The ethicolegal framework relevant to human faecal microbiota transplants in South Africa: Part 1. A legal vacuum

M Labuschaigne,1 BA Hons, MA, DLitt, LLB, LLDE M Slabbert,1 BA Hons, HED, BProc, LLB, LLDE S Budree,2 MB ChB, DCH, FC Paed (SA), Cert Gastroenterology (SA) Paed; E Hoosien,2 MB ChB, MMed Path (Med Microbiol), DTM&H; A Brink,9 MB ChB, MMed (Clin Microbiol); M Blockman,9 BPharm, MB ChB, PG Dip Int Res Ethics, MMed (Clin Pharmacol)

1 Department of Jurisprudence, School of Law, University of South Africa
2 Department of Paediatrics, Faculty of Health Sciences, University of Cape Town, South Africa; and OpenBiome, Cambridge, Mass., USA
3 Clinical Microbiology Department, Ampath Laboratories, Centurion, South Africa
4 Division of Medical Microbiology, Department of Pathology, Faculty of Health Sciences, University of Cape Town, South Africa; and National Health Laboratory Service, Cape Town, South Africa
5 Division of Clinical Pharmacology, Department of Medicine, Faculty of Health Sciences, University of Cape Town, South Africa

Corresponding author: M Labuschaigne (slabbmn@unisa.ac.za)

The legal regulation of faecal microbiota transplantation (FMT) in South Africa (SA) is currently unclear. The purpose of this article, the first of three in a series, is to explore the nature, role and clinical application of FMT in SA in order to determine, from a legal perspective, the appropriate regulatory pathways governing FMT as a procedure that may combine approaches for the treatment of drugs, human tissue for transplantation, or clinical treatment as part of the practice of medicine. FMT has been shown to be a novel, safe and effective treatment for recurrent Clostridioides difficile infection (CDI). Stool banks are instrumental in enabling access to FMT for patients and clinicians and help to catalyse research in the microbiome. However, the regulatory landscape in SA remains unclear. Microbial therapies such as FMT are necessary, especially in a time of rising microbiome-associated inflammatory diseases and increasing resistance to traditional antibiotics. FMT is now considered as part of the standard of care for recurrent CDI overseas, but is currently only being used for research purposes in a minority of clinical cases of CDI in SA. This article, which lays the foundation for consideration of this question in three parts, suggests that the relevant regulatory system would depend on the categorisation of human stool as tissue, the exact composition of the FMT, how it is administered to patients, and the relevant levels of manipulation of the stool for FMT-derived products.

The human body hosts a series of dynamic and diverse microbial communities consisting of bacteria, archaea, viruses and eukaryotes, collectively known as the human microbiome. The microbiome plays a critical role in health and disease and has been shown to influence various medical conditions, ranging from diabetes to multiple sclerosis, autism, cancer and inflammatory bowel disease.24

Globally there has been a rapid increase in research and scientific publications predominantly on the gastrointestinal (GIT) microbiome and its role in health, immunology and nutrition. Therapeutic applications are of great interest, and a successful example is faecal microbiota transplantation (FMT). This process involves the introduction of gut bacteria from healthy donors into patients, with the aim of repairing the abnormal microbial community in the recipient. The efficacy of FMT for treatment of recurrent Clostridioides difficile infection (CDI) is a crucial proof of the concept that therapeutic modification of the GIT microbiome can be an effective treatment modality for human disease.25 Other indications for FMT are also currently being explored.26

The legal regulation of microbiome-based therapy is difficult to categorise, however, as it poses an uncomfortable fit in terms of the South African (SA) legal framework governing the use of human tissue, as well as the legal regulation of medicines and medical devices.26 More specifically, human stool banking or creation of a repository is critical to an FMT service. The purpose of this article is to explore the use of FMT as a microbiome-based therapy in SA by exploring the nature of FMT, followed by the role of FMT in clinical practice and then indicating the legal vacuum in which FMT currently exists.

The need for legal clarity on the regulation of FMT is underscored by a recent safety alert issued by the US Food and Drug Administration (FDA) regarding the transmission of a multidrug-resistant organism that resulted in two immunocompromised patients developing bacteraemia following an FMT.14 This incident led to the death of one patient, and subsequent investigation found that the donor had not been screened for extended-spectrum beta-lactamase organisms.23

Understanding the role of the microbiome

Microbial cells outnumber human cells by approximately three to one.30 The combined microbial genome encodes 500 times more genes than the human genome. The human microbiota refers to all the micro-organisms residing in the human body. The microbiome includes all the microbiota, and their genetic products – proteins, metabolites and RNA.18,31

The human gut has the largest collection of microbes, producing signals that influence host metabolic, immune and physiological functions. The five most abundant bacterial phyla found in healthy individuals are Bacteroidetes, Firmicutes, Actinobacteria, Proteobacteria and Verrucomicrobia. Within each phylum there are hundreds of bacterial species, each found in different abundances.
in healthy individuals. Gut bacterial lineages are highly variable, making a healthy core microbiome difficult to describe at organism phyllum level. Shared microbial genes in the gut of different individuals indicate a common functional core microbiome at the gene level, implying that different microbes have similar metabolites or functions and the absence of a single bacterial species may not be impactful.

A healthy microbiome consists of a diverse assembly of microbial species in specific anatomical niches carrying out their specific biomolecular function. High microbial alpha diversity is associated with host health and microbial stability to environmental pressures. Alpha diversity can be described as the number of different microbial taxa and their abundance in an ecosystem. Dysbiosis is a term that refers to disruption of the normal gut microbiome (e.g. a decrease in microbial alpha diversity) and is associated with various diseases such as obesity, inflammatory bowel disease, autoimmune disease, allergy and diabetes. Broad-spectrum antibiotics can decrease microbial diversity with variable recovery dynamics. When healthy gut microbes are diminished, they are unable to compete against pathogens wanting to colonise the gut. In addition, certain beneficial microbes that produce beneficial metabolites (e.g. short-chain fatty acids (SCFAs)) perform important metabolic functions (e.g. metabolism of bile salts), produce anti-inflammatory metabolites or enhance gut epithelial function, and may be lost due to antibiotics or disease.

There are several factors that influence the microbiome during early life. Significant ones include mode of delivery (normal vaginal delivery v. caesarean section), gestational age at delivery, infant feeding patterns (breastfed v. formula fed) and weaning foods, antibiotic use, and maternal factors such as health and diet. A recently published study that included the largest dataset of children born by either caesarean section or vaginal delivery demonstrated persistent significant differences in the microbial community composition associated with mode of delivery. The gut microbiome undergoes substantial changes or maturation during the first 3 years of life and contributes to the development and maturation of the child's immune system. By the age of 3, the child's microbiome resembles that of an adult.

The ongoing composition of the microbiome is influenced by host genetics to a limited extent and more significantly by environmental factors. Age, sex, geographical location, chronic disease, diet, medication (specifically antibiotics) and other environmental exposures (chemical or microbial) all play a role. Diet greatly shapes gut microbiota and may explain the geographical variation observed between populations on different continents.

Gut-derived signalling molecules influence peripheral organs and metabolism. Signalling molecules can be part of the bacterial structure, such as lipopolysaccharide, which interacts with host receptors to produce an immunological response. Other important signalling molecules are byproducts of bacterial metabolism and are influenced by diet, such as SCFAs. Gut microbiota can be considered as a separate organ that influences the immune system, brain, lung, liver, skin, adipose tissue and metabolism. The microbiome-gut-brain axis refers to the myriad effects that the gut microbiota has on the brain and its functions.

FMT is one modality that is used to manipulate the microbiome. Other techniques include probiotics, diet, phage therapy and antibiotics. Extensive data suggest that FMT can restore the healthy intestinal microbiome in a patient by enabling engraftment of donor microbes into recipients.

C. difficile is an anaerobic, spore-forming, Gram-positive intestinal pathogen that is present in our environment and can asymptptomatically colonise a healthy host. CDI occurs when bacterial toxins cause diarrhoea, and in certain patients this progresses to a life-threatening colitis. Dysbiosis of the gut microbiome caused by antibiotic use is the main risk factor for CDI. The first-line treatment for CDI is administration of antibiotics directed against CDI, but despite various dosing regimens for oral vancomycin, 20% of patients go on to develop recurrent CDI. Recent US and European guidelines now support the use of FMT to treat recurrent CDI. Meta-analysis of randomised controlled trials and case series indicates a cure rate of 80 - 90% with a single treatment, but some patients will require more than one FMT. Current evidence suggests that FMT is a safe and well-tolerated procedure in patients with recurrent CDI.

To date, a single USA-based stool bank, OpenBiome, has treated >50 000 patients with recurrent CDI and currently supports >30 enrolling FMT clinical trials globally. This stool banking model (based on the blood banking model) is critical to an FMT service from both doctor/patient access and research perspectives. Prior to the expansion of stool banks, patients were required to identify their own donors who would need to be consented and screened by the treating doctor, a process that was prohibitively expensive and introduced ethical concerns around coercion and disclosure from patient-selected donors. In addition, the collection and processing of the material was unpleasant and beyond the scope of most doctors. These factors were reported as major barriers to access to FMT for both patients and doctors. The stool bank model removed many of these barriers and enabled almost universal access to FMT in the USA. In the stool bank model, prospective donors undergo a detailed clinical assessment aimed at ruling out those who are an infectious risk or have a personal or immediate family history of microbiome-mediated diseases such as autoimmune disease, allergy, neuropsychiatric disease, malignancy, etc. Thereafter, prospective donors undergo extensive blood and stool tests for potential pathogens.

Those who pass this process are onboarded as stool donors and undergo repeat clinical and laboratory screening at 2-monthly intervals, in addition to brief clinical assessment at each donation, to ensure that there has been no acute change in the donor's health status. Stool is frozen in a cryopreservative buffer and stored at ~80°C until needed. In addition, a safety aliquot is retained from each sample that is donated. This safety aliquot can be used for pathogen testing during investigation of an adverse event. Owing to the theoretical risk of transfer of inflammatory disease from donors to recipients, stool banks aim to retain data and samples from donors for extended periods. The American Gastroenterological Association has established an FMT registry in the USA that aims to prospectively follow up patients, adult and paediatric, up to 10 years after FMT, with the aim of establishing long-term safety data.

**Regulatory challenges to date**

Regulatory challenges in the USA and Europe are one of the main reasons why FMT is not widely adopted among doctors. In 2013, the FDA announced that FMT would be regulated as a biological and as an investigational drug. An initial regulation was that an investigational new drug (IND) application was required for access for all disease indications. Following the landmark paper by Van Nood et al., demonstrating significantly superior efficacy of FMT over placebo or standard of care in recurrent CDI, and subsequent pressure from providers and patients, the FDA chose to exercise enforcement discretion regarding the requirement for an IND requirement to treat recurrent CDI, thereby allowing clinicians and patients to access FMT. The use of FMT to treat any other disease indication would, however, require an IND application.
Another significant barrier for doctors and patients was access to material for treatment. The study by Lee et al.,[31] which demonstrated similar efficacy of frozen v. fresh FMT, catalysed the development of the universal donor stool bank. In this model, donors could be thoroughly screened and material banked for convenient future use by doctors.[33]

In 2016, a draft guidance statement from the FDA raised a safety concern related to the use of banked stool samples.[30] It proposed that stool banks would be required to submit an IND application to obtain and distribute stool. The draft guidance would not affect clinicians who were collecting and screening patient-selected donor stools. A safety concern of the FDA was that centralised stool banking may lead to a relatively small number of donors providing material to multiple patients. Potential transmission of infectious agents and other unidentifiable risks related to the microbiome of the few donors would reach many recipients.[31]

Criticism of the proposed regulation change was raised among both clinicians and consumers, and the draft guidance was never implemented. Owing to the ease of access to safe material using a stool bank, most medical facilities in the USA choose to use stool bank material for FMT. Stools for banking undergo extensive investigation for transmissible pathogens, as opposed to direct donation, where quality control is not possible.[31] Stool banks offer a centralised point of reference for safety monitoring and regulatory oversight and reduce costs through economies of scale.

A recent framework has been proposed in the USA that allows the degree of manipulation of the donor faecal material to determine the regulatory category.[34,35] Faecal material from a donor that is not manipulated or minimally manipulated (e.g. addition of a cryopreservative buffer or saline homogenisation) would be considered practice of medicine and not subject to federal regulation. When stools are manipulated and consist of defined bacterial populations, faecal material would fall under a traditional drug category. Stool banks would fall under a regulatory category like that of a tissue or blood bank.

Based on the successful US model of a stool bank of having treated >50 000 patients with only two adverse events, the ideal scenario would be to replicate this model in the SA and African context, while at the same time adapting it to the local context and disease burden. In order to ensure that FMT is performed safely, and is accessible to doctors, patients and researchers, SA must carefully consider how best to regulate stool banks and FMT.

**Nature of FMT for the purpose of legal regulation**

It seems as if stool donations would fall within the scope of the definition of ‘biological material’ as defined in the 2012 Regulations on the Use of Human Biological Material,[36] promulgated in terms of the National Health Act 61 of 2003 (NHA), which refers to biological material ‘material from a human being’. The use of stool samples in FMT is not covered directly by these regulations or any other provisions in the NHA itself. The nature of FMT is equally unclear, as it is difficult to determine whether the procedure should be regarded in law as a medicine or a medical treatment.

The nature of FMT is unique and different from conventional medicines or therapies. It may include characteristics and components of a ‘health service’ in terms of the NHA, which include medical treatment, or a ‘medicine’ as described in the Medicines and Related Substances Control Act 101 of 1965. Each of these has different legal and ethical considerations, depending on the nature of the procedure.

If administered orally by way of swallowing capsules, FMT will arguably be regarded as a medicine. The administration of FMT via a naso-duodenal tube or colonoscopy resembles medical treatment. The confusion regarding the exact nature of FMT is also evident from the FDA’s position on the issue, which has shifted a few times in recent years. As mentioned above, the FDA currently classifies FMT as a drug (e.g. a live biotherapeutic product), subject to enforcement discretion of not requiring an IND for doctors performing the procedure for patients with recurrent CDI unresponsive to standard therapies.[31]

Regulatory systems may decide to regulate FMT as both a product and a procedure that combines approaches for the treatment of drugs, blood and human tissue for transplantation, or view it as part of the practice of medicine (e.g. clinical treatment). The regulatory pathway would depend on the exact composition of the FMT and the relevant formulations or levels of manipulation of the stool, which may range from basic frozen stool to capsules containing frozen or lyophilised stool, or defined consortia of bacteria.[31] Stool-based products containing specific bacterial strains aimed at treating specific diseases involve a higher level of manipulation and therefore a more target-specific approach.

In the most elementary FMT procedure, faeces are liquefied, and saline or glycerol (e.g. a cryopreservative) added, which points to minimal manipulation. The FDA’s definition of ‘minimal manipulation’ of cells or non-structural tissue in the case of the transplantation of human cells and tissues refers to ‘processing that does not alter the relevant biologic characteristics of cells or tissue’.[37] This prompts the question regarding what the original relevant characteristics of stool are, and how one knows that these have changed. Osssorio and Zhou[38] rightly remark that if the scientific community has not adequately characterised the health-related functions of microbes in complex communities, considering that ‘people’s microbial communities are highly dynamic, then clinicians and regulators lack metrics for determining when a microbial community’s relevant properties have been altered by manipulation’.

According to Hoffmann et al.,[39] the degree of manipulation of stool may vary as follows from the least to the most manipulated: (i) fresh stool transferred from an individual donor; (ii) frozen filtered stool from a stool bank; (iii) concentrated stool microbiota in capsule form; (iv) biologically sourced, purified microbial groups of specific bacterial strains; and (v) cultured bacterial mixture delivered in oral pill form.

In the USA, stool that is extensively manipulated, isolated or combined with other components, or used for novel functions, would therefore be regulated under section 351 of the Public Health Services Act (2012) and the Federal Food, Drug, and Cosmetic Act as biological drugs or devices requiring premarket approval under Title 21 of the Code of Federal Regulations (1271.20), as stated above. Determining whether manipulated stool is a product or a procedure may also require characterisation of the stool (e.g. to document the structure and properties of the material). The regulatory process for live biotherapeutic products traditionally requires characterisation of all the microbes in the faecal material, which would require whole-community genome sequencing of all the microbes contained in the stool.[24] This would not only be scientifically challenging but impossible, because human stool differs biologically and chemically considerably from one person to the next, rendering the regulatory requirement of consistency in product composition unfeasible.[24]

There is no doubt that FMT using single-donor stool (e.g. from a friend or relative) creates fewer risks, and it is for this reason that the recent draft FDA guidance document makes provisions for an ‘enforcement discretion’ when ‘the stool donor and stool are qualified by screening and testing performed under the direction of the licensed healthcare provider for the purpose of providing the FMT product for treatment of the patient’. The same cannot be said for
public health risks associated with universal donors or pooled stool. In the case of pooled stool from multiple donors, it is possible that interactions between different microbial communities and increased risk of infection may arise. The complexity of pooled stool is that it is difficult to trace the source of infection back to the actual donor. Osorio and Zhou, however, in response to Hoffmann et al., maintain that 'if stool composition is dynamic and difficult to characterize, then a minimally-manipulated stool product for allogeneic transplant is possibly the most dangerous type of microbial product for a recipient. Compared to highly-manipulated but better characterized stool-derived products, stool from an asymptomatic donor is more likely to contain novel or uncharacterized viruses, bacteria, or protozoa that could be pathogenic, particularly for an already sick recipient.' However, the role of minimally manipulated stool (transplantation of the entire microbial community from a healthy donor) should not be dismissed, as there may be currently uncharacterized features of stool (e.g., viral, fungal components or metabolites) that could convey beneficial effects in some diseases. In addition, the use of whole-stool FMT will provide scientific data to inform the future selection of specific microbial bacteria necessary to develop defined consortia.

It is clear from this discussion that the uncertainties regarding the nature of stool and stool-derived products will continue to hamper their effective classification.

Conclusions
FMT has been shown to be a novel, safe and effective treatment for recurrent CDI. Stool banks are instrumental in enabling access to FMT for patients and clinicians and help to catalyse research in the microbiome. However, the regulatory landscape in SA remains unclear. Microbial therapies such as FMT are necessary, especially in a time of rising microbiome-associated inflammatory diseases and increasing resistance to traditional antibiotics. FMT is now considered as part of standard of care for recurrent CDI overseas, but is currently only being used for research purposes or in a minority of clinical cases of CDI in SA. The advantages of the procedure are not disputed, but what is needed is clarity on the regulatory framework in which FMT is performed. This article recommends that the relevant regulatory pathway for FMT would depend on the exact composition of the FMT and the relevant formulations or levels of manipulation of the stool. In part 2 of this series of articles, the legal categorisation of human stool as human tissue will be considered, as well as the application thereof in clinical practice. This will involve an analysis of the relevant legal framework to determine how FMT will be regulated in SA in future. The fact that FMT is still in a research phase in SA also has peculiar legal consequences that need to be explored. If the stool donations are processed and transformed into products, such as capsules, the legal framework may change completely. These issues will be interrogated further in subsequent parts of this series of articles.

Declaration. None.
Acknowledgements. None.
Author contributions. Equal co-authors.
Funding. None.
Conflicts of interest. SB is employed by Finch Therapeutics, Somerville, Mass., USA, a company developing microbial therapies related to the treatments described in these articles.

1. Wang H, Wei C-X, Min L, Ling-Yun Zhu L-Y. Good or bad: Gut bacteria in human health and diseases. Bacterial Biotechnol Exp 2018;32(5):1075-1080. https://doi.org/10.1093/mae/bex030
2. Fedorova E, Hovesten E, Brint K. A tale for stool banks in South Africa. S Afr Med J 2019;109(5):546-547. https://doi.org/10.7196/SAMJ.2019.v109i5.14169
3. Farrow P, Bhuyan S, Scholer A, et al. Scaling safe access to fecal microbiota transplantation: Past, present, and future. Curr Gastroenterol Rep 2018;20(1):4. https://doi.org/10.1007/s11894-018-0189-8
4. US Food and Drug Administration. FDA in brief: FDA warns about potential risk of serious infections caused by multi-resist drug-resistant organisms. (accessed 30 September 2019).
5. Keisler EJ, Allegretti J, Hawkey P, et al. A necessary discussion of transmission of multidrug-resistant organisms through faecal microbiota transplantation. Lancet Infect Dis 2019;19(11):1561-1562. https://doi.org/10.1016/S1473-3099(19)30545-6
6. DeFilipp Z, Bloom PP, Soto MT, et al. Drug-resistant E. coli bacterium transmittable by local fecal microbiota transplantation. N Engl J Med 2019;381(21):2021-2026. https://doi.org/10.1056/NEJMc1913947
7. Blaser MJ. Fecal microbial transplantation for dysbiosis: predictable risk? N Engl J Med 2019;381(22):2064-2066. https://doi.org/10.1056/NEJMc1913947
8. Romo J. Ten times more microbial cells than body cells in humans! Microbe 2019;14(2):47-47. https://doi.org/10.1287/micro.19.0072
9. Human Microbiome Jumpsart Reference Strains Consortium. Wilson KE, Winters-Giuliano CA, Baldwin SL, et al. A catalog of reference genomes from the human microbiome. Science 2013;339(6125):944-949. https://doi.org/10.1126/science.1228136
10. Cortez PL, Eldred V, Aurangzeb M, et al. Enterotypes in the landscape of gut microbiome composition. Nat Microbiol 2018;3(8):1-16. https://doi.org/10.1038/s41564-018-0072-8
11. Turmingh PJ, Gordon JI. The core gut microbiome: energy balance and obesity. J Physiol 2009;587(Pt 1):143-158. https://doi.org/10.1111/j.1469-7588.2008.02127.x
12. Human Microbiome Project Consortium. Structure, function and diversity of the healthy human microbiome. Nature 2012;486(7409):207-214. https://doi.org/10.1038/nature11234
13. Knight R, Callaway C, Marcoe M, et al. The microbiome and human biology. Am Rev Genetomics Human Genet 2017;18:165-86. https://doi.org/10.1111/gene.12458
14. Hoffmann DE, Palumbo FB, Ravel J, Rowthorn V, von Rosenvinge E. A proposed definition of microbiota dysbiosis in health and disease. Curr Gastroenterol Rep 2018;20(4):14. https://doi.org/10.1007/s11894-018-0619-8
15. Kassam Z, Dubois N, Ramakrishna B, et al. Donor screening for fecal microbiota transplantation. N Engl J Med 2019;381(13):1360-1. https://doi.org/10.1056/NEJMc1913670
16. Gibson MIK, Wang B, Almada S, et al. Developmental dynamics of the preterm infant gut microbiota and enterotypes. Nat Microbiol 2016;1:1624. https://doi.org/10.1038/nmicrobiol.2016.134
17. Willard RC, Clemente JC, Peter L, Loss JF. The preclinical gut microbiome. Are we colonized with bacteria in utero? Pediatr Obstet Gynecol 2017;32(Suppl 1):1-17. https://doi.org/10.1097/POG.0000000000000217
18. Zhang M, Dörschlend MG, Benjamin-Jones IE, Osthau T, Heyo C, Madler NT. Association of prenatal antibiotics with measures of infant adiposity and the gut microbiome. Ann Clin Microbiol Antimicrob 2019;18(1):18. https://doi.org/10.1186/s12937-019-0316-1
19. Shao Y, Forner GC, Tiabhi E, et al. Stunted microbiota and opportunistic pathogen colonization in caesarean-section born infants. Nature 2019;575(7781):480-484. https://doi.org/10.1038/s41586-019-1170-4
20. Enrich et al. The Integrative Human Microbiome Project, what’s next for the microbiome community? Nat Microbiol 2016;1(4):779-799. https://doi.org/10.1038/nmicrobiol.2016.117-w
21. Lloyd-Price J, Alo A-G, Hutteranner C. The healthy human microbiome. Genome Med 2016;8(6):51. https://doi.org/10.1186/s13073-016-0165-7
22. Wahl L, Valli G, Peeters H, Van Pauwenberge M, et al. The role of minimally manipulated bacteria, or protozoa that could be pathogenic, particularly for an already sick recipient. However, the role of minimally manipulated stool (transplantation of the entire microbial community from a healthy donor) should not be dismissed, as there may be currently uncharacterized features of stool (e.g., viral, fungal components or metabolites) that could convey beneficial effects in some diseases. In addition, the use of whole-stool FMT will provide scientific data to inform the future selection of specific microbial bacteria necessary to develop defined consortia.

23. Ossorio and Zhou, however, in response to Hoffmann et al., maintain that ‘if stool composition is dynamic and difficult to characterize, then a minimally-manipulated stool product for allogeneic transplant is possibly the most dangerous type of microbial product for a recipient. Compared to highly-manipulated but better characterized stool-derived products, stool from an asymptomatic donor is more likely to contain novel or uncharacterized viruses, bacteria, or protozoa that could be pathogenic, particularly for an already sick recipient.’ However, the role of minimally manipulated stool (transplantation of the entire microbial community from a healthy donor) should not be dismissed, as there may be currently uncharacterized features of stool (e.g., viral, fungal components or metabolites) that could convey beneficial effects in some diseases.

24. In the case of pooled stool from multiple donors, it is possible that treatments described in these articles.

25. The complex breakdown of the stool is that it is difficult to trace the source of infection back to the actual donor. Osorio and Zhou, however, in response to Hoffmann et al., maintain that ‘if stool composition is dynamic and difficult to characterize, then a minimally-manipulated stool product for allogeneic transplant is possibly the most dangerous type of microbial product for a recipient. Compared to highly-manipulated but better characterized stool-derived products, stool from an asymptomatic donor is more likely to contain novel or uncharacterized viruses, bacteria, or protozoa that could be pathogenic, particularly for an already sick recipient.’ However, the role of minimally manipulated stool (transplantation of the entire microbial community from a healthy donor) should not be dismissed, as there may be currently uncharacterized features of stool (e.g., viral, fungal components or metabolites) that could convey beneficial effects in some diseases.

26. In the case of pooled stool from multiple donors, it is possible that treatments described in these articles.

27. In the case of pooled stool from multiple donors, it is possible that treatments described in these articles.