies:** Beyond whole stool-based FMT, future therapies may involve targeted manipulation of the gut microbiome using defined microbial consortia, microbial metabolites, or microbial-derived products. These approaches could offer more precise and controlled

interventions with potentially fewer risks compared to traditional FMT.

**Regulatory Considerations:** As FMT becomes more widely used and novel applications emerge, regulatory frameworks may evolve to ensure the safety, quality, and ethical standards of fecal microbiota transplantation. This includes establishing guidelines for donor screening, standardized protocols for FMT procedures, and oversight of FMT-related research and clinical practice.

**Integration with Other Therapies:** FMT may be integrated with other therapeutic modalities, such as antibiotics, probiotics, prebiotics, dietary interventions, and immunomodulatory agents, to optimize treatment outcomes and address underlying disease mechanisms comprehensively. Combining FMT with complementary therapies could enhance its efficacy and reduce the risk of disease recurrence.

**Public Awareness and Acceptance:** Continued education and outreach efforts aimed at healthcare providers, patients, and the public will be crucial to raise awareness about FMT, dispel misconceptions, and foster acceptance of this



innovative treatment modality. Increased awareness may also facilitate greater participation in donor recruitment efforts and clinical trials.

### Acknowledgements

We extend our gratitude to all the authors who have contributed their work and data to this topic.

### Author contributions

All authors contributed significantly to the paper's conception and design, data collection, and writing the paper; agreed to submit the manuscript to the current journal; granted final approval of the version to be published.

### Availability of data and materials

Not applicable.

### Declaration of conflicting interests

The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

### Funding

The author(s) received no financial support for the research, authorship, and/or publication of this article.

### Ethical approval

Not applicable.

### Informed consent

Not applicable.

### Consent for publication

Not applicable.

### ORCID iDs

Zenawork Sahle  <https://orcid.org/0000-0002-9296-993X>

Demissew Shenkute Gebreyes  <https://orcid.org/0000-0003-2245-2609>

Tsegahun Asfaw Abebe  <https://orcid.org/0000-0002-7410-5484>

### References

1. Thursby E and Juge N. Introduction to the human gut microbiota. *Biochem J* 2017; 474(11): 1823–1836.
2. Bull MJ and Plummer NT. Part 1: the human gut microbiome in health and disease. *Integr Med (Encinitas)* 2014; 13(6): 17–22.
3. Piché ME, Tchernof A and Després JP. Obesity phenotypes, diabetes, and cardiovascular diseases. *Circ Res* 2020; 126(11): 1477–1500.
4. Sitkin SI, Tkachenko EI and Vakhitov TY. Metabolic dysbiosis of the gut microbiota and its biomarkers. *Eksp Klin Gastroenterol* 2016; 12(12): 6–29.
5. Crudele L, Gadaleta RM, Cariello M, et al. Gut microbiota in the pathogenesis and therapeutic approaches of diabetes. *EBioMedicine* 2023; 97: 104821.
6. Strati F, Lattanzi G, Amoroso C, et al. Microbiota-targeted therapies in inflammation resolution. *Semin Immunol* 2022; 59: 101599.
7. Mutalub YB, Abdulwahab M, Mohammed A, et al. Gut microbiota modulation as a novel therapeutic strategy in cardiometabolic diseases. *Foods* 2022; 11(17): 2575.
8. Olofsson LE and Bäckhed F. The metabolic role and therapeutic potential of the microbiome. *Endocr Rev* 2022; 43(5): 907–926.
9. Allegretti JR, Mullish BH, Kelly C, et al. The evolution of the use of faecal microbiota transplantation and emerging therapeutic indications. *Lancet* 2019; 394(10196): 420–431.
10. Burke KE and Lamont JT. Fecal transplantation for recurrent *Clostridium difficile* infection in older adults: a review. *J Am Geriatr Soc* 2013; 61(8): 1394–1398.
11. Zhang X, Luo X, Tian L, et al. The gut microbiome dysbiosis and regulation by fecal microbiota transplantation: umbrella review. *Front Microbiol* 2023; 14: 1286429.
12. Belvoncikova P, Maronek M and Gardlik R. Gut dysbiosis and fecal microbiota transplantation in autoimmune diseases. *Int J Mol Sci* 2022; 23(18): 10729.
13. Ademe M. Benefits of fecal microbiota transplantation: a comprehensive review. *J Infect Develop Countries* 2020; 14(10): 1074–1080.
14. Sandhu A and Chopra T. Fecal microbiota transplantation for recurrent *Clostridioides difficile*, safety, and pitfalls. *Therap Adv Gastroenterol* 2021; 14: 17562848211053105.
15. Panchal P, Budree S, Scheeler A, et al. Scaling safe access to fecal microbiota transplantation: past, present, and future. *Curr Gastroenterol Rep* 2018; 20: 14.
16. Osman M, Budree S, Kelly CR, et al. Effectiveness, and safety of fecal microbiota transplantation for *Clostridioides difficile* infection: results from a 5344-patient cohort study. *Gastroenterology* 2022; 163(1): 319–322.
17. Kellermayer R. Fecal microbiota transplantation: great potential with many challenges. *Transl Gastroenterol Hepatol* 2019; 4: 40.
18. Bowman KA, Broussard EK and Surawicz CM. Fecal microbiota transplantation: current clinical efficacy and future prospects. *Clin Exp Gastroenterol* 2015; 8: 285–291.
19. Hyland N and Stanton C (eds.) *The gut-brain axis: dietary, probiotic, and prebiotic interventions on the microbiota*. Amsterdam, The Netherlands: Elsevier, 2023.
20. Sommer F and Bäckhed F. The gut microbiota—masters of host development and physiology. *Nat Rev Microbiol* 2013; 11(4): 227–238.
21. Arrieta MC, Stiemsma LT, Amenogbe N, et al. The intestinal microbiome in early life: health and disease. *Front Immunol* 2014; 5: 105813.
22. Shin NR, Whon TW and Bae JW. Proteobacteria: microbial signature of dysbiosis in gut microbiota. *Trends Biotechnol* 2015; 33(9): 496–503.
23. Mariat D, Firmesse O, Levenez F, et al. The firmicutes/bacteroidetes ratio of the human microbiota changes with age. *BMC Microbiol* 2009; 9: 1–6.
24. Rinninella E, Raoul P, Cintoni M, et al. What is the healthy gut microbiota composition? A changing ecosystem across age, environment, diet, and diseases. *Microorganisms* 2019; 7(1): 14.

25. Jandhyala SM, Talukdar R, Subramanyam C, et al. Role of the normal gut microbiota. *World J Gastroenterol* 2015; 21(29): 8787.
26. Marotz CA and Zarrinpar A. Focus: microbiome: treating obesity and metabolic syndrome with fecal microbiota transplantation. *Yale J Biol Med* 2016; 89(3): 383.
27. Xue L, Deng Z, Luo W, et al. Effect of fecal microbiota transplantation on non-alcoholic fatty liver disease: a randomized clinical trial. *Front Cell Infect Microbiol* 2022; 12: 759306.
28. Ghorbani Y. Fecal microbiota transplant from healthy lean donors to individuals with obesity: effect on insulin resistance, metabolic parameters, appetite and metabolites. *Doctoral Dissertation, University of Toronto, Canada*, 2023.
29. Schluter J and Foster KR. The evolution of mutualism in gut microbiota via host epithelial selection. *PLoS Biol* 2012; 10(11): e1001424.
30. Elzinga J, van der Oost J, de Vos WM, et al. The use of defined microbial communities to model host-microbe interactions in the human gut. *Microbiol Mol Biol Rev* 2019; 83(2): 10–128.
31. Zheng D, Liwinski T and Elinav E. Interaction between microbiota and immunity in health and disease. *Cell Res* 2020; 30(6): 492–506.
32. Pickard JM, Zeng MY, Caruso R, et al. Gut microbiota: role in pathogen colonization, immune responses, and inflammatory disease. *Immunol Rev* 2017; 279(1): 70–89.
33. Chung H and Kasper DL. Microbiota-stimulated immune mechanisms to maintain gut homeostasis. *Curr Opin Immunol* 2010; 22(4): 455–460.
34. Song Q, Wang Y, Huang L, et al. Review of the relationships among polysaccharides, gut microbiota, and human health. *Food Res Int* 2021; 140: 109858.
35. Cockburn DW and Koropatkin NM. Polysaccharide degradation by the intestinal microbiota and its influence on human health and disease. *J Mol Biol* 2016; 428(16): 3230–3252.
36. Hijova E. Gut bacterial metabolites of indigestible polysaccharides in intestinal fermentation as mediators of public health. *Bratisl Lek Listy* 2019; 120(11): 807–812.
37. Rowland I, Gibson G, Heinken A, et al. Gut microbiota functions: metabolism of nutrients and other food components. *Euro J Nutr* 2018; 57: 1–24.
38. Ramakrishna BS. Role of the gut microbiota in human nutrition and metabolism. *J Gastroenterol Hepatol* 2013; 28: 9–17.
39. Ahlawat S and Asha Sharma KK. Gut–organ axis: a microbial outreach and networking. *Lett Appl Microbiol* 2021; 72(6): 636–668.
40. Borody TJ and Khoruts A. Fecal microbiota transplantation and emerging applications. *Nat Rev Gastroenterol Hepatol* 2012; 9(2): 88–96.
41. Brandt LJ. Fecal transplantation for the treatment of clostridium difficile infection. *Gastroenterol Hepatol (N Y)* 2012; 8(3): 191–194.
42. Bibbò S, Settanni CR, Porcari S, et al. Fecal microbiota transplantation: screening and selection to choose the optimal donor. *J Clin Med* 2020; 9(6): 1757.
43. Nicco C, Paule A, Konturek P, et al. From donor to patient: collection, preparation and cryopreservation of fecal samples for fecal microbiota transplantation. *Diseases* 2020; 8(2): 9.
44. Hamilton MJ, Weingarden AR, Sadowsky MJ, et al. Standardized frozen preparation for transplantation of fecal microbiota for recurrent clostridium difficile infection. *Am J Gastroenterol* 2012; 107(5): 761–767.
45. Kelly CR, Kahn S, Kashyap P, et al. Update on fecal microbiota transplantation 2015: indications, methodologies, mechanisms, and outlook. *Gastroenterology* 2015; 149(1): 223–237.
46. Choi HH and Cho Y-S. Fecal microbiota transplantation: current applications, effectiveness, and future perspectives. *Clin Endosc* 2016; 49(3): 257–265.
47. Rose C, Parker A, Jefferson B, et al. The characterization of feces and urine: a review of the literature to inform advanced treatment technology. *Crit Rev Environ Sci Technol* 2015; 45: 1857.
48. Bojanova DP and Bordenstein SR. Fecal transplants: what is being transferred? *PLoS Biol* 2016; 14(7): e1002503.
49. Matijašić M, Meštrović T, Paljetak HČ, et al. Gut microbiota beyond bacteria-mycobiome, virome, archaeome, and eukaryotic parasites in IBD. *Int J Mol Sci* 2020; 21(8): 2668.
50. Khaledi M, Poureslamfar B, Alsaab HO, et al. The role of gut microbiota in human metabolism and inflammatory diseases: a focus on elderly individuals. *Ann Microbiol* 2024; 74(1): 1.
51. Kleerebezem M and Vaughan EE. Probiotic and gut lactobacilli and bifidobacteria: molecular approaches to study diversity and activity. *Ann Rev Microbiol* 2009; 63: 269–290.
52. Rodríguez C, Romero E, Garrido-Sanchez L, et al. Microbiota insights in Clostridium difficile infection and inflammatory bowel disease. *Gut Microbes* 2020; 12(1): 1725220.
53. Rohlke F and Stollman N. Fecal microbiota transplantation in relapsing Clostridium difficile infection. *Therap Adv Gastroenterol* 2012; 5(6): 403–420.
54. Filip M, Tzaneva V and Dumitrascu DL. Fecal transplantation: digestive and extradigestive clinical applications. *Clujul Med* 2018; 91(3): 259.
55. Patangia DV, Anthony Ryan C, Dempsey E, et al. Impact of antibiotics on the human microbiome and consequences for host health. *Microbiologyopen* 2022; 11(1): e1260.
56. Mattila E, Uusitalo–Seppälä R, Wuorela M, et al. Fecal transplantation, through colonoscopy, is effective therapy for recurrent Clostridium difficile infection. *Gastroenterology* 2012; 142(3): 490–496.
57. Vyas D, Aekka A and Vyas A. Fecal transplant policy and legislation. *World J Gastroenterol* 2015; 21(1): 6.
58. Song Y, Garg S, Girotra M, et al. Microbiota dynamics in patients treated with fecal microbiota transplantation for recurrent Clostridium difficile infection. *PLoS One* 2013; 8: e81330.
59. Drekonja D, Reich J, Gezahegn S, et al. Fecal microbiota transplantation for clostridium difficile infection: a systematic review. *Ann Intern Med* 2015; 162: 630.
60. Chopra T. A profile of the live biotherapeutic product RBX2660 and its role in preventing recurrent Clostridioides difficile infection. *Exp Rev Anti Infect Therap* 2023; 21(3): 243–253.
61. Zuo T, Wong SH, Lam K, et al. Bacteriophage transfer during faecal microbiota transplantation in Clostridium difficile infection is associated with treatment outcome. *Gut* 2018; 67(4): 634–643.
62. Rea MC, Alemayehu D, Ross RP, et al. Gut solutions to a gut problem: bacteriocins, probiotics and bacteriophage for control of Clostridium difficile infection. *J Med Microbiol* 2013; 62(9): 1369–1378.

63. Nuding S, Frasch T, Schaller M, et al. Synergistic effects of antimicrobial peptides and antibiotics against *Clostridium difficile*. *Antimicrob Agents Chemother* 2014; 58(10): 5719–5725.
64. Xu B, Shaoyong W, Wang L, et al. Gut-targeted nanoparticles deliver specifically targeted antimicrobial peptides against *Clostridium perfringens* infections. *Sci Adv* 2023; 9(39): eadf8782.
65. Hing TC, Ho S, Shih DQ, et al. The antimicrobial peptide cathelicidin modulates *Clostridium difficile*-associated colitis and toxin A-mediated enteritis in mice. *Gut* 2013; 62(9): 1295–1305.
66. Ben-Horin S, Margalit M, Bossuyt P, et al. Combination immunomodulator and antibiotic treatment in patients with inflammatory bowel disease and *Clostridium difficile* infection. *Clin Gastroenterol Hepatol* 2009; 7(9): 981–987.
67. Leylabadlo HE, Heravi FS, Soltani E, et al. The role of gut microbiota in the treatment of irritable bowel syndrome. *Rev Res Med Microbiol* 2022; 33(1): e89–e104.
68. Huang HL, Chen HT, Luo QL, et al. Relief of irritable bowel syndrome by fecal microbiota transplantation is associated with changes in diversity and composition of the gut microbiota. *J Dig Dis* 2019; 20(8): 401–408.
69. Iacob T, Țăulescu DF and Dumitrașcu D. Therapy of the postinfectious irritable bowel syndrome: an update. *Clujul Med* 2017; 90(2): 133.
70. Petrof EO and Khoruts A. From stool transplants to next-generation microbiota therapeutics. *Gastroenterology* 2014; 146(6): 1573–1582.
71. Gupta M, Kapoor B and Gulati M. Bacterial consortia-The latest arsenal to inflammatory bowel disease bacteriotherapy. *Med Microecol* 2024; 31: 100107.
72. Kingsley MJ and Abreu MT. A personalized approach to managing inflammatory bowel disease. *Gastroenterol Hepatol* 2016; 12(5): 308.
73. Liu J, Li X, Zhang X, et al. Gut lumen-targeted oral delivery system for bioactive agents to regulate gut microbiome. *J Future Foods* 2022; 2(4): 307–325.
74. Woo CW, Tso P and Yiu JH. Commensal gut microbiota-based strategies for oral delivery of therapeutic proteins. *Trends Pharmacol Sci* 2022; 43(12): 1004–1013.
75. Lacy BE, Chey WD and Lembo AJ. New and emerging treatment options for irritable bowel syndrome. *Gastroenterol Hepatol* 2015; 11(4 Suppl 2): 1.
76. Gagnière J, Raisch J, Veziat J, et al. Gut microbiota imbalance and colorectal cancer. *World J Gastroenterol* 2016; 22(2): 501.
77. Zur Hausen H. The search for infectious causes of human cancers: where and why. *Virology* 2009; 392(1): 1–10.
78. Schwabe RF and Jobin C. The microbiome and cancer. *Nat Rev Cancer* 2013; 13(11): 800–812.
79. Costello SP, Hughes PA, Waters O, et al. Effect of fecal microbiota transplantation on 8-week remission in patients with ulcerative colitis: a randomized clinical trial. *JAMA* 2019; 321(2): 156–164.
80. Suskind DL, Brittnacher MJ, Wahbeh G, et al. Fecal microbial transplant effect on clinical outcomes and fecal microbiome in active Crohn's disease. *Inflam Bowel Dis* 2015; 21(3): 556–563.
81. Greenberg A, Aroniadis O, Shelton C, et al. Long-term follow-up study of fecal microbiota transplantation (FMT) for inflammatory bowel disease (IBD): 1791. *Am J Gastroenterol* 2013; 108: S540.
82. Barra M, Danino T and Garrido D. Engineered probiotics for detection and treatment of inflammatory intestinal diseases. *Front Bioeng Biotechnol* 2020; 8: 265.
83. Oka A and Sartor RB. Microbial-based and microbial-targeted therapies for inflammatory bowel diseases. *Digest Dis Sci* 2020; 65(3): 757–788.
84. Hua S, Marks E, Schneider JJ, et al. Advances in oral nano-delivery systems for colon targeted drug delivery in inflammatory bowel disease: selective targeting to diseased versus healthy tissue. *Nanomedicine* 2015; 11(5): 1117–1132.
85. Zhang S, Langer R and Traverso G. Nanoparticulate drug delivery systems targeting inflammation for treatment of inflammatory bowel disease. *Nano Today* 2017; 16: 82–96.
86. Alshehri D, Saadah O, Mosli M, et al. Dysbiosis of gut microbiota in inflammatory bowel disease: current therapies and potential for microbiota-modulating therapeutic approaches. *Bosn J Basic Med Sci* 2021; 21(3): 270.
87. Aberra FN and Lichtenstein GR. Monitoring of immunomodulators in inflammatory bowel disease. *Aliment Pharmacol Therap* 2005; 21(4): 307–319.
88. Chan HC and Ng SC. Emerging biologics in inflammatory bowel disease. *J Gastroenterol* 2017; 52: 141–150.
89. He Z, Cui B-T, Zhang T, et al. Fecal microbiota transplantation cured epilepsy in a case with Crohn's disease: the first report. *World J Gastroenterol* 2017; 23(19): 3565.
90. Wrona D. Neural-immune interactions: an integrative view of the bidirectional relationship between the brain and immune systems. *J Neuroimmunol* 2006; 172(1–2): 38–58.
91. Kalyan M, Tousif AH, Sonali S, et al. Role of endogenous lipopolysaccharides in neurological disorders. *Cells* 2022; 11(24): 4038.
92. Zhao L. The gut microbiota and obesity: from correlation to causality. *Nat Rev Microbiol* 2013; 11(9): 639–647.
93. Gérard P. Gut microbiota and obesity. *Cell Mol Life Sci* 2016; 73(1): 147–162.
94. Liu BN, Liu XT, Liang ZH, et al. Gut microbiota in obesity. *World J Gastroenterol* 2021; 27(25): 3837.
95. Garcia G, Sunil TS and Hinojosa P. The fast food and obesity link: consumption patterns and severity of obesity. *Obes Surg* 2012; 22: 810–818.
96. Machado PP, Steele EM, Levy RB, et al. Ultra-processed food consumption and obesity in the Australian adult population. *Nutr Diabetes* 2020; 10(1): 39.
97. Matsuda M and Shimomura I. Increased oxidative stress in obesity: implications for metabolic syndrome, diabetes, hypertension, dyslipidemia, atherosclerosis, and cancer. *Obes Res Clin Pract* 2013; 7(5): e330–e341.
98. Nguyen NT, Magno CP, Lane KT, et al. Association of hypertension, diabetes, dyslipidemia, and metabolic syndrome with obesity: findings from the National Health and Nutrition Examination Survey, 1999 to 2004. *J Am College Surg* 2008; 207(6): 928–934.
99. Yu EW, Gao L, Stastka P, et al. Fecal microbiota transplantation for the improvement of metabolism in obesity: The FMT-TRIM double-blind placebo-controlled pilot trial. *PLoS Med* 2020; 17(3): e1003051.
100. Mocanu V, Zhang Z, Deehan EC, et al. Fecal microbial transplantation and fiber supplementation in patients with severe

- obesity and metabolic syndrome: a randomized double-blind, placebo-controlled phase 2 trial. *Nat Med* 2021; 27(7): 1272–1279.
101. Ng SC, Xu Z, Mak JW, et al. Microbiota engraftment after faecal microbiota transplantation in obese subjects with type 2 diabetes: a 24-week, double-blind, randomised controlled trial. *Gut* 2022; 71(4): 716–723.
  102. Allegretti JR, Kassam Z, Mullish BH, et al. Effects of fecal microbiota transplantation with oral capsules in obese patients. *Clin Gastroenterol Hepatol* 2020; 18(4): 855–863.
  103. Lahtinen P, Juuti A, Luostarinen M, et al. Effectiveness of fecal microbiota transplantation for weight loss in patients with obesity undergoing bariatric surgery: a randomized clinical trial. *JAMA Network Open* 2022; 5(12): e2247226.
  104. Vallianou NG, Kounatidis D, Tsilingiris D, et al. The role of next-generation probiotics in obesity and obesity-associated disorders: current knowledge and future perspectives. *Int J Mol Sci* 2023; 24(7): 6755.
  105. Green M, Arora K and Prakash S. Microbial medicine: prebiotic and probiotic functional foods to target obesity and metabolic syndrome. *Int J Mol Sci* 2020; 21(8): 2890.
  106. Daryabor G, Atashzar MR, Kabelitz D, et al. The effects of type 2 diabetes mellitus on organ metabolism and the immune system. *Front Immunol* 2020; 11: 546198.
  107. Mauri-Obradors E, Estrugo-Devesa A, Jané-Salas E, et al. Oral manifestations of diabetes mellitus. A systematic review. *Med Oral Patol Oral Cir Bucal* 2017; 22(5): e586.
  108. Zhao M, Liao D and Zhao J. Diabetes-induced mechanophysiological changes in the small intestine and colon. *World J Diabetes* 2017; 8(6): 249.
  109. Larsen N, Vogensen FK, Van Den Berg FW, et al. Gut microbiota in human adults with type 2 diabetes differs from non-diabetic adults. *PLoS One* 2010; 5(2): e9085.
  110. Chen L, Guo L, Feng S, et al. Fecal microbiota transplantation ameliorates type 2 diabetes via metabolic remodeling of the gut microbiota in db/db mice. *BMJ Open Diabetes Res Care* 2023; 11(3): e003282.
  111. Wu L, Li MQ, Xie YT, et al. Washed microbiota transplantation improves patients with high blood glucose in South China. *Front Endocrinol* 2022; 13: 985636.
  112. Kootte RS, Levin E, Salojärvi J, et al. Improvement of insulin sensitivity after lean donor feces in metabolic syndrome is driven by baseline intestinal microbiota composition. *Cell Metab* 2017; 26(4): 611–619.
  113. Ji J, Jin W, Liu SJ, et al. Probiotics, prebiotics, and postbiotics in health and disease. *MedComm* 2023; 4(6): e420.
  114. Brown JM and Hazen SL. Targeting of microbe-derived metabolites to improve human health: the next frontier for drug discovery. *J Biol Chem* 2017; 292(21): 8560–8568.
  115. Chung KF. Airway microbial dysbiosis in asthmatic patients: a target for prevention and treatment? *J Allerg Clin Immunol* 2017; 139(4): 1071–1081.
  116. Maksimova O, Gervazieva V and Zverev V. Intestine microbiota and allergic diseases. *J Microbiol Epidemiol Immunobiol* 2014; 91(3): 49–60.
  117. Reynolds LA and Finlay BB. A case for antibiotic perturbation of the microbiota leading to allergy development. *Exp Rev Clin Immunol* 2013; 9(11): 1019–1030.
  118. Park SY and Seo GS. Fecal microbiota transplantation: is it safe? *Clin Endosc* 2021; 54(2): 157–160.
  119. Wang JW, Kuo CH, Kuo FC, et al. Fecal microbiota transplantation: review and update. *J Formosan Med Assoc* 2019; 118: S23–S31.
  120. Bénard MV, de Bruijn CM, Fenneman AC, et al. Challenges and costs of donor screening for fecal microbiota transplantations. *PLoS One* 2022; 17(10): e0276323.
  121. Woodworth MH, Carpentieri C, Sitchenko KL, et al. Challenges in fecal donor selection and screening for fecal microbiota transplantation: a review. *Gut Microbes* 2017; 8(3): 225–237.
  122. Liu SX, Li YH, Dai WK, et al. Fecal microbiota transplantation induces remission of infantile allergic colitis through gut microbiota re-establishment. *World J Gastroenterol* 2017; 23(48): 8570.
  123. Rapoport EA, Baig M and Puli SR. Adverse events in fecal microbiota transplantation: a systematic review and meta-analysis. *Ann Gastroenterol* 2022; 35(2): 150–163.
  124. Serrano-Villar S, Talavera-Rodríguez A, Gosalbes MJ, et al. Fecal microbiota transplantation in HIV: a pilot placebo-controlled study. *Nat Commun* 2021; 12(1): 1139.