${\bf S}$ Table 1. Recent trials conducted to investigate the effects of probiotics on clinical outcomes of cancer patients

| ClinicalTrials.g ov Identifier | Location | Disease | Numbe r of Particip ants | Intervention/Treatment | Duration | Outcome | Reference |
|-----------------------------------|----------|---|-----------------------------------|---|-----------|--|-----------|
| NCT03358 511 | USA | Breast cancer | 20 | Primal Defense Ultra® Probiotic (Saccharomyces boulardii, Lactobacillus plantarum, Bacillus subtilis, Bifidobacterium lactis, Bifidobacterium bifidum, Lactobacillus rhamnosus, Bifidobacterium breve, lactobacillus casei, Lactobacillus salivarius, Lactobacillus acidophilus, Lactobacillus brevis, Bifidobacterium longum, and Lactobacillus paracasei) | 2-4 weeks | Recruiting subjects. This study examines if pre-operative probiotics will help the body's immune system react to breast cancer. | N/A |
| NCT03290 651 | Canada | Women at risk of breast cancer | 40 | Lactobacillus rhamnosus GR-1 and Lactobacillus reuteri RC-14 | 90 days | Recruiting subjects. This study examines if probiotics reduce harmful bacteria, as well as inflammation, in the breast, and lower the chances of developing breast cancer. Changes in breast microbiota will be determined by nextgeneration sequencing. | N/A |
| NCT02819 960 | Slovakia | Colon cancer | 100 | PROBIO-FIX INUM® (Lactobacillus rhamnosus GG, Bifidobacterium animalis subsp. lactis BB-12) | 6 weeks | Recruiting subjects. This trial evaluates potential of probiotics to prevent grade 3-4 diarrhea in patients treated with irinotecan. | N/A |

| NCT03705 442 | Croatia | Metastatic colorectal cancer | 76 | OMNi-BiOTiC® 10AAD (Lactobacillus acidophilus, Lactobacillus acidophilus, Lactobacillus paracasei, Lactobacillus rhamnosus, Enterococcus faecium, Lactobacillus salivarius, Lactobacillus | 84 days | Recruiting subjects. This study examines if probiotics can be used as adjuvant therapy to reduce diarrhea in cancer patients receiving chemotherapy. | N/A |
|-----------------|----------|------------------------------------|----|---|------------|---|------------|
| | | | | plantarum, Bifidobacterium bifidum, Bifidobacterium lactis, Bifidobacterium longum) | | | |
| NCT03782 428 | Malaysia | Colon cancer | 52 | Lactobacillus acidophilus, Lactobacillus lactis, Lactobacillus casei subsp, Bifidobacterium longum, Bifidobacterium bifidum and Bifidobacterium infantis | 6 months | Postoperative probiotics reduced the level of proinflammatory cytokines, including TNF-α, IL-6, IL-10, IL-12, IL-17A, IL17C and IL-22, in patients. | Ref. (113) |
| NCT03072 641 | Sweden | Colon cancer | 36 | ProBion Clinica (Bifidobacterium lactis BI-04, Lactobacillus acidophilus NCFM and inulin) | 31±28 days | Probiotics increased level of butyrate-producing bacteria,including Faecal ibacterium and Clostridia les spp in the tumor, nontumor mucosa and faecal microbiota of patients while reducing the level of Fusobacterium and Peptostreptococcus in the faecal microbiota of patients. | Ref. (106) |
| NCT01468 779 | Brazil | Colon cancer | 91 | Lactobacillus acidophilus, Lactobacillus rhamnosus, Lactobacillus casei, Bifidobacterium and fructooligosaccharide | 19 days | Perioperative administration of symbiotics significantly reduced postoperative infection rates in patients. | Ref. (100) |

| NCT01609 660 | Brazil | Colon cancer | 33 | Saccharomyces boulardii | 7 days | Preoperative probiotics downregulated proinflammatory cytokines IL-1β and IL-23A, as well as anti-inflammatory cytokine IL-10, in the intestinal colonic mucosa, with no statistical impact on postoperative infection rates. | Ref. (105) |
|-----------------|----------|-----------------|----|---|----------|---|---------------|
| NCT01479 907 | Greece | Colon cancer | 75 | Synbiotic Forte TM (Pediococcus pentosaceus, Leuc onostoc mesenteroides, La ctobacillus paracasei ssp. paracasei, Lactob acillus plantarum, b-glucan, inulin, pectin and resistant starch) | 15 days | Postoperative synbiotics ameliorated postcolectomy gastrointestinal function and diarrhea in patients. | Ref. (101) |
| NCT01410 955 | Slovakia | Colon cancer | 46 | Colon Dophilus™ | 12 weeks | Probiotics reduced the incidence of irinotecaninduced diarrhea in | Ref. (102) |

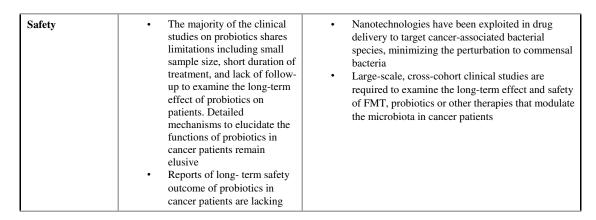
| | | | | | | colorectal cancer patients. | |
|-----------------|--------|--|----|--|--|--|------------|
| NCT01790 035 | USA | Gastroin testinal, abdomin al, or pelvic cancer | 23 | Lactobacillus rhamnosus GG (LGG) | LGG administered at least 3 days prior, during, and 2 weeks following radiation treatment | In vitro and in vivo models were used to identify the mechanism of LGG radioprotection. Preliminary result suggested LGG releases radioprotective lipoteichoic acid (LTA), which primes the protection of epithelial stem cells by triggering an immune signaling cascade involving macrophages and PGE2 secreting mesenchymal stem cells. | Ref. (112) |
| NCT02654 652 | Brazil | Head and neck cancer | 40 | LactoFos (Lactoba cillus paracasei LPC-31, L. rhamnosus HN00l, L. acidophilus NCF M, and Bifidobacterium lactis HN019 plus fructooligosaccharides) | 5-7 days | Postoperative symbiotics did not affect intestinal function and postoperative outcomes of head and neck surgical patients. | Ref. (104) |

| NCT03829 111 | USA | Kidney cancer | 30 | Clostridium butyricum CBM 588 | Treatment every 21 days for 4 cycles; treatment every 28 days beginning cycle 5 | Recruiting subjects. This phase I trial examines if CBM 588 and immunotherapy (nivolumab and ipilimumab) exert a synergistic effect on patients with advanced kidney cancer. | N/A |
|-----------------|-------|-------------------|----|--|---|---|------------|
| NCT02771 470 | China | Lung cancer | 41 | Clostridium butyricum | 3 weeks | Clostridium butyricum promoted beneficial genera, including Blautia, Clostridium, Faecalibacterium and La ctobacillus, while decreasing the abundance of pathogenic bacteria Escherichia-Shigella. C butyricum reduced not only chemotherapyinduced diarrhea but also systemic inflammatory response in patients. However, there was no significant alteration in lymphocyte subsets, immunoglobulin and albumin expression levels. | Ref. (103) |
| NCT03574 051 | China | Thyroid cancer | 30 | Bifidobacterium infantis, Lactobacillus acidophilus and Enterococcus faecalis | 30 days | Recruiting subjects. This study examine if probiotics prevent adverse effects of hypothyroidism and improve side effects caused by 131I treatment. | N/A |

Legend for S Table 1: Several trials report an improved clinical outcome in cancer patients receiving probiotics whereas other trials deny the clinical benefits of probiotics. Probiotics have been shown to modulate the production of pro-inflammatory cytokines, to lower the incidence of chemotherapy-induced diarrhea, as well as postoperative infection, and to protect the intestine from radiation by releasing lipoteichoic acid. On the contrary, strands of evidence indicate that probiotic treatment does not improve intestinal function of cancer patients. Patients receiving the probiotics and those receiving the placebo had similar rate of postoperative infection and displayed comparable level of inflammatory markers and diamine oxidase, an indicator of gut permeability. The majority of the trials assesses the effect of probiotics on cancer treatment-induced toxicities rather than their direct impact on mediating the efficacy of anti-cancer drugs. Ongoing and future trials that examine the interactions between probiotics and cancer therapy will predict patient's response to treatment and provide new insights into personalized medicine tailored according to patient's microbial profile.

\boldsymbol{S} Table 2. Limitations of current microbiome research and efforts to address these concerns

| Manipulating the microbiome to prevent/ treat cancer | Current state | Ongoing work and future trend |
|---|---|---|
| Conception | The genomic content and functions of bacterial strains can be changed during the lifespan of individuals by forces of evolution, including host-associated factors, environmental factors or microbe-microbe interactions Microbial profiles are determined by 16S rRNA gene amplicon and metagenomic sequencing, with limited resolution of bacterial species at strainlevel | Efforts are directed to identify bacterial strains that play a significant role in cancer, and to elucidate the associated mechanisms Functions of bacterial strains should be examined in the context of the entire microbiota as bacteria interact with one another, as well as the host and other microorganisms, in this complex ecosystem Findings are to be validated extensively with preclinical models and large-scale, cross-cohort clinical trials |
| Research approach | Microbial profiles are determined by 16S rRNA gene amplicon and metagenomic analysis of stool and mucosa samples of cancer patients Inferred functions of microbes based on association analysis Correlation-based | The role of bacteria in cancer should be examined by integrative analysis involving host genome/metabolome/ immune profile, oral/gut/faecal microbiome, fungal and viral communities The role of exogenous factors, such as diet and medications should be included to elucidate the complex interactions between the host, the microbes, and anti-cancer drugs Causation/Mechanism-based |
| Clinical translation | Candidate bacteria related to cancer diagnosis or prognosis can potentially be used as novel biomarkers for cancer Modulating the microbiota via FMT improves the outcome of patients with Clostridium difficile infection, inflammatory bowel disease or other diseases Probiotic treatment improves cancer treatment-induced toxicities, including diarrhea and infection | Precision therapy to be developed based on host and microbiome characteristics Diet modification or probiotic/prebiotic supplementation may augment the anti-tumor effect of chemotherapy/ immunotherapy and improve patient outcomes |



Legend for S Table 2: Various aspects of microbiome research and their implications in cancer are examined. With advances in sequencing technology and development of powerful computational tools, the research paradigm has shifted from association-based approach to mechanism-based approach. Unravelling causal links between bacteria and cancer has become an area of active research. In an attempt to generate a holistic view of the microbiota-host interaction in the context of cancer, researchers examine the interactions between the gut microbiome, the host, anti-cancer drugs and other exogenous factors through integrative analysis of multiple types of sequencing data. Regardless of the previous findings, it remains challenging to assess the clinical benefits of FMT/probiotics in cancer patients. The majority of the clinical trials on probiotics shares limitations, such as small sample size, short duration of treatment and lack of follow-up to examine the long-term effect of probiotics on patients. As such, well-designed studies that explore the direct interactions between the microbiota, tumor cells and anticancer drugs are critical for the evaluation of FMT/probiotic treatment in cancer patient. FMT, fecal microbiota transplant.