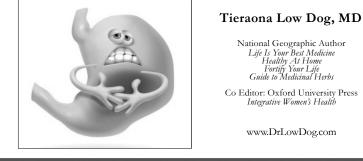
Gut Check: Understanding the Microbiome



Vieraona Low Dog, M.D.

Objectives

1. Identify examples of how diet, lifestyle, and the environment influence the human microbiome.

2. Discuss the relationship between the oral microbiota and health.

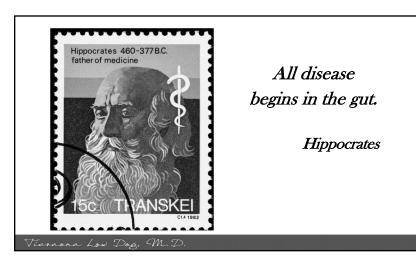
3. Discuss the relationship between the gut microbiota and health.

4. Identify how certain medications, such as proton pump inhibitors and antibiotics, impact the microbiota.

5. Describe the role of diet, dietary fiber, prebiotics, and probiotics in optimizing the microbiota.

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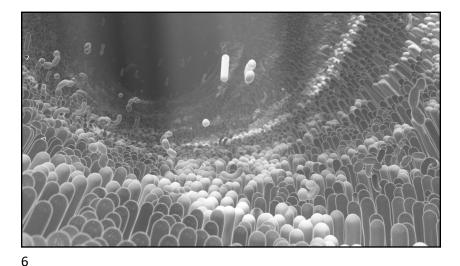




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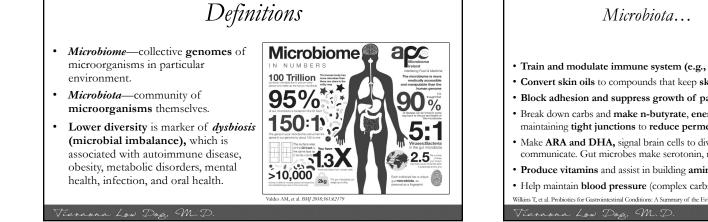
Human Microbiome Project

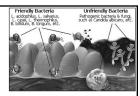
- · Massive research initiative of NIH cataloging the microorganisms living in and on our body starting in 2007.
- Led to rapidly growing appreciation for incredible and diverse impact these organisms have on our health and well-being.
- · Gut bacteria produce vitamins and break down our food; their presence or absence has been linked to obesity, inflammatory bowel disease, IBS, anxiety, depression, food allergies, neuroinflammation, GI infections, high blood pressure, diabetes, metabolic syndrome, and more.
- Our resiliency—our ability to recover quickly from stressors—may be a function of which bacteria inhabit or don't inhabit our gut.







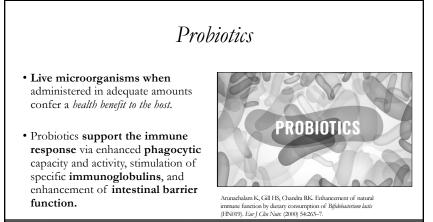




- Train and modulate immune system (e.g., skin, gut).
- · Convert skin oils to compounds that keep skin supple and lower pH.
- · Block adhesion and suppress growth of pathogenic bacteria.
- · Break down carbs and make n-butyrate, energy for intestinal cells, but also crucial for maintaining tight junctions to reduce permeability.
- · Make ARA and DHA, signal brain cells to divide (infants). Gut and brain neurons communicate. Gut microbes make serotonin, melatonin, GABA, and others.
- · Produce vitamins and assist in building amino acids.

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• Help maintain **blood pressure** (complex carbs — formate — impact salt processing) Wilkins T, et al. Probiotics for Gastrointestinal Conditions: A Summary of the Evidence. Am Fam Physician 2017 Aug 1;96(3):170-178.



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Obesity Metabolic Syndrome

Diabetes C. difficile Infection

Colorectal Cancer

Diseases

Allergies

Psoriasis

Skin Cancer

Atopic Dermatitis

Ectodermal Dysplasia

Acne

Inflammatory Bowe

Psychiatric Disorders

Actinobacter

Bacteroide

Fusobacteria

Proteobacter

Gut Microbiom

Gut stool

Skin Microbiom

Skin: antecubital fossa

Oral Microbiom

Mouth: tonsils

Placenta Microt

Placenta: term birth

Vagina Microbiome

Caries Periodontal Diseases

Gingivitis

Pre-term Birth Chorioamnionitis

TORCH Infections

Sexually Transmitted

Graphic from:

Belizario, JE, et al. Front Microbiol

2015; October 6

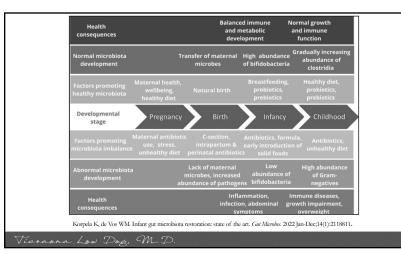
Villitis

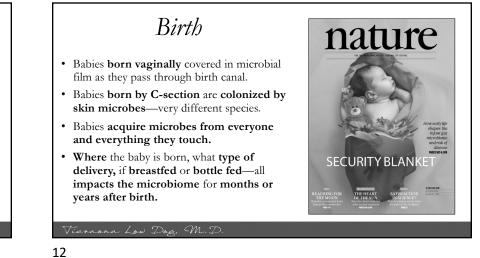
Vaginosis

Diseases

Yeast Infection

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Neonatal Microbiome



- Greatest insults to the natural assembly of neonatal microbiome: C-section delivery, antibiotic use, and formula feeding.
- Differences in microbial species observed between C-section and vaginally delivered babies up to 7 years after birth.¹
- Intrapartum antibiotics associated with lower abundance of good bacteria (Lactobacilli and Bifidobacterium) in neonatal gut.²
- Formula feeding associated with increased prevalence of pathogenic bacteria (C. difficile, Bacteroides fragilis, E. coli) and decreased prevalence of Bifidobacterium.³

Salminen S, et al. Gut. 2004;53:1388–1389; 2. Aloisio I, et al. Appl Microbial Biotechnol. 2014;98:6051–6060.
Mueller NT. et al. Trends Mol Med 2015; 21(2): 109-17

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Probiotics & Birth Mode

- Mothers given probiotic or placebo during pregnancy and then infants given same.
 - **Placebo group:** birth mode and/or antibiotics significantly altered microbiota composition and function, reducing *Bifidobacterium*.
 - Probiotic group: effects of birth mode and/or antibiotics either completely eliminated or dramatically reduced.

(Probiotic: Bifidobacterium breve, Propionibacterium freundenreichii subsp. shermanii JS, Lactobacillus rhamnosus Le705, and L. rhamnosus GG)

Korpela K, et al. Probiotic Supplementation Restores Normal Microbiota Composition and Function in Antibiotic-Treated and in Caesarean-Born Infants. *Microbiome* 2018; 6(1): 182

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Breast Milk.

- Prospective, 12-month longitudinal study, bacterial composition identified in breast milk, areolar skin, and infant stool samples of 107 healthy mother-infant pairs in Los Angeles, California and St Petersburg, Florida.
- During first 30 days of life, infants who obtained 75% or more of daily milk intake from breastmilk received 27.7% of the bacteria from breast milk and 10.3% from arcolar skin – almost 40% of bacteria was from mother to infant.
- Infant gut microbial communities more closely related to infant's mother's milk and skin compared with random mother (P < .001).
- Bacterial diversity (p=0.003) and composition changes associated with proportion of daily breast milk intake in a dose-dependent manner, even after the introduction of solid foods.
- Breastfeeding confers protection against respiratory and gastrointestinal tract infections and allergic diseases, and reduces risk of chronic diseases (e.g., diabetes, obesity, and inflammatory bowel disease). Likely through modulation of microbiome. Buttinger K, et al. JAMA Philar. 2017 Jul 1;171(7):647-654.

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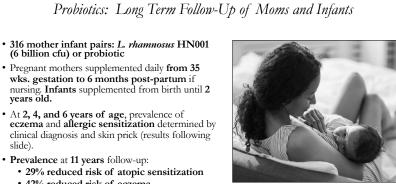
Birth to 3 Years

- Within weeks, microbial specialization occurs. Different populations in mouth, gut, skin, etc.
- Microbial populations in infant are similar to people they live with.
- Microbiota dramatically altered by **new foods**, **antibiotics**, **PPI use**, etc.
- Number and types of species increase and change with age.
 - Example: babies have more folate *producing* microbes—adults have more folate *barvesting* microbes.

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Azad MB, et al. Gut microbiota of healthy Canadian infants: profiles by mode of delivery and infant diet at 4 months. *Can Medical Association Journal*, 2013; 185(5), 385-394.



- 42% reduced risk of eczema
- 24% reduced risk of wheeze



Wickens K, et al. Clin Exp Allergy 2013; 43(9):1048-57. Wickens K, et al. Pediatr Allergy Immunol 2018; 29(8): 808-14



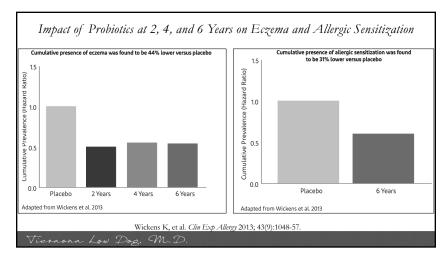


Age 3 to Old Age

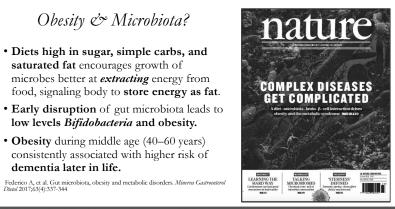
- Microbiome becomes stable by age 3-5.
- · Even with disruptions (medications, disease, dietary changes)-it usually returns to baseline.
- After age 65, microbe populations decrease, species become more similar (dysbiosis), may explain/contribute to some diseases of aging.

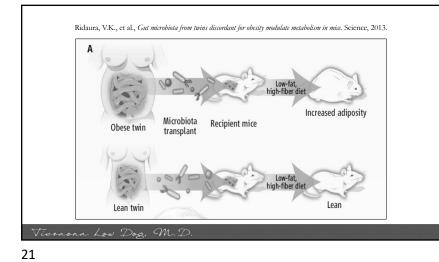
Yatsunenko T, et al. Human gut microbiome viewed across age and geography. Nature 2012; 486:222-228. The Human Microbiome Project Consortium (2012). Structure, function and diversity of the healthy human microbiome. Nature 2012; 86, 207-214.

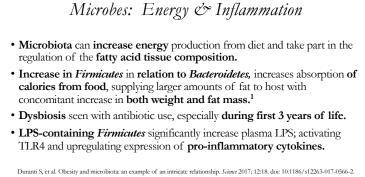




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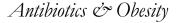
ntt S, et al. Obesity and microbiota: an example of an intricate relationship. Science 2017; 12:18. doi: 10.1186/s12265-017-0566 Fessler MB, et al. Curr Opin Lipidal 2009; DOI: 10.1097/MOL.0b013e32832fa5c4

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Title of the study	Year	Subjects of the study	Final result(s) gathered	Reference	
Childhood overweight after establishment of the gut microbiota: the role of delivery mode, pre-pregnancy weight and early administration of antibiotics.	2011	28354 mother-child	Antibiotics in infancy influences the risk of overweight in later childhood	Ajslev et al., 2011	
Infant antibiotic exposures and early-life body mass.	2013	11532 children	Exposure to antibiotics during the first 6 months of life was associated with increases in body mass.	Trasande et al., 2013	
Antibiotic treatment during infancy and increased body mass index in boys: an international cross-sectional study.	2014	74946 children	Exposure to antibiotics during the first 12 months of life is associated with a small increase in BMI in boys aged 5–8 years	Murphy et al., 2014	
Infant antibiotic exposure and the development of childhood overweight and central adiposity	2014	1047 children	Antibiotic use in the first year of life was associated with overweight	Azad et al., 2014	
Association of antibiotics in infancy with early childhood obesity.	2014	64580 children	Repeated exposure to broad-spectrum antibiotics was associated with early childhood obesity	Bailey et al., 2014	
Prenatal exposure to antibiotics, cesarean section and risk of childhood obesity.	2015	436 mother-child dyads	Exposure to antibiotics in the second or third trimester of pregnancy were associated with higher risk of childhood obesity.	Mueller et al., 2015	
Prenatal exposure to systemic antibacterials and overweight and obesity in Danish schoolchildren: a prevalence study.	2015	9886 children	Prenatal exposure to systemic antibacterials was associated with an increased risk of overweight and obesity at school age	Mor et al., 2015	
Antibiotic exposure in infancy and risk of being overweight in the first 24 months of life.	2015	6114 boys and 5948 girls	Antibiotic exposure before 6 months was associated with increased body mass	Saari et al., 2015	
Early Life Antibiotic Exposure and Weight Development in Children.	2016	979 children	Repeated exposure to antibiotics early in life, especially β-lactam agents, is associated with increased weight and height.	Mbakwa et al., 2016	Del Fiol FS, et al. Obesity: A new adverse
Antibiotic Use and Childhood Body Mass Index Trajectory.	2016	142824 children	Body Mass Index increase	Schwartz et al., 2016	effect of antibiotics? Front Pharmacol 2018:
Administration of Antibiotics to Children Before Age 2 Years Increases Risk for Childhood Obesity.	2016	21714 children	Administration of 3 or more courses of antibiotics before age of 2 years was associated with an increased risk of early childhood obesity	Scott et al., 2016	https://doi.org/10.338 fphar.2018.01408

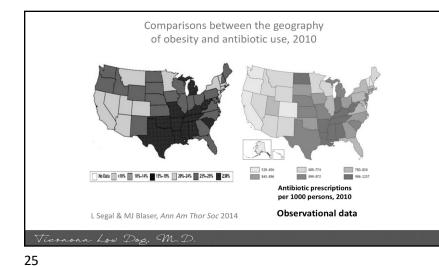




- American children up to 2 years of age, on average receive 3 full doses of antibiotics; up to 10 years of age received 10 full doses; and 17 full doses antibiotic by age 20.¹
- Four or more courses of antibiotics given between ages 2 to 3 years independently associated with obesity at age 5. (OR: 1.6).²

 Cox I.M. Antibiotics in early life and obesity. Nat. Rev. Endocrinol 2015; 11, 182–190.
Kelly D, et al. Antibiotic use in early childhood and risk of obesity: longitudinal analysis of a national cohort. World J Pediatrics 2019;15(4):390-397.

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Antibiotics and Microbes

- Disrupt existing microbiota; linked to antibioticassociated diarrhea, pseudomembranous colitis, and increased susceptibility to subsequent disease.¹
- Extent of change depends on antibiotic type, duration, and dose.
- Systematic review: changes in gut microbiome from metronidazole and clarithromycin lasted longest (4 years), clindamycin (2 years), and ciprofloxacin (1 year).²

 Abeles SR, et al. Microbial diversity in individuals and their household contacts following typical antibiotic courses. *Minrhimm* 2016; 4: 39–51.
Zimmermann P, Curtis N. The effect of antibiotics on the composition of the intestinal microbiota - a systematic review. *J Inf Scars*. 2019;79(6):471–89.

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Antibiotic ProphylaxisImage: Subscript of the state of the sta

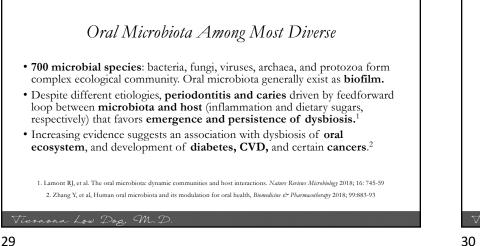
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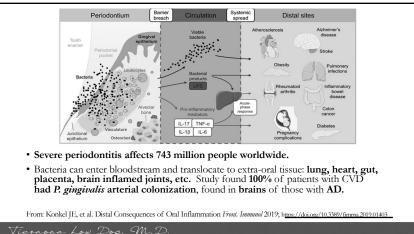
Oral Health & Systemic Disease

- **1891:** first oral microbiologist Willoughby D. Miller put forward theory of **oral focal infections**, suggesting that **oral microbial infection** can affect other parts of the body, related to a **variety of systemic diseases**.
- 1912: Frank Billings speculated that infection of the teeth may be the cause of rheumatoid arthritis, nephritis, endocarditis, and other diseases.
- Periodontal inflammation leads to loss of connective tissues/bones. Extensive inflammatory cell infiltration appears in connective tissue near periodontal pocket epithelium. This low-grade inflammation may disturb the health of body or worsen other systemic diseases.

Miller WD. The human mouth as a focus of infection. Lanat 1891; 138, 340–342 Billings E Chronic focal infections and their etiologic relations to arthritis and nephritis. Arch. Intern. Med 1912; IX, 484–498 (1912).

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Oral Microbiota & Gut Inflammation

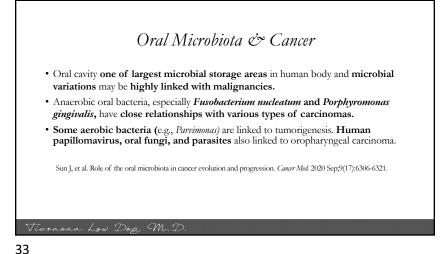
- Adults produce >1000 mL/d of saliva, carrying oral microbes to the GI tract. Bacteria can also enter GI tract via bloodstream.
- Inflammation caused by P. gingivalis in oral cavity can alter intestinal microbial communities, disrupt intestinal barrier, induce endotoxemia, and trigger a systemic inflammatory response.
- F. nucleatum can migrate to intestine, inhibiting the immune response mediated by T cells, and promoting progression of IBD.

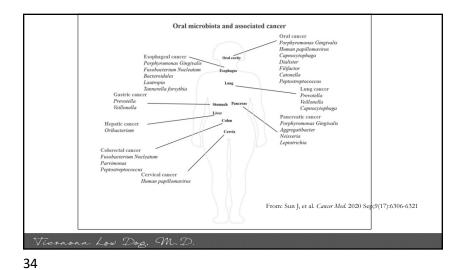
Peng X, et al. International Journal of Oral Science 2022; 14, 14.

Oral Microbiota & Cardiovascular Disease

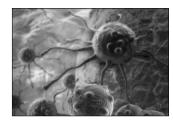
- Cross-sectional studies, case analyses, epidemiological investigations: periodontitis is important risk factor for CVD.
- Periodontal disease-related bacteria stimulate cells to produce inflammatory factors (e.g., IL-1 β , IL-6, TNF- α) and enter circulation from damaged periodontal tissue, resulting in inflammation and vascular endothelial damage & formation of atherosclerotic plaques.
- After periodontal treatment, **CRP and other inflammatory markers** are significantly reduced, further linking oral health to CVD.

Herrera D, et al. Periodontology 2020; 83: 66-89 Schoffer C. J. Periodontol 2021; 92: 793-802





Esophageal Cancer



Gao, S, et al. Infect Agent Cancer 2016; 11: 3–12. Chen, X. et al. PLoS ONE 2015; 10: e0143603.

• Esophageal squamous cell carcinoma closely related to tooth loss and tooth brushing frequency. Incidence of metastasis in periodontitis patients significantly higher than that in non-periodontal patients

- P. gingivalis detected
 - 61% cancerous tissues
 - 12% adjacent tissues
- 0% of normal esophageal mucosa.
- *F. nucleatum* is also a promotor.

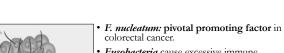
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Nosho K, et al. World J Gastroenterol 2016; 22: 557-566

Liu Y, et al. J Gastroenterol 2019 Jan;54(1):33-41



Colorectal Cancer

- *Fusobacteria* cause excessive immune response, turning on cancer growth genes in CR cancer.¹
- Have specific surface molecules allowing them to attach and invade colorectal cells.¹
- Higher levels *F. nucleatum* DNA in tumor, associated with poorer survival.
- May contribute to **chemo-resistance in GI cancers.**²

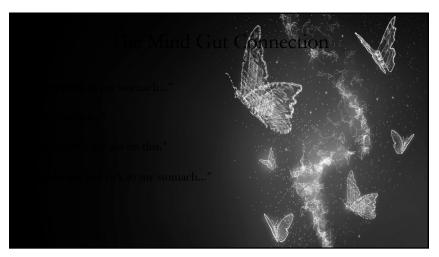


Pancreatic Cancer

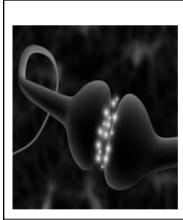
- Meta-analysis 8 studies: RR for periodontitis and pancreatic cancer was 1.74 (95% CI 1.41-2.15) and 1.54 for edentulism (95% CI 1.16-2.05). No heterogeneity or publication bias. Reports from 3 continents, association is generalizable.
- Associations between pancreatic cancer and abundance of *P. gingivalis* in oral wash samples, and pancreatic cancer and increased levels of antibodies against *P. gingivalis* have been well-documented.

Maisonneuve P, et al. Periodontal disease, edentulism, and pancreatic cancer: a meta-analysis. Ann Onu/2017 May 128(5):985-995 Charg, JS, et al. Investigating the association between periodontal disease and risk of pancreatic cancer Panara 2016; 45:134-141. Fan X et al. Human onal microbiome and prospective risk for pancreatic cancer: a population-based neural case-control study. (cm: 2018 Jan¢7(1):12)-127.

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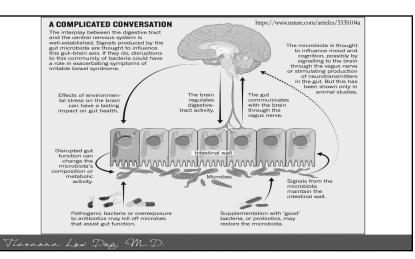
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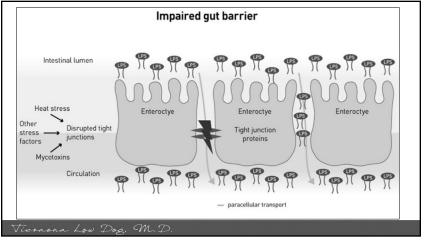


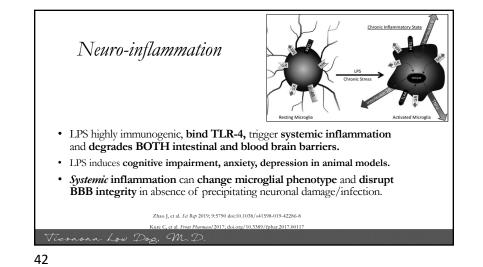
Gut-Brain Communication

- Gut bacteria produce neurotransmitters: dopamine, serotonin, norepinephrine, GABA, acetylcholine, melatonin; critical for mood, sleep, anxiety, concentration, reward, and motivation. Acts in many ways like an endocrine-like organ.
- Gut microbiota can cause changes in how our brains react to events/stressors
- Serotonin associated with **depression and happiness**; *90%* is made in the digestive tract—not the brain.
- Gut microbiota regulate brain function through gut-brain axis, and *dysbiosis* may trigger anxiety and depression.

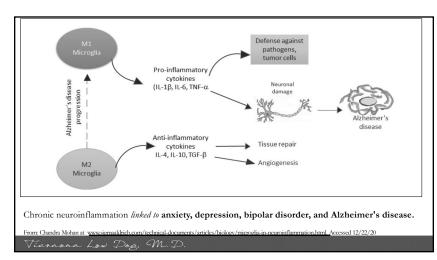
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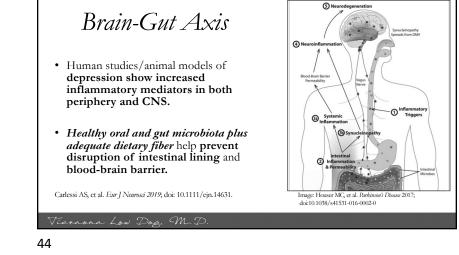


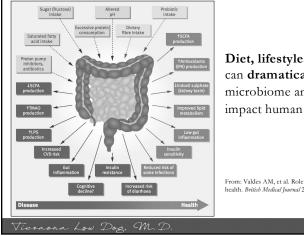












Diet, lifestyle, and medications can dramatically impact the microbiome and ultimately impact human health.

From: Valdes AM, et al. Role of gut microbiota in nutrition and health. British Medical Journal 2018;361:j2179

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More Fiber, Less Sugar • Diets high in fiber and low in sugar increase Bifidobacteria, preventing toxins from passing through intestinal wall into bloodstream.

• Strong evidence for optimizing intestinal barrier function with

dietary fiber. • Aim for 25-35 grams/d

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Impact of Certain Diets

- 21 healthy people had substantially altered gut microbiota profiles after four weeks on gluten-free diet; significant reduction in key beneficial microbe species.
- Low FODMAP diets lead to significant reduction in Bifidobacterium and profound changes in the microbiota and metabolome; duration and clinical relevance are not known.



diet on the human gut microbiome. Genome Med 2016;8:45 McIntosh K, et al. FODMAPs alter symptoms and the metabolome of patients with IBS: a randomised controlled trial. Gut 2017;66:1241-51.

Table 1 | Examples of foods, nutrients, and dietary patterns that influence human health linked to their effect Effect on health outcomes mediated by **Dietary element** Effect on gut microbiome gut microbiome Low FODMAP diet Low FODMAP diet increased Actinobacteria; high FODMAP diet Reduced symptoms of irritable bowel decreased abundance of bacteria involved in gas consumption58 syndrome⁵ Cheese Increased Bifidobacteria, 97.98 which are known for their positive Potential protection against health benefits to their host through their metabolic activities.9 pathogens.¹⁰⁰ Increased production of Decrease in Bacteroides and Clostridia, some strains of which are SCFA and reduced production of TMAO⁹¹ associated with intestinal infections⁵ Increased microbiota diversity and SCFA production²²¹⁰¹¹⁰² Fibre and prebiotics Reduced type 2 diabetes22 and cardiovascular disease10 Induced glucose intolerance¹⁰ Overgrowth of Proteobacteria and Escherichia coli.104 Bacteroides, Artificial sweeteners Clostridia, and total aerobic bacteria were significantly lower, and faecal pH was significantly higher⁴ Polyphenols (eg, from Increased intestinal barrier protectors (Bifidobacteria and Gut micro-organisms alter polyphenol tea, coffee, berries, Lactobacillus), butyrate producing bacteria (Faecalibacterium bioavailability resulting in reduction prausnitzii and Roseburia) and Bacteroides vulgatus and and vegetables such of metabolic syndrome markers and as artichokes, olives, Akkermansia muciniphila.107 Decreased lipopolysaccharide cardiovascular risk markers¹⁰⁸ and asparagus) producers (E coli and Enterobacter cloacae)10 Very modest differences in composition and diversity in humans Vegan Some studies show benefit of vegetarian over omnivore diet, 109 others fail to find a and strong differences in metabolomic profile compared with omnivore diet in humans50 difference¹ Valdes AM, et al. Role of gut microbiota in nutrition and health. British Medical Journal 2018;361:j2179



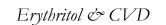
Nettleton JE, et al. Reshaping the gut microbiota: Impact of low calorie sweeteners and the link to insulin resistance? *Physiol Behav* 2016;164(Pt B):488-93.

Ruiz-Ojeda FJ, et al. Effects of Sweeteners on the Gut Microbiota: A Review of Experimental Studies and Clinical Trials, *Adv Nutr* 2019; 10(1): \$31-48

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Sugar Substitutes

- Sugar substitutes frequently *1000 times sweeter* than sucrose.
- Despite GRAS status by regulatory agencies, sugar substitutes can have negative effects on gut microbiota.
- Sucralose, saccharin, stevia—all shown to disrupt balance and diversity of gut microbiota.
- Erythritol, mannitol, sorbitol have no adverse effect.



- As a sweetener, levels > 1,000-fold greater than levels found naturally in foods.
- Increasing blood erythritol levels speeds up blood clot formation and artery blockage in mice.
- NIH-funded research: people with highest erythritol levels (top 25%) were twice as likely to have cardiovascular events over three years of follow-up as those with the lowest (bottom 25%).
- Blood erythritol levels measured in 8 healthy volunteers after drinking a beverage sweetened with erythritol. Erythritol levels increased 1,000-fold and remained substantially elevated for several days. For at least two days, levels high enough to trigger changes in platelet function.

Witkowski, M, et al. Nature Medicine 2023 https://doi.org/10.1038/s41591-023-02223-9

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PPIs, Dysbiosis, and Infection

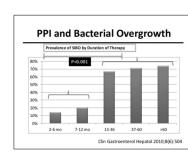
- Stomach acid directly destroys harmful pathogens.
- When acid is shut down, ~50% of salivary and ingested bacteria survive by slipping past this "gastric acid trap."
- Translocated bacteria disrupt gut microbiota, leading to dysbiosis, SIBO, and dyspepsia.
- 70% of immune system resides in GI tract: critical line of defense.
- By altering balance between beneficial and pathogenic microbes, the risk for infection is increased.

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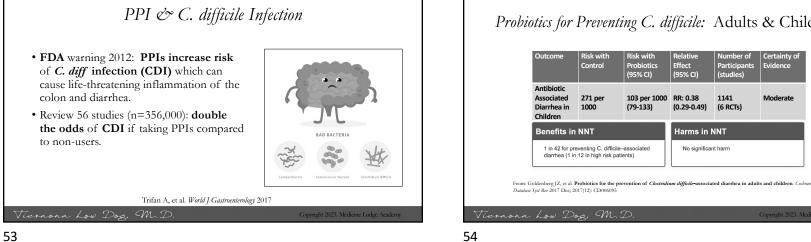
Dysbiosis & SIBO

- Meta analysis 19 studies (n=7055): statistically significant association between increased risk of SIBO and PPI use (OR 1.71).¹
- Dysbiosis and SIBO increase intestinal permeability, allowing bacteria and other substances to pass directly through the intestinal mucosa into the blood stream.
- PPIs may have more prominent effect on microbiota composition on population basis than any other drug.²

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1. Su T, et al. J Gastroenterol 2018; Jan;53(1):27-36 2. Imhann F, et al. Gut. 2016; 65: 740–748



Probiotics for Preventing C. difficile: Adults & Children

(95% CI)

RR: 0.38

(0.29-0.49)

Harms in NNT

No significant harm

1141

(6 RCTs)

Moderate

(95% CI)

(79-133)

103 per 1000

Risk wit Control

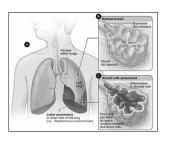
271 per

1000

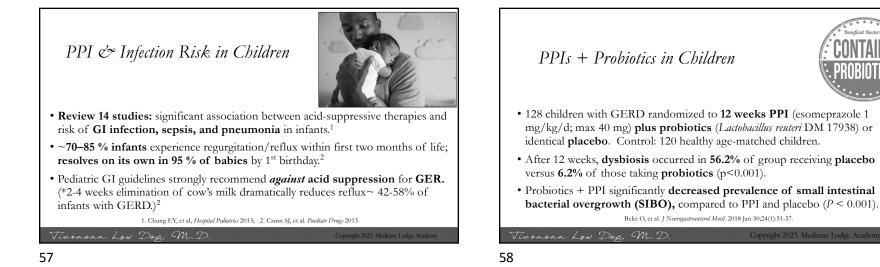
	PPIs & SARS-CoV-2
	• GI tract significant portal of entry for virus. Bind to widely expressed ACE-2 receptors in intestine, replicating rapidly.
	No stomach acid means no viral inactivation.
	• Korean study: current use of PPIs conferred 79% greater risk of severe clinical outcomes; risk climbed to 90% if PPI use started within 30 days of confirmed Covid 19 infection. ¹
•	• US study: mortality from COVID-19 was 2.3 times higher in PPI users, compared to non-users. ²
	1. Lee SW, et al. <i>Gut</i> 2021
	2. Ramachandran P, et al. Eur J Gastroenterol Hepatol 2021
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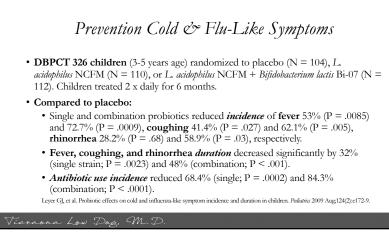
PPI & Community Acquired Pneumonia

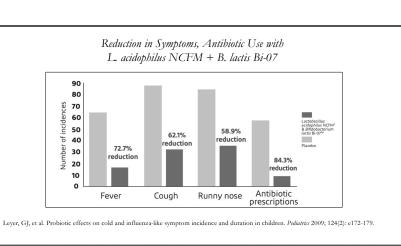
- Without adequate stomach acid, there can be overgrowth of oropharyngeal bacteria, which can increase the risk for infection.
- Review of 26 studies: 1.5-fold increase in risk for community-acquired pneumonia, with the highest risk occurring within 30 days of starting PPI.



Lambert AA, et al. PloS One 2015



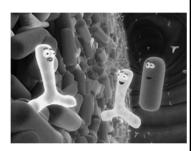




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Acute Infectious Diarrhea

- High quality evidence support probiotics in acute infectious diarrhea, common for those traveling, kids going to daycare, etc. Note: start probiotics first sign of diarrhea and 1-2 weeks beyond; if traveling, start 2 days before travel and duration of trip.¹
- Meta-analysis **17 RCTs** (2,102 children): significant **reduction in duration** of diarrhea with *S. boulardii* use (20 fewer hours).²
- Meta-analysis **8 RCTs** (1,229 children): *L renteri* DSM 17938 reduced duration of diarrhea (25 fewer hours), increased cure rate on days 1 and 2.³

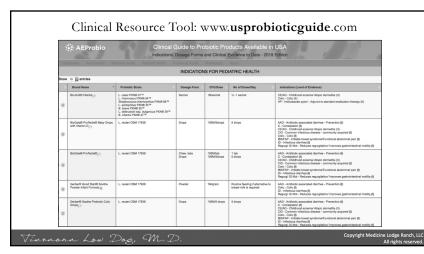


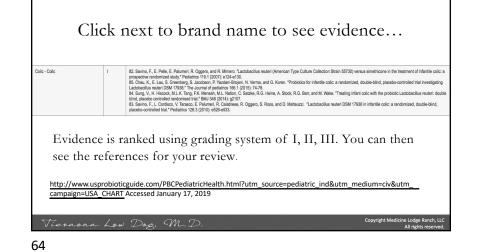
- https://www.aafp.org/afp/2017/0801/p170.html, Accessed December
- 22, 2020 Feizizadeh S, et al. Efficacy and safety of *Saccharomyces boulardii* for acut diarrhea. *Pediatriss*. 2014;134(1):e176–e191.
- Urbańska M, et al. Systematic review with meta-analysis: Ladobacillus ratteri DSM 17938 for diarrhoeal diseases in children. Aliment Pharmaco Ther. 2016;43(10):1025–1034.

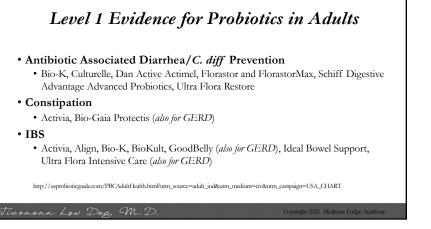
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Outcome	Reference	No of studies/ participants	Evidence of benefit?	Prevention and treatment of	Saez Lara et al (2015) ¹¹²	14/821 ulcerative colitis	Yes]
Clostridium difficile associated	Goldenberg et al (2017) ¹¹¹	39/9955	Yes	Crohn's disease and ulcerative colitis		8/374 Crohn's disease		
liarrhoea in adults				Pulmonary exacerbations in	Ananathan et al (2016) ¹²³	9/275	Yes	
Necrotising enterocolitis	Al Faleh et al (2014) ¹¹² Rees et al (2017) ¹¹³	17/5338	Yes	children with cystic fibrosis				From: Valdes AM, et al. Rol of gut microbiota in nutrition and health. <i>BMJ</i>
Antibiotic associated diarrhoea in children	Goldenberg et al (2015) ¹¹⁴	26/3898	Yes	Type 2 clabetes Abarr et al (2016) ¹²⁴ (filtering glucose, p)/clabetes p)/cla	Akbari et al (2016) ¹²⁶	13/805	Yes	
Probiotics for preventing acute upper respiratory	Hao et al (2015) ¹¹⁵	12/3720	Yes			7/425	Yes	
tract infections Urinary tract infections	Schwenger et al (2015) ¹¹⁶	9/735	No		Athalye-Jape et al (2016) ¹²⁶	6/1778	Yes	
Prevention of asthma and wheeze in infants	Azad et al (2013) ¹¹⁷	6/1364	No		Mazidi et al (2017) ¹²⁷	19/935	Yes	
Prevention of eczema in infants and children	Mansfield et al (2014)	16/2797	Yes	cardiovascular risk factors in patients	Hendijani et al (2017) ¹²⁸	11/641	Yes	
Prevention of invasive fungal	Agrawal et al (2015) ¹¹⁹ 19/4912 Unc	Unclear	with trans 7 dishetses				2018;361:j21	
infections in preterm neonates				Reduction of total cholesterol and low	Wu et al (2017) ¹²⁹	15/976	Yes	
Prevention of nosocomial infections	Manzanares et al (2015) ¹²⁰	30/2972	Yes	density lipoprotein cholesterol				_
Intections .	Ahmadi et al (2015) ¹²¹	14/1149	Yes	Depressive symptoms	Wallace and Milev (2017) ^{79,130}	6/1080	Yes	
rotavirus diarrhoea in infants and children				Vulvovaginal candidiasis in non- pregnant women	Xie et al (2018) ¹³¹	10/1656	Yes	

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Level 1 Evidence for Probiotics in Children

• Antibiotic Associated Diarrhea/C. diff Prevention

 BioGaia® Protectis, Culturelle Kids, FlorastorKids, Gerber Good Start Grow Toddler Probiotic + Gentle Infant Formula + Soothe Vitamin D and Probiotic Drops, Pedia-Lax® Probiotic Yums

Constipation

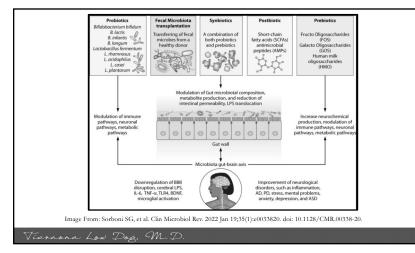
 BioGaia® Protectis, Gerber Good Start Grow Toddler Probiotic + Gentle Infant Formula + Soothe Vitamin D and Probiotic Drops, Pedia-Lax® Probiotic Yums

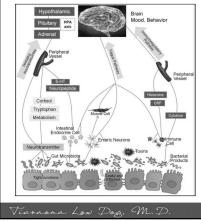
IBS/Functional Abdominal Pain

 BioGaia® Protectis, Culturelle Kids, Gerber Good Start Grow Toddler Probiotic + Gentle Infant Formula + Soothe Vitamin D and Probiotic Drops, Pedia-Lax® Probiotic Yums, Visbiome + Extra Strength

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Sleep & Stress

- Disruption of circadian rhythm alters gut microbiome equilibrium. Microbes and humans *share* circadian clock.
- Emotional and physiological stress negatively affect gut microorganisms, impacting immune and nervous systems.
- Lactobacillus and Bifidobacterium probiotic strains improve stress response.

Farre N, et al. Sleep and circadian alterations and the gut microbiome: associations or causality. *Current Sleep Mult Report* 2018, 4(1):50-57 Li, Y, et al. The role of microbiome in insomnia, circadian disturbance and depression. *From Psychiatr* 2018, doi: 10.3389/fpst/2018.00669

