The Gut Microbiome and Inflammation

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What is the Microbiome

- Collective genomes of **microbial flora** harbored by humans
- Microflora is the community of microorganisms, including algae, fungi, and bacteria that live in or on another living organism or in a particular habitat

The Microbiome

SOURCE: N. POPGEORGIEV ET AL/JOURNAL OF INFECTIOUS DISEASES 2013



- The ratio of the amount of microbes to cells in our bodies is 10:1
- Most identified DNA sequences floating in our blood plasma belong to viruses
- We are more them than we are us

http://linxc10.wixsite.com/microbes/the-microbiome

Depiction of the human body and bacteria that predominate





Grice and Serge 2011

Ecosystem: Skin histology with microorganisms



Grice and Serge 2011



Young VB 2017

Prebiotic

Nutrients that favor the growth and predominance of beneficial microbes and their inherent functions. Most of these have been carbohydrates that cannot be broken down by the human digestive machinery but are metabolized by specific members of the microbiota

Probiotic

Commonly defined as "live micro-organisms which, when administered in adequate amounts, confer a health benefit on the host." In the past, these were often organisms that were first recognized in fermented food products. Currently, there is interest in identifying potential probiotics that are members of the microbiota of healthy people

All diseases begin in the gut

The bacterial cells harbored within the human gastrointestinal tract (GIT) outnumber the host's cells by a factor of 10 and the genes encoded by the bacteria resident within the GIT outnumber their host's genes by more than 100 times.

Approximately 100 trillion micro-organisms in the gut



Hippocrates The Father of Western Medicine 460 BC-370 BC



Composition and luminal numbers of dominant microbial species in the human gastrointestinal tract.



Sartor and Mazmanian 2012

The gut microbiota and diseases

- Lower bacterial diversity has been reproducibly observed in people with inflammatory bowel disease, psoriatic
 arthritis, type 1 diabetes, atopic eczema, coeliac disease, **obesity**, type 2 diabetes, and arterial stiffness, than in
 healthy controls.
- Germ-free mice that receive faecal microbes from obese humans gain more weight than mice that receive microbes from healthy weight humans

IBD: Ulcerative Colitis and Crohn's disease

Colitis is reduced or absent in germ free animals and those treated with antibiotics

More inflammation



Arthritis is reduced or absent in germ free animals



Clemente et al., 2018

s of foods, nutrients, and dietary patterns that influen	ce human health linked to their effe	ect on the gut microb	iota
Effect on gut microbiome	Effect on health outcomes mediated by gut microbiome	Human observational studies	Human interventional studies
Low FODMAP diet increased Actinobacteria; high FODMAP diet decreased abundance of bacteria involved in gas consumption ⁵⁸	Reduced symptoms of irritable bowel syndrome ⁵⁶	Yes	Yes
Increased <i>Bifidobacteria</i> , ^{97 98} which are known for their positive health benefits to their host through their metabolic activities. ⁹⁹ Decrease in <i>Bacteroides</i> and <i>Clostridia</i> , some strains of which are associated with intestinal infections ⁹⁸	Potential protection against pathogens. ¹⁰⁰ Increased production of SCFA and reduced production of TMAO ⁹⁹	Yes	Yes
Increased microbiota diversity and SCFA production ^{22 101 102}	Reduced type 2 diabetes ²² and cardiovascular disease ¹⁰³	Yes	Yes
Overgrowth of Proteobacteria and <i>Escherichia coli</i> . ¹⁰⁴ <i>Bacteroides</i> , <i>Clostridia</i> , and total aerobic bacteria were significantly lower, and faecal pH was significantly higher ⁴⁷	Induced glucose intolerance ¹⁰⁵	No	No
Increased intestinal barrier protectors (<i>Bifidobacteria</i> and <i>Lactobacillus</i>), butyrate producing bacteria (<i>Faecalibacterium prausnitzii</i> and <i>Roseburia</i>) and <i>Bacteroides vulgatus</i> and <i>Akkermansia muciniphila</i> . ¹⁰⁷ Decreased lipopolysaccharide producers (<i>E coli</i> and <i>Enterobacter cloacae</i>) ¹⁰⁶	ut micro-organisms alter polyphenol Yes oavailability resulting in reduction metabolic syndrome markers and ardiovascular risk markers ¹⁰⁸		Yes
Very modest differences in composition and diversity in humans and strong differences in metabolomic profile compared with omnivore diet in humans ⁵⁰	Some studies show benefit of vegetarian over omnivore diet, ¹⁰⁹ others fail to find a difference ¹¹⁰	Yes	Yes
	 of foods, nutrients, and dietary patterns that influen Effect on gut microbiome Low FODMAP diet increased Actinobacteria; high FODMAP diet decreased abundance of bacteria involved in gas consumption⁵⁸ Increased <i>Bifidobacteria</i>, ^{97 98} which are known for their positive health benefits to their host through their metabolic activities.⁹⁹ Decrease in <i>Bacteroides</i> and <i>Clostridia</i>, some strains of which are associated with intestinal infections⁹⁸ Increased microbiota diversity and SCFA production²²¹⁰¹¹⁰² Overgrowth of Proteobacteria and <i>Escherichia coli</i>.¹⁰⁴<i>Bacteroides</i>, <i>Clostridia</i>, and total aerobic bacteria were significantly lower, and faecal pH was significantly higher⁴⁷ Increased intestinal barrier protectors (<i>Bifidobacteria</i> and <i>Lactobacillus</i>), butyrate producing bacteria (<i>Faecalibacterium prausnitzii</i> and <i>Roseburia</i>) and <i>Bacteroides vulgatus</i> and <i>Akkermansia muciniphila</i>.¹⁰⁷ Decreased lipopolysaccharide producers (<i>E coli</i> and <i>Enterobacter cloacae</i>)¹⁰⁶ Very modest differences in composition and diversity in humans and strong differences in metabolomic profile compared with omnivore diet in humans⁵⁰	of foods, nutrients, and dietary patterns that influence human health linked to their effect Effect on gut microbiomeEffect on gut microbiomeEffect on health outcomes mediated by gut microbiomeLow FODMAP diet increased Actinobacteria; high FODMAP diet decreased abundance of bacteria involved in gas consumption58Reduced symptoms of irritable bowel syndrome56Increased Bifidobacteria, ^{97 98} which are known for their positive health benefits to their host through their metabolic activities. ⁹⁹ Decrease in Bacteroides and Clostridia, some strains of which are associated with intestinal infections98Potential protection against pathogens. ¹⁰⁰ Increased production of SCFA and reduced production of TMAO99Increased microbiota diversity and SCFA production22101102Reduced type 2 diabetes22 and cardiovascular disease ¹⁰³ Overgrowth of Proteobacteria and Escherichia coli. ¹⁰⁴ Bacteroides, Clostridia, and total aerobic bacteria were significantly lower, and faecal pH was significantly higher ⁴⁷ Induced glucose intolerance ¹⁰⁵ Increased intestinal barrier protectors (Bifidobacteria and Lactobacillus), butyrate producing bacteria (Faecalibacterium prausnitzii and Roseburia) and Bacteroides vulgatus and Akkermansia muciniphila. ¹⁰⁷ Decreased lipopolysaccharide producers (E coli and Enterobacter cloacae) ¹⁰⁶ Gut micro-organisms alter polyphenol bioavailability resulting in reduction of metabolic syndrome markers and cardiovascular risk markers ¹⁰⁸ Very modest differences in composition and diversity in humans and strong differences in metabolomic profile compared with omnivore diet in humans ⁵⁰ Some studies show benefit of vegetarian over omnivore diet, ¹⁰⁹ others fail to find a difference ¹¹⁰	of foods, nutrients, and dietary patterns that influence human health linked to their effect on the gut microbioneHuman observational studiesEffect on gut microbiomeEffect on health outcomes mediated by gut microbiomeHuman observational studiesLow FODMAP diet increased Actinobacteria; high FODMAP diet decreased abundance of bacteria involved in gas consumptionsReduced symptoms of irritable bowel syndromesYesIncreased Bifidobacteria, ^{97,98} which are known for their positive health benefits to their host through their metabolic activities, ⁵⁹ Decrease in Bacteroides and Clostridia, some strains of which are associated with intestinal infections ⁹⁸ Potential protection against pathogens. ¹⁰⁰ Increased production of TMAO ⁹⁹ YesIncreased microbiota diversity and SCFA production feacal pH was significantly higher ⁴⁷ Reduced type 2 diabetes ²² and cardiovascular disease ¹⁰³ YesIncreased intestinal barrier protectors (Bifidobacteria and Lactobacillus), butyrate producing bacteria (Faccalibacterium prousnizii and Roseburia) and Bacteroides vulgatus and Akkermansia muciniphila. ¹⁰⁰ Decreased lipopolysaccharide producers (<i>E coli</i> and <i>Enterobacter cloacae</i>)Gut micro-organisms alter polyphenol bioavailability resulting in reduction of metabolic syndrome markers and cardiovascular risk markers ¹⁰⁸ YesVery modest differences in composition and diversity in humans and strong differences in metabolomic profile compared with omnivore diet, in humans ⁵⁰ Some studies show benefit of vegetarian over omnivore diet, ¹⁰⁹ others fail to find a difference ¹¹⁰

FODMAP=fermentable oligosaccharides, disaccharides, monosaccharides and polyols; SCFA=small chain fatty acids; TMAO= trimethylamine N-oxide

Table 2 Summar	y of systematic reviews a	nalysing the role	e of probiotic	cs on clinical outcomes
Outcome	Reference	No of studies/ participants	Evidence of benefit?	Results/conclusions
<i>Clostridium difficile</i> associated diarrhoea in adults and children	Goldenberg et al (2017) ¹¹¹	39/9955	Yes	Moderate quality evidence that probiotics are safe and effective for preventing <i>C difficile</i> associated diarrhoea. (RR 0.30, 95% CI 0.21 to 0.42)
Necrotising enterocolitis	Al Faleh et al (2014) ¹¹² Rees et al (2017) ¹¹³	17/5338	Yes	Enteral supplementation of probiotics prevents severe necrotising enterocolitis (RR 0.43, 95%Cl 0.33 to 0.56) and all cause mortality in preterm infants (RR 0.65, 95% Cl 0.25 to 0.81)
Antibiotic associated diarrhoea in children	Goldenberg et al (2015) ¹¹⁴	26/3898	Yes	Moderate evidence of a fall in the incidence of antibiotic associated diarrhoea in the probiotic v control group (RR 0.46, 95% Cl 0.35 to 0.61; l^2 =55%, 3898 participants)
Probiotics for preventing acute upper respiratory tract infections	Hao et al (2015) ¹¹⁵	12/3720	Yes	Probiotics were better than placebo in reducing the number of participants experiencing episodes of acute upper respiratory tract infections, the mean duration of an episode, antibiotic use, and related school absence (12 trials, 37 20 participants including children, adults, and older people)
Urinary tract infections	Schwenger et al (2015) ¹¹⁶	9/735	No	No significant benefit for probiotics compared with placebo or no treatment
Prevention of asthma and wheeze in infants	Azad et al (2013) ¹¹⁷	6/1364	No	No evidence to support a protective association between perinatal use of probiotics and doctor diagnosed asthma or childhood wheeze
Prevention of eczema in infants and children	Mansfield et al (2014)	16/2797	Yes	Probiotic supplementation in the first several years of life did have a significant impact on development of eczema (RR 0.7 4, 95% CI 0.67 to 0.82)

Prevention of invasive fungal infections in preterm neonates	Agrawal et al (2015) ¹¹⁹	19/4912	Unclear	Probiotic supplementation reduced the risk of invasive fungal infections (RR 0.50, 95% Cl 0.34 to 0.73, I ² =39%) but there was high heterogeneity between studies. Analysis after excluding the study with a high baseline incidence (75%) showed that probiotic supplementation had no significant benefits (RR 0.89, 95% Cl 0.44 to 1.78)
Prevention of nosocomial infections	Manzanares et al (2015) ¹²⁰	30/2972	Yes	Probiotics were associated with a significant reduction in infections (RR 0.80, 95%Cl 0.68 to 0.95, P=0.009; I ² =36%, P=0.09). A significant reduction in the incidence of ventilator associated pneumonia was found (RR 0.74, 95% Cl 0.61 to 0. 90, P=0.002; I ² =19%)
Treatment of rotavirus diarrhoea in infants and children	Ahmadi et al (2015) ¹²¹	14/1149	Yes	Probiotic supplementation resulted in a mean difference of –0.41 (Cl 95% –0.56 to –0.25; P<0.001) in the duration of diarrhoea. Probiotics exert positive effect on reducing the duration of acute rotavirus diarrhoea compared with control
Prevention and treatment of Crohn's disease and ulcerative colitis	Saez Lara et al (2015) ¹²²	14/821 ulcerative colitis 8/374 Crohn's disease	Yes	The use of probiotics and/or synbiotics has positive effects in the treatment and maintenance of ulcerative colitis, whereas in Crohn's disease clear effectiveness has only been shown for synbiotics (no meta- analysis was performed)
Pulmonary exacerbations in children with cystic fibrosis	Ananathan et al (2016) ¹²³	9/275	Yes	Significant reduction in the rate of pulmonary exacerbation (two parallel group randomised controlled trials and one crossover trial: RR 0.25, 95% CI 0.15 to 0.41; P< 0.00001)
Type 2 diabetes (fasting glucose, glycated haemoglobin test)	Akbari et al (2016) ¹²⁴	13/805	Yes	Probiotics significantly reduced fasting blood glucose compared with placebo (8 studies; standardised mean difference -1.583 ; 95% Cl -4.18 to 4.18 ; P=0.000). Significant reduction in HbA _{1c} was also seen (6 studies; SMD -1.779 ; 95% Cl -2.657 to -0.901 ; P=0.000)
Type 2 diabetes (insulin resistance, insulin levels)	Zhang et al (2016) ¹²⁵	7/425	Yes	Probiotic therapy significantly decreased homeostasis model assessment of insulin resistance (HOMA-IR) and insulin concentration (WMD: –1.08, 95% Cl – 1.88 to –0.28; and weighted mean difference –1.35mlU/L, 95% Cl – 2.38 to –0.31, respectively

7./				
Necrotising enterocolitis in pre-term neonates with focus on Lactobacillus reuteri	Athalye-Jape et al (2016) ¹²⁶	6/1778	Yes	Probiotic reduced duration of hospitalisation (mean difference = -10.77 days, 95% Cl -13.67 to -7.86 ; in 3 randomised controlled trials), and late onset sepsis (RR 0.66; 95% Cl, 0.52 to 0.83; 4 RCTs) were reduced in the
Reduction of serum concentration of C reactive protein	Mazidi et al (2017) ¹²⁷	19/935	Yes	Significant reduction in serum C reactive protein after probiotic administration with a WMD –1.35 mg/L, (95% Cl – 2.15 to –0.55, I ² 65.1%)
Cardiovascular risk factors in patients with type 2 diabetes	Hendijani et al (2017) ¹²⁸	11/641	Yes	Probiotic consumption significantly decreased systolic blood pressure (-3.28 mm Hg; 95% Cl -5.38 to -1.18), diastolic (WMD -2.13 mm Hg; 95% Cl -4.5 to 0.24), low density lipoprotein cholesterol (WMD 8.32 mg/dL; 95% Cl -15.24 to -1.4), total cholesterol (WMD -12.19 mg/ dL; 95% Cl -17.62 to -6.75) and triglycerides(WMD -24.48 mg/dL; 95% Cl -33.77 to -11.18) compared with placebo
Reduction of total cholesterol and low density lipoprotein cholesterol	Wu et al (2017) ¹²⁹	15/976	Yes	Lactobacillus consumption significantly reduced total cholesterol by 0.26 mmol/L (95% CI -0.40 to -0.12) and LDL-C by 0.23 mmol/L (95% CI, -0.36 to -0.10)
Depressive symptoms	Wallace and Milev (2017) ^{79,130}	6/1080	Yes	No quantitative analysis was performed. Most studies found positive results, and the authors conclude that compelling evidence shows that probiotics alleviate depressive symptoms
Vulvovaginal candidiasis in non- pregnant women	Xie et al (2018) ¹³¹	10/1656	Yes	Probiotics increased the rate of short term clinical cure (RR 1.14, 95% Cl 1.05 to 1.24, low quality evidence) and mycological cure (RR 1.06, 95% Cl 1.02 to 1.10, low quality evidence) and decreased relapse rate at one month (RR 0.34, 95% Cl 0.17 to 0.68, low quality evidence)
Chronic periodontitis	Ikram et al (2018) ¹³²	7/220	Yes	The overall mean difference for gaining clinical attachment level gain between probiotics and placebo was significant (weighted mean difference 1.41, 95% Cl 0.15 to 2.67, P=0.028)
RR=risk ratio, SBP sys difference' CI=confide	tolic blood pressure, DBP= diastolio nce interval	blood pressure,	TC= total chole	sterol, TG=serum triglycerides, SMD=standardised mean difference, WMD=weighted mean

Consensus and uncertainties

What we know

- Probiotic supplementation has several beneficial effects on human health
- The microbes in our gut influence and human energy metabolism²²⁻⁴⁵
- Diet and medication have a strong influence on gut microbiota composition
- Microbiota composition influences response to chemotherapy and immunotherapy⁹⁶
- Microbiome composition defines glucose response to foods and can be used to personalise diet⁹⁴
- Dietary fibre intake influences gut microbiota composition and is related to better health^{86 87 104}

What we don't know

- Are natural probiotics in food better than probiotic supplements? Should we take them preventively?
- Can microbes influence food choices and appetite?
- Do low dose antibiotics in food affect human health?
- What is the effect of pesticides in food on the gut microbiome? Is organic food better for the gut microbiota?
- Should all new drugs and food chemicals be tested on the gut microbiota?

Potential strategies for therapeutic microbiome manipulation

Approach	Rational	Example
Antibiotics	Target specific members or groups of the microbiota and suppress them, allowing expansion of desirable species	Small intestinal bacterial overgrowth Hepatic encephalopathy
Bacteriophages	Use naturally occurring bacterial viruses to target specific members of a community that are disruptive or are carrying out pathogenic processes	Pathogen targeted therapy to spare the microbiota from collateral damage
Probiotics (single species)	Replace a presumably missing organism and thus a missing function in the form of an organism	Sacchromyces boulardii for prevention of antibiotic associated diarrhea
Multispecies/designer communities	As with single species probiotics, use a collection of organisms to replace a missing function in the microbiome	Feces derived communities, multispecies
Prebiotics	Supply a complex food product (often carbohydrates) that is not digestible by the host to stimulate specific members of the microbiota. The prebiotic is meant to be metabolized by the microbiota to compounds beneficial to the host	Inulin, resistant starch
Synbiotics	Supply a complex of a microbe or microbes along with a prebiotic that is meant to be used by these organisms replace a missing function in a microbiome	
Nutritional therapy	Complete redesign of a diet to promote beneficial microbial communities and function	Low FODMAP diet for IBS. Exclusive enteral nutritional therapy for IBD
Community replacement, "microbiota restoration"	To restore a "deficient" microbiota, harvest a presumably normal microbiota from a healthy person and administer it to a patient	Fecal transplantation

FODMAP=fermentable oligosaccharides, disaccharides, monosaccharides, and polyols; IBS=irritable bowel syndrome; IBD=inflammatory bowel disease.



Table 2 Pipeline o	n microbiome derived therapeutics in inflammatory bowel disease and syste	emic autoimmunity	
Company	Microbiota product and indication	Development stage	Pharmaceutical partner/s
AvidBiotics	Targeted antibacterial bacteriocins (Avidocin/Purocin)	Preclinical	DuPont Nutrition & Health
CIPAC	Standardized approach to FMT for CDI and IBD (Full-Spectrum Microbiota)	Undisclosed	2 <u>4</u> 1
Enterome	Anti-Escherichia coli small molecule for IBD (EB-8018; EB110)	Phasel	Takeda, Janssen, Bristol-Myers Squibb, Nestlé
4D Pharma	Therapies from microbiome based molecules for IBS and IBD indication (Blautix; Thetanix)	Phasel	Publicly traded
Rebiotix Inc	Prescreened stool offered to health providers for FMT (microbiota restoration therapy for recurrent CDI; RBX2660)	Phase III (FDA breakthrough therapy designation for CDI); phase I (pediatric UC)	Private
OselInc	Single strains of native and genetically engineered bacteria for urogenital and gastrointestinal disease indications (Lactin V; CBM588)	Phase II	Private
Second Genome	Application of microbiome science for discovery of new therapies (eg, IBD; SGM-1019)	Phasel	Janssen, Pfizer, Roche, Monsanto
Seres Health	Therapeutics to catalyze restoration of healthy microbiome in CDI (SER-109) and UC (SER-287)	Phase III (FDA orphan drug designation for SER-109); phase I (SER-287)	Nestlé Health Science, publicly traded
Symbiotix	Bacteroides fragilis derived polysaccharide A for IBD and multiple sclerosis	Preclinical	
Vedanta Biosciences	Human microbiome consortia (Clostridia cocktail for IBD/allergy indications; VE-202)	Phase I/II	Janssen
CD⊨Clostridium difficile	infection: FDA=Food and Drug Administration: FMT=fecal microbiota transplant: IBD=inflammat	ory bowel disease: IBS=irritable bowel syndrom	e: LIC=ulcerative colitis

Clemente et al., 2018

