What Happens In The Gut Does Not Stay In the Gut -

The influence of gut microbiome on dry eye disease

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Disclosures

- B&L
- BioTissue
- J&J
- Lumenis
- Sight Sciences
- SUN

Agenda



Review gut health, the microbiome, and dysbiosis.



Inform on the evidence of the relationship between the gut microbiome and dry eye disease



Discuss how ECPs can educate patients on the importance of gut health and nutritional choices that support their eye health and overall wellbeing.

Inflammation

Malabsorption

Barrier Breakdown

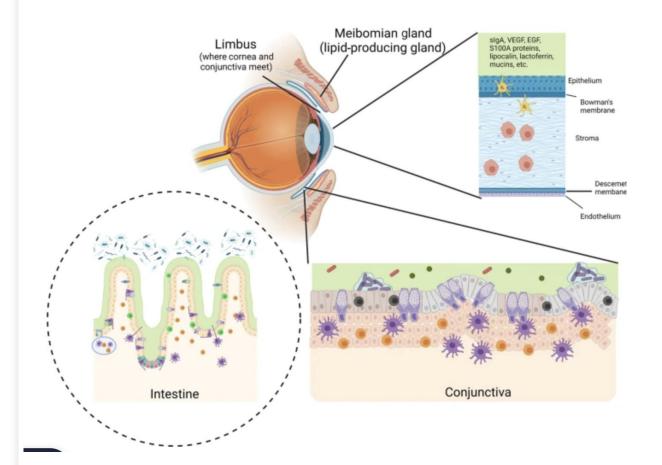
Pathogens and Biofilm

Toxins

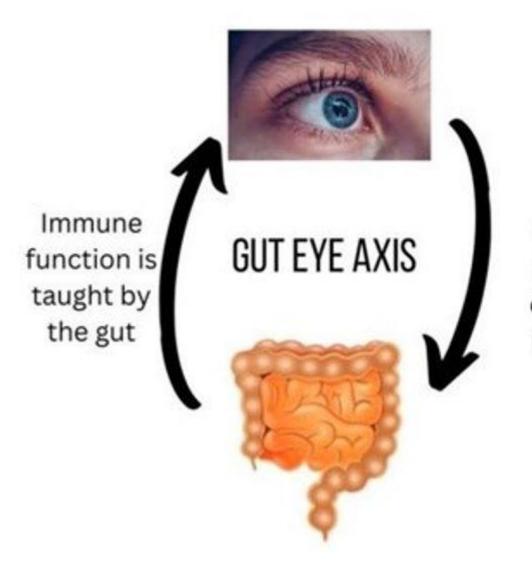
Food Reactions

Enteric Nervous System Dysfunction

Gastrointestinal Environment



- Similar function of mucins and glycoproteins on the ocular surface and the GI tract.
- They share specialized cells and the signaling cascades evolved to maintain environmental defenses in mucosal layers, an example being innate lymphoid cells (ILC's).



80% of Immune cells live in the GI tract • Chronic inflammation in the gut can lead to release of TNF-a, IL-6, IL-1β

 Modulating the gut microbiome has been shown to also modulate the proteins expressed by the lacrimal glands, with IL-1β and IL-6 decreasing, and IL-10 increasing

Immune Branches and Balance



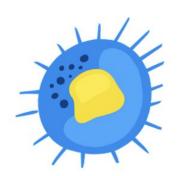
T Helper Type 1

Bacteria and viruses, intracellular



T Helper Type 2

Parasites, allergens, mast cells & histamine



T Helper Regulatory

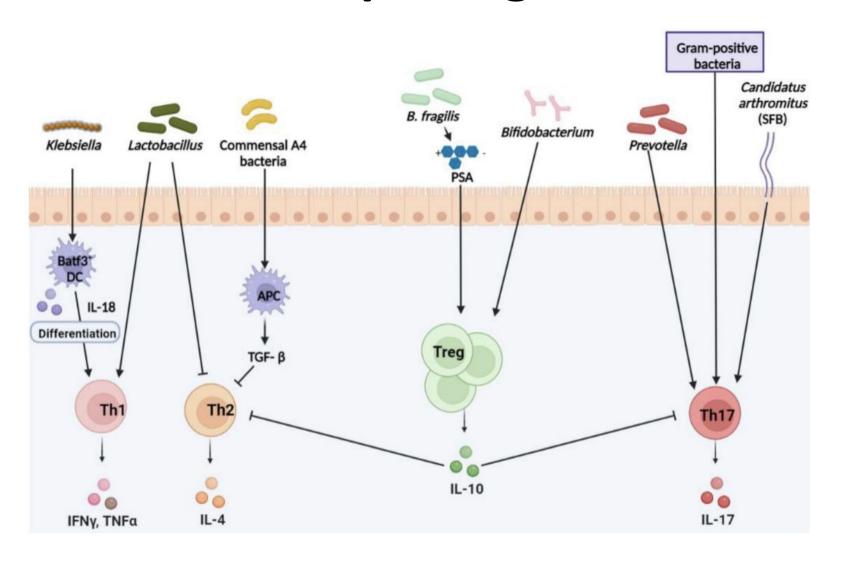
Immune suppression & tolerance

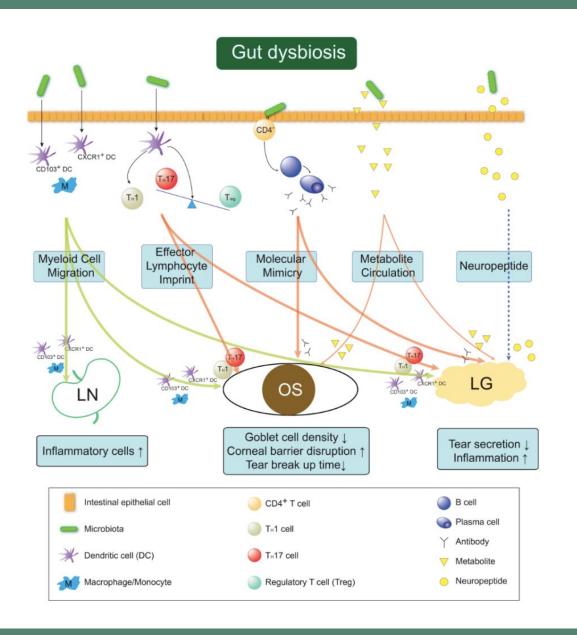


T Helper Type 17

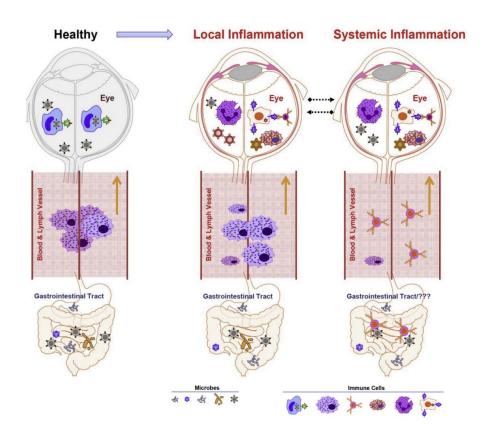
Mucosal immunity, autoimmune tissue damage

Microbiota mediates T cell differentiation in homeostatic and pathogenic conditions





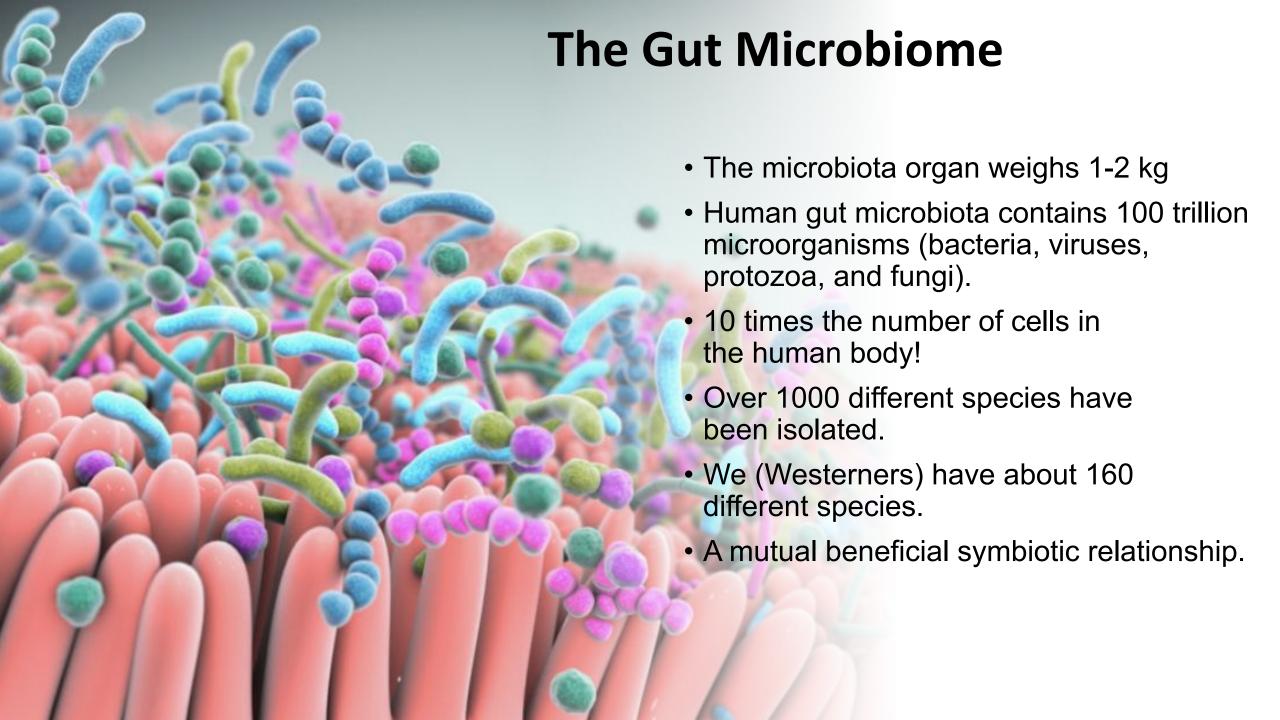
Intraocular Inflammation



Gut Dysbiosis and Ocular Disease

- AMD
- Glaucoma
- Diabetic Retinopathy
- Uveitis
- Dry Eye Disease

Wen, X., Hu, X., Miao, L., Ge, X., Deng, Y., Bible, P. W., & Wei, L. (2018). Epigenetics, microbiota, and intraocular inflammation: New paradigms of immune regulation in the eye. *Progress in retinal and eye research*, 64, 84–95.



The Roles of the Human Microbiota

- Gut motility
- Modulates the immune system
 - IgE
- Impact on inflammation
- Regulates metabolism and glucose regulation
- Mood management
- Improves nutritional status
 - B vitamins
 - Vitamin K
 - Mineral absorption Ca, Mg, Zn
 - Production of SCFAs

Short Chain Fatty Acids (SCFAs)

- The most common SCFAs include acetate, propionate, and butyrate
- Group of organic compounds that are produced by the fermentation of dietary fibers by gut bacteria in the colon
- Improve gut barrier function, increase gut motility, and modulate immune function in the gut.
- Regulation of appetite and metabolism
- SCFAs have anti-inflammatory effects
- Butyrate has been shown to have a protective effect against colon cancer by inducing cell differentiation and apoptosis.

Commensal Microbes

- extract nutrients and energy from our diets,
- maintain gut barrier function,
- produce vitamins (biotin and vitamin K),
- and protect against colonization by potential pathogens

Bifidobacterium species

Lactobacillus species

Faecalibacterium prausnitzii

Akkermansia muciniphila

COMMENSAL BACTERIA	
Bacteroides fragilis	Gram-negative species of the <i>Bacteroidetes</i> phylum. Immune-modulating normal gut species. Believed to be involved in microbial balance, barrier integrity, and neuroimmune health (Hsiao 2013). High levels may result from reduced digestive capacity or constipation. Low levels may contribute to reduced anti-inflammatory activity in the intestine.
Bifidobacterium spp.	Gram-positive genus in the <i>Actinobacteria</i> phylum. Present in breast milk. Colonizes the human GI tract at birth. Common in probiotics. Thrives on a wide variety of prebiotic fibers. Low levels may result from low fiber intake or reduced mucosal health. High levels are more common in children than in adults.
Enterococcus spp.	Gram-positive genus of lactate-producing bacteria in the <i>Firmicutes</i> phylum. High levels may be due to reduced digestive capacity, constipation or small intestinal bacterial overgrowth. Low levels may indicate insufficiency of beneficial bacteria.
Escherichia spp.	Gram-negative genus in the <i>Proteobacteria</i> phylum. Normal gut flora. <i>Escherichia coli</i> (<i>E. coli</i>) is the primary species in this genus. Most <i>E. coli</i> are nonpathogenic (<i>pathogenic E. coli strains are measured separately in "Pathogens" section of the GI-MAP</i>). High levels may be indicative of increased intestinal inflammatory activity. Low levels may indicate reduced mucosal health and decreased protection against pathogenic <i>E. coli</i> .
Lactobacillus spp.	Gram-positive genus of lactate-producing bacteria in the <i>Firmicutes</i> phylum. Many strains used as probiotics. High levels may result from reduced digestive capacity or excessive intake of carbohydrates. Low levels may be due to low carbohydrate intake or high salt intake, and may also indicate reduced mucosal health.
Enterobacter spp.	Gram-negative genus in the <i>Proteobacteria</i> phylum. Closely related to <i>E. coli (in the same taxonomic family)</i> . High levels may indicate increased intestinal inflammatory activity. Low levels may indicate reduced mucosal health.
Akkermansia municiphila	Keystone species and primary mucus degrader. Generates mucus-derived sugars and metabolic products that support the growth and energy needs of other gut microbes. Promotes mucosal health and mucus production. Low levels associated with obesity and metabolic dysfunction. High levels linked to multiple sclerosis.
Faecalibacterium prausnitzii	Widely recognized as an important keystone species in the Clostridia class, as well as a major butyrate producer. Promotes anti-inflammatory processes and mucosal homeostasis. Reduced levels have been associated with a wide range of chronic inflammatory and autoimmune diseases.
Roseburia spp.	A genus of Gram-positive anaerobic bacteria in the <i>Clostridia</i> class that inhabit the human colon. The <i>Roseburia</i> genus has five well-characterized species, all of which produce short-chain fatty acids (SCFAs), such as acetate, propionate, and butyrate. <i>Roseburia</i> can also produce butyrate from acetate promoting balance in energy homeostasis. The genus is widely recognized to influence colonic motility, support immunity, and suppress inflammation. Low levels are associated with several disease (including irritable bowel syndrome, obesity, Type 2 diabetes, nervous system conditions and allergies).

Pathogenic Microbes

- Salmonella species
- Escherichia coli (some strains)
- Clostridium difficile
- Campylobacter jejuni
- Helicobacter pylori

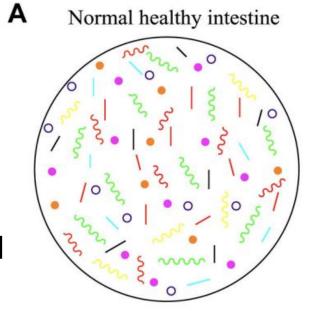
DYSBIOTIC & OVERGROWTH BACTERIA		
Bacillus spp.	Common group of gram-positive bacteria in the <i>Firmicutes</i> phylum. Some strains are used as probiotics. High levels may result from reduced digestive function, SIBO, or constipation.	
Enterococcus faecalis Enterococcus faecium	Gram-positive species in the <i>Firmicutes</i> phylum. High levels may result from reduced stomach acid, PPI use, compromised digestive function, SIBO or constipation. High natural resistance to some antibiotics, which may result in overgrowth.	
Morganella spp.	Gram-negative group in the <i>Proteobacteria</i> phylum. May produce histamine. High levels may indicate increased intestinal inflammatory activity. High levels may cause diarrhea, and may also be associated with SIBO.	
Pseudomonas spp. Pseudomonas aeroginosa	Gram-negative bacteria in the <i>Proteobacteria</i> phylum. High levels may indicate increased intestinal inflammatory activity and may cause abdominal cramping and loose stools. Some strains of <i>P. aeroginosa</i> may produce toxins that can damage cells.	
Staphylococcus spp. Staphylococcus aureus	Gram-positive bacteria in the <i>Firmicutes</i> phylum. High levels may result from reduced digestive capacity, and intestinal inflammatory activity. Some strains may produce toxins and contribute to loose stools or diarrhea.	
Streptococcus spp.	Gram-positive bacteria in the <i>Firmicutes</i> phylum. <i>Streptococcus</i> spp. colonize skin and mucous membranes throughout the body; High levels in the intestine may result from low stomach acid, PPI use, reduced digestive capacity, SIBO or constipation; Elevated levels may also be indicative of intestinal inflammatory activity, and may cause loose stools.	

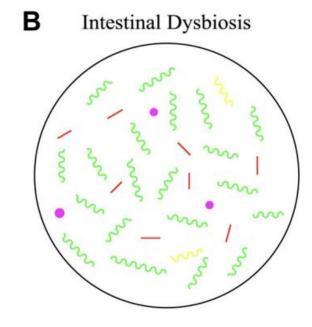
INFLAMMATORY & AUTOIMMUNE-RELATED BACTERIA		
Citrobacter spp. Citrobacter freundii.	Gram-negative bacteria in the <i>Proteobacteria</i> phylum. High levels may indicate increased intestinal inflammatory activity.	
Klebsiella spp. Klebsiella pneumoniae	Gram-negative bacteria in the <i>Proteobacteria</i> phylum. Common residents of the oral cavity and respiratory tract. May cause diarrhea, gas, abdominal pain, and bloating; Common after long-term antibiotic use; May release histamine in the gut; High levels may indicate increased intestinal inflammatory activity.	
Mycobacterium avium subsp. paratuberculosis	Bacterial species in the <i>Actinobacteria</i> phylum. Higher levels have been associated with Crohn's disease and rheumatoid arthritis.	
Proteus spp. Proteus mirabilis	Gram-negative bacteria in the <i>Proteobacteria</i> phylum. High levels may indicate increased intestinal inflammatory activity; May contribute to loose stools or diarrhea Pets or wild animals can be a source.	

Gut Dysbiosis

•Disruption to the microbiome resulting in an imbalance in the microbiota.

Overgrowth
 of opportunistic colonies to fill the void





Changes in the functional composition and metabolic activities

Linked to metabolic, cardiovascular, autoimmune, and inflammatory diseases.

Causes of Dysbiosis

- Standard American Diet (SAD)
- C-Section
- Formula feeding
- Xenobiotics
 – drugs,
 food additives, pesticides,
 and herbs/supplements
- Mold exposure
- Infections
- Alcohol abuse
- Lack of exercise
- Stress



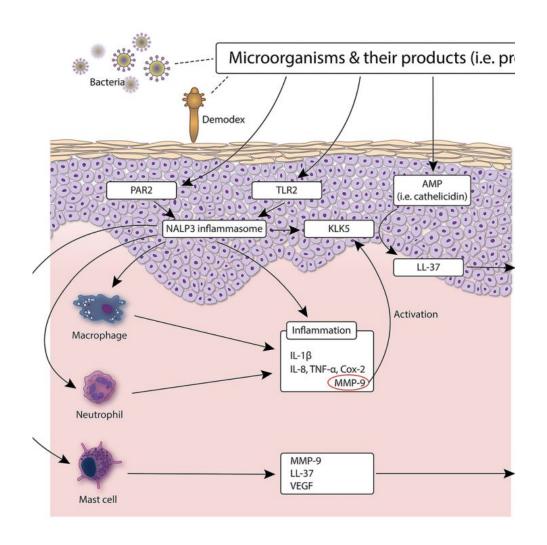
Diseases Associated with GI Dysbiosis

GI Conditions

- Alcoholic liver disease
- Antibiotic-associated diarrhea
- Chemotherapy-associated diarrhea
- Clostridium difficile-associated disease
- Celiac disease
- Crohn's disease
- Diverticular disease
- Irritable bowel syndrome
- Liver cirrhosis
- Non-alcoholic fatty liver disease
- Radiotherapy-associated diarrhea
- SIBO
- Ulcerative colitis

Non-GI Condition

- Alzheimer's disease
- Anxiety
- Asthma
- Atopic eczema
- Autism
- Chronic Fatigue Syndrome
- Depression
- Kidney stones
- Metabolic syndrome
- Multiple sclerosis
- Obesity
- Parkinson's disease
- Rheumatoid arthritis
- Type I diabetes
- Type II diabetes



Rosacea – Cutaneous Microbiome Imbalances

Imbalances in cutaneous organisms: Cutibacterium acnes, Staphylococcus epidermidis, Bacillus oleronius, and Demodex folliculorum

Potential cutaneous effects of alterations in the gastrointestinal (GI) microbiome.

Associations with GI pathologies

Rosacea and the Gut

Inflammatory Bowel Disease (IBD)

Celiac Disease

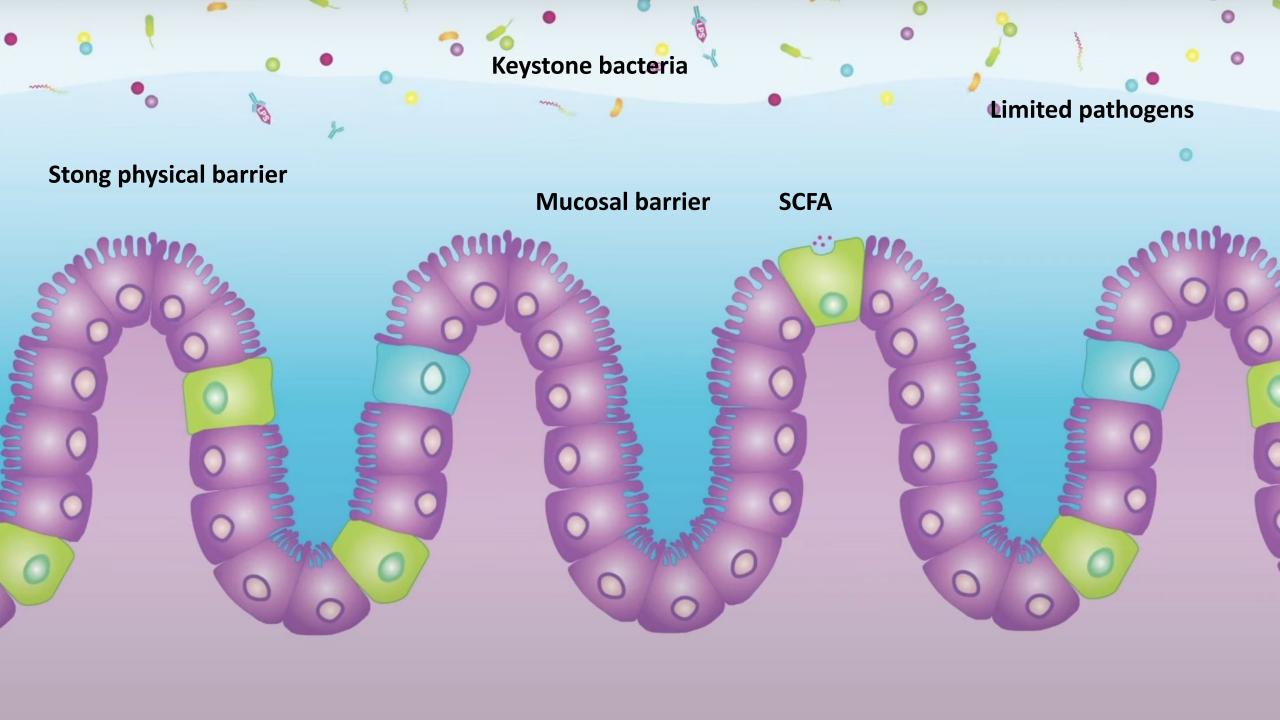
Irritable Bowel Syndrome (IBS)

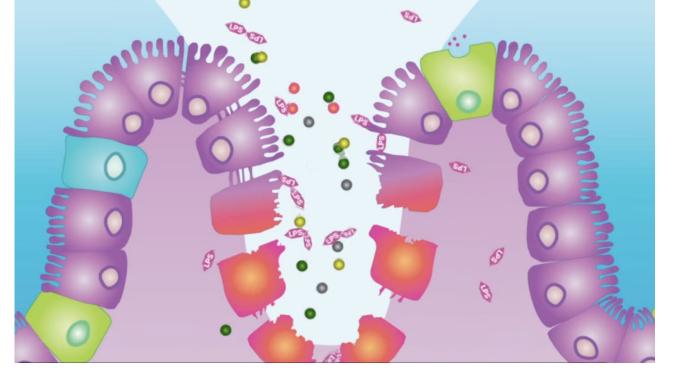
Gastroesophageal Reflux Disease (GERD)

Helicobacter Pylori (HP) infection

Small Intestine Bacterial Overgrowth (SIBO)

• Seen in 46-66% of rosacea patients



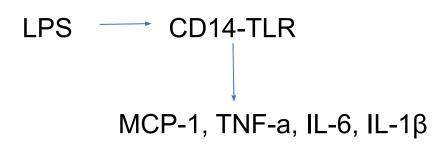


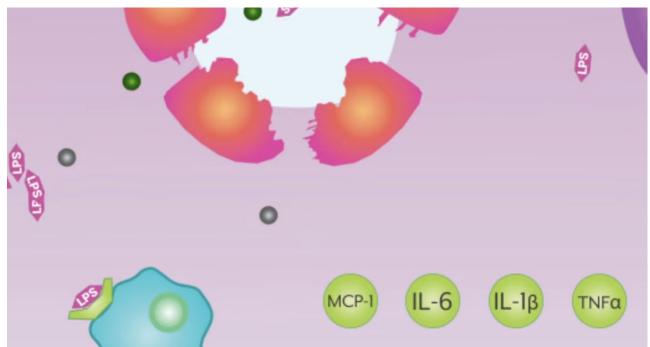
Unbalanced Microbial Population

Harmful bacteria feed on glycans or polysaccharides, and degrade the mucosal barrier and loosen the tight junctions

Leaky Gut = migration of toxins from the intestines into circulation

lipopolysaccharide (LPS)





Systemic Inflammation

- Brain
- Fat stores
- Nerves
- Joints
- Eyes

"Leaky" barrier Foreign molecules Inflammatory foods **Immune Attack**

The 5Rs Functional Medicine Approach to Healing the Gut





Remove

Remove anything irritating the gut



Replace

Replace any missing digestive elements



Reinoculate

Reinoculate the gut microbiome with good bacteria



Repair

Repair intestinal cells



Rebalance

Rebalance lifestyle elements



Remove allergic foods, parasites, bacteria or yeast (berberine, caprylic acid, garlic oil, oil of oregano, uva ursi, olive leaf extract)



Replace digestive secretions: Digestive support: Betaine hydrochloride, apple cider vinegar, herbal bitters, ox bile, lactase, pancreatic enzymes (amylase, lipase, protease), pepsin.



Reinoculate with good bacterial: Supplementation with probiotics, along with the use of prebiotics helps re-establish the proper microbial balance.



Repair the lining of the GI tract: Restore the integrity of the gut mucosa by giving support to healthy mucosal cells, as well as immune support.



Rebalance – lifestyle choices: Sleep, diet, exercise, and stress management

- Probiotics Live
 microorganisms with beneficial health effects
 on the host. Help digest food, support immune
 cells and produce vitamins.
- Prebiotics What your bugs eat
- <u>Postbiotics</u> are the metabolites (acetate, propionate, and butyrate) of the beneficial bacteria, and these bioactive compounds help fight infections, reduce inflammation, and have anti-cancer effects.





Probiotic Nomenclature

Lactobacillus acidophilus M23 Genus Species Strain

The strain of the bacteria determines the quality and clinical efficacy.

Match the action of specific probiotic strains to the condition or an imbalance being treated.

Strains	Origin
Lactobacillus sp. JNU 8829	Infant feces
Lactobacillus acidophilus KU41	Infant feces
Lactobacillus acidophilus M23	Infant feces
Lactobacillus brevis CH7	Kimchi
Lactobacillus casei MB3	Kimchi
Lactobacillus fermentum NS2	Kimchi
Lactobacillus plantarum M13	Kimchi
Lactobacillus plantarum NS3	Kimchi
Lactobacillus sakei CH8	Fermented olive
Lactobacillus sakei MA9	Fermented olive

Probiotics Prescribing

Upregulation of non-specific Immunity

 Lactobacillus acidophilus NCFM + Bifidobacterium lactis Bi07

Upregulating GIT production of slgA:

 Saccharomyces cerevisiae var boulardii CNCM I-745, Lactobacillus rhamnosus GG, Lactobacillus acidophilus La5

Down regulation of pathogens

• L.reuteri (DMS 17938) has been shown to eradicate H.pylori

Dry Eye Disease

- E. faecium LMG S-28935 and Saccharomyces boulardii MUCL 53837
- S. boulardii MUCL 53837 & E. faecium LMG S-28935
- B. lactis and B. bifido
- E. faecium WB2000
- MULTIBIOTIC[™] probiotic contains 21.075 billion CFU of bacteria per capsule, including Streptococcus, Lactobacillus, and Bifidobacterium species



APPLE CIDER VINEGAR

contain healthy acids that encourage a pH in your body that supports the growth of probiotics.



FERMENTED DAIRY

is probiotic-rich and helps improve gut health.



SAUERKRAUT

is rich in Lactobacillus. It's high in vitamin C and in digestive enzymes.



KIMCHI

the Korean cousin to sauerkraut, made with Chinese cabbag and some other food and spices.



NATTO

is a Japanese dish of fermented soybeans high in *Bacillus* subtilis.



KVASS

is a common beverage in Eastern Europe made from fermented barley or rye.



MISO

is a major component of Japanese medicine.



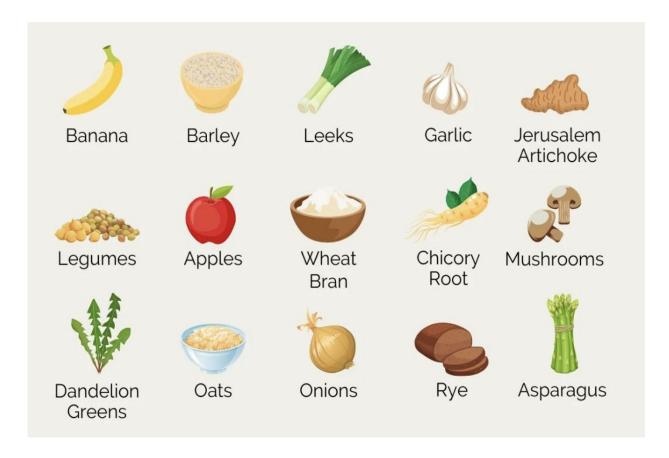
KOMBUCHA

is black tea fermented using a symbiotic colony of bacteria and yeast.

Probiotic Supplementation

Use a broad-spectrum, diverse probiotic formula, 50–450 billion CFUs/day containing Lactobacillus acidophilus, Bifidobacterium bifidum, Bifidobacterium longum, Lactobacillus rhamnosus, Bifidobacterium breve, Lactobacillus casei, Streptococcus thermophilus

- Lactobacillus rhamnosus GG
- Saccharomyces boulardii (probiotic yeast)
- Akkermansia muciniphila



It's Not Just What We Eat

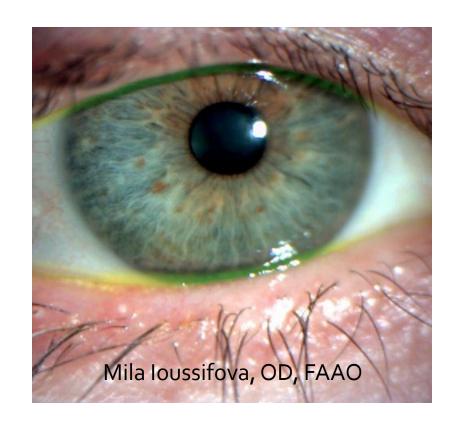
- Sunlight exposure
- Exercise
- Stress
- Sleep

Your bugs are listening to your thoughts!



Case Report

- 30 yr old Caucasian male
- Ocular history- Dry eye, ocular rosacea, blepharitis.
 On maintenance IPL treatments
- Medical history
 - GERD (dx 8/2013), Gallstones, Gallbladder removed in 2015
 - Migraines, Allergies, Rosacea
- Medications
 - Omeprazole daily since 2015
 - Antibiotics frequently for strep



- GI symptoms stomach pain, bloating, and nausea
- Food aversion acidic foods, red meat, eggs, and fish
- No food allergies

GI- Map

pylori	_		
Extension 1	Result		Normal
elicobacter pylori	5.4e2		<1.0e3
rulence Factor, babA	Negative		Negative
rulence Factor, cagA	Negative		Negative
rulence Factor, dupA	Negative		Negative
rulence Factor, iceA	Negative		Negative
rulence Factor, olpA	Negative		Negative
rulence Factor, vacA	Negative		Negative
rulence Factor, virB	Negative		Negative
rulence Factor, virD	Negative		Negative
rmai Bacterial Flora			
	Result		Normal
acteroides fragilis	7.30e9		1.6e9 - 2.5e11
fidobacterium spp.	1.28e10		>6.7e7
nterococcus spp.	2.47e7		1.9e5 - 2.0e8
scherichia spp.	2.07e8		3.7e6 - 3.8e9
actobacillus spp.	3.98e7		8.6e5 - 6.2e8
Clostridia (class)	3.4107		5.0e6 - 5.0e7
nterobacter spp.	7.00e7	High	1.0e6 - 5.0e7
kkermansia muciniphila	<dl< td=""><td></td><td>1.0e1 - 5.0e4</td></dl<>		1.0e1 - 5.0e4
aecalibacterium prausnitzii	6.14e5		1.0e3 - 5.0e8
nyla Microbiota	Result		Normal
acteroidetes	3.31e13	High	8.6e11 - 3.3e12
rmicutes	1.56e12	High	5.7e10 - 3.0e11
rmicutes:Bacteroidetes Ratio	0.05		<1.0

Enterobacter spp. - High

Gram-negative genus in the Proteobacteria phylum. Closely related to E. coli (in the same taxonomic family). High levels may indicate increased intestinal inflammatory activity. Low levels may indicate reduced mucosal health.

Firmicutes/Bacteroidetes - High Poor diet, Dysbiosis, Maldigestion or hypochlorhydria

Additional Dysbiotic/Overgrowth Bacteria Result Normal Bacillus spp. 2,72e6 High <1,50e5 Enterococcus faecalis <1.00e4 <1,00e4 Enterococcus faecium <1 <1,00e4 Morganella spp. <1 <1,00e3 Pseudomonas spp. <1 <1,00e4 Pseudomonas aeruginosa <1 <1,00e4 Staphylococcus spp. <1,00e4 <1,00e4 Staphylococcus aureus 2,39e2 <5,00e2 Streptococcus spp. 9,57e4 High <1,00e3 Methanobacteriaceae (family) 2,84e8 <5,00e2 Streptococcus spp. 1,70e4 <5,00e3 Citrobacter spp. 1,70e4 <5,00e3 Citrobacter spp. <1,70e4 <5,00e3 Klebsiella spp. <1 <5,00e3 Klebsiella pneumoniae <1 <5,00e3 Klebsiella pneumoniae <1 <5,00e3 Proteus mirabilis <1 <1,00e3 Fusobacterium spp. <1,00e3 <					
Enterococcus faecalis Enterococcus faecium Adi Enterococcus faecium Adi Morganella spp. Pseudomonas spp. Adi Pseudomonas spp. Adi Staphylococcus spp. Staphylococcus spp. Staphylococcus spp. Streptococcus spp. Methanobacteriaceae (family) Potential Autoimmune Triggers Citrobacter spp. Citrobacter freundii Klebsiella spp. Klebsiella pneumoniae M. avium subsp. paratuberculosis Proteus mirabilis Fusobacterium spp. Result Candida spp. Candida sp	Additional Dysbiotic/Overgrowth Bacteria	Result		Normal	
Enterococcus faecains Enterococcus faecains Enterococcus faecains Anorganella spp. Pseudomonas spp. Yesudomonas spp. Pseudomonas aeruginosa Staphylococcus spp. Staphylococcus aureus Staphylococcus sureus Streptococcus spp. Methanobacteriaceae (family) Potential Autoimmune Triggers Citrobacter spp. Citrobacter freundii Klebsiella spp. Klebsiella pneumoniae M. avium subsp. paratuberculosis Proteus mirabilis Proteus mirabilis Furogli/Yeast Candida spp. Ca	Bacillus spp.	2.72e6	High	<1.50e5	
Morganella spp.	Enterococcus faecalis	<dl< td=""><td></td><td><1.00e4</td><td></td></dl<>		<1.00e4	
Pseudomonas spp. Image: Pseudomonas aeruginosa Image: Pseudomonas Image: Pseudomon	Enterococcus faecium	<dl< td=""><td></td><td><1.00e4</td><td></td></dl<>		<1.00e4	
Pseudomonas aeruginosa <di <di="" =""><5.00e2 <di <di="" =""><1.00e4 <di <<="" <di="" td="" =""><td>Morganella spp.</td><td><dl< td=""><td></td><td><1.00e3</td><td></td></dl<></td></di></di></di></di></di></di></di></di>	Morganella spp.	<dl< td=""><td></td><td><1.00e3</td><td></td></dl<>		<1.00e3	
Staphylococcus spp. <di< td=""><td>Pseudomonas spp.</td><td><dl< td=""><td></td><td><1.00e4</td><td></td></dl<></td></di<>	Pseudomonas spp.	<dl< td=""><td></td><td><1.00e4</td><td></td></dl<>		<1.00e4	
Staphylococcus area 2.39e2 5.00e2	Pseudomonas aeruginosa	<dl< td=""><td></td><td><5.00e2</td><td></td></dl<>		<5.00e2	
Streptococcus surieus 9.57e4 High <1.00e3 Methanobacteriaceae (family) 2.84e8 <5.00e9	Staphylococcus spp.	<dl< td=""><td></td><td><1.00e4</td><td></td></dl<>		<1.00e4	
Methanobacteriaceae (family) 2.84e8 <5.00e9 Potential Autoimmune Triggers Result Normal Citrobacter spp. 1.70e4 <5.00e6	Staphylococcus aureus	2.39e2		<5.00e2	
Normal Normal	Streptococcus spp.	9.57e4	High	<1.00e3	
Citrobacter spp. 1.70e4 <5.00e6	Methanobacteriaceae (family)	2.84e8		<5.00e9	
Citrobacter freundii <dl< td=""> <5.00e5</dl<>	Potential Autoimmune Triggers	Result		Normal	
Klebsiella spp. <dl< td=""> <5.00e3</dl<>	Citrobacter spp.	1.70e4		<5.00e6	
Klebsiella pneumoniae	Citrobacter freundii	<dl< td=""><td></td><td><5.00e5</td><td></td></dl<>		<5.00e5	
M. avium subsp. paratuberculosis <dl< td=""> <5.00e3</dl<>	Klebsiella spp.	<dl< td=""><td></td><td><5.00e3</td><td></td></dl<>		<5.00e3	
Proteus spp. <dl< td=""> <5.00e4</dl<>	Klebsiella pneumoniae	<dl< td=""><td></td><td><5.00e4</td><td></td></dl<>		<5.00e4	
Proteus mirabilis <dl< td=""> <1.00e3</dl<>	M. avium subsp. paratuberculosis	<dl< td=""><td></td><td><5.00e3</td><td></td></dl<>		<5.00e3	
### Froteus mirabilis #### Fusobacterium spp. #### Prevotella spp. #### Prevotella spp. #### Result #### Candida spp. #### Candida albicans #### Candida albicans ##### Candida albicans ##### Candida albicans ##### Candida albicans ###### Candida albicans ###################################	Proteus spp.	<dl< td=""><td></td><td><5.00e4</td><td></td></dl<>		<5.00e4	
Prevotella spp. 2.26e8 High <1.00e8 Fungi/Yeast Result Normal Candida spp. <dl< td=""> <5.00e3</dl<>	Proteus mirabilis	<dl< td=""><td></td><td><1.00e3</td><td></td></dl<>		<1.00e3	
Result Normal	Fusobacterium spp.	8.82e7		<1.00e8	
Result Normal	Prevotella spp.	2.26e8	High	<1.00e8	
Candida spp. <dl< td=""> <5.00e3</dl<>	Fungi/Yeast				
Candida albicans <dl< td=""> <5.00e2</dl<>		Result		Normal	
Cartolida albicaris	Candida spp.	<dl< td=""><td></td><td><5.00e3</td><td></td></dl<>		<5.00e3	
Microsporidium spp. <dl< td=""> <5.00e3</dl<>	Candida albicans	<dl< td=""><td></td><td><5.00e2</td><td></td></dl<>		<5.00e2	
Rhodotorula spp. Cytomegalovirus Cytomegalovirus Color Result Normal Cytomegalovirus Color Result Color Colo	Geotrichum spp.	<dl< td=""><td></td><td><3.00e2</td><td></td></dl<>		<3.00e2	
Viruses Result Normal Cytomegalovirus All Control Control Cytomegalovirus Result Normal Cytomegalovirus Result Normal Cytomegalovirus	Microsporidium spp.	<dl< td=""><td></td><td><5.00e3</td><td></td></dl<>		<5.00e3	
Result Normal Cytomegalovirus <dl <1.00e5<="" td=""><td>Rhodotorula spp.</td><td><dl< td=""><td></td><td><1.00e3</td><td></td></dl<></td></dl>	Rhodotorula spp.	<dl< td=""><td></td><td><1.00e3</td><td></td></dl<>		<1.00e3	
Cytomegalovirus <dl <1.00e5<="" td=""><td>Viruses</td><td></td><td></td><td></td><td></td></dl>	Viruses				
Cytomegalovirus		Result		Normal	
-dl	Cytomegalovirus	<dl< td=""><td></td><td><1.00e5</td><td></td></dl<>		<1.00e5	
Epstein-Barr Virus	Epstein-Barr Virus	<dl< td=""><td></td><td><1.00e7</td><td></td></dl<>		<1.00e7	

Bacillus spp. - High

Common group of gram-positive bacteria in the Firmicutes phylum. High levels may result from reduced digestive function, SIBO, or constipation.

Streptococcus spp. - High

Gram-positive bacteria in the Firmicutes phylum. Streptococcus spp. colonize skin and mucous membranes throughout the body; High levels in the intestine may result from low stomach acid, PPI use, reduced digestive capacity, SIBO or constipation; Elevated levels may also be indicative of intestinal inflammatory activity, and may cause loose stools.

Prevotella spp.- High

Gram-negative species in the Bacteroidetes phylum. Associated with rheumatoid arthritis. High levels may result from reduced digestive capacity, or a high-starch diet.

Parasites		
Protozoa	Result	Normal
Blastocystis hominis	<dl< td=""><td><2.00e3</td></dl<>	<2.00e3
Chilomastix mesnili	<dl< td=""><td><1.00e5</td></dl<>	<1.00e5
Cyclospora spp.	<dl< td=""><td><5.00e4</td></dl<>	<5.00e4
Dientamoeba fragilis	<dl< td=""><td><1.00e5</td></dl<>	<1.00e5
Endolimax nana	<dl< td=""><td><1.00e4</td></dl<>	<1.00e4
Entamoeba coli	<dl< td=""><td><5.00e6</td></dl<>	<5.00e6
Pentatrichomonas hominis	<dl< td=""><td><1.00e2</td></dl<>	<1.00e2
Worms	Result	Normal
Ancylostoma duodenale	Not Detected	Not Detected
Ascaris lumbricoides	Not Detected	Not Detected
Necator americanus	Not Detected	Not Detected
Trichuris trichiura	Not Detected	Not Detected
Taenia spp.	Not Detected	Not Detected
Intestinal Health		
Digestion	Result	Normal
Steatocrit	<dl< td=""><td><15 %</td></dl<>	<15 %
Elastase-1	724	>200 ug/g
GI Markers	Result	Normal
β-Glucuronidase	1334	<2486 U/mL
Occult Blood - FIT	0	<10 ug/g
mmune Response	Result	Normal
Secretory IgA	255 Lo	w 510 - 2010 ug/g
Anti-gliadin IgA	20	<175 U/L
Inflammation	Result	Normal
Calprotectin	66	<173 ug/g
Add-on Test	Result	Normal
Zonulin	80.8	<175 ng/g

Low Fecal SIgA –

The gut immune system is suppressed. Possible underlying causes: chronic dysbiosis, antigen exposure, chronic stress, immunocompromised patient, or even protein malnutrition.

Dietary and supplement recommendations

Remove

- refined carbohydrates, sugar, processed foods and saturated fats
- Avoid eating out (meal plan for lunch and dinner)

Replace

 Bio-Gest Digestive Enzymes Thorne (Betaine Hydrochloride, L-Glutamic Acid Hydrochloride, Ox Bile, Pancreatin, Pepsin)

Reinoculate

Ortho Biotic, Ortho Molecular Products

Proprietary Blend ... 50 Billion CFU
 Lactobacillus acidophilus (La-14)
 Lactobacillus paracasei (Lpc-37)
 Bifidobacterium lactis (Bl-04)
 Bifidobacterium bifidum (Bb-06)
 Lactobacillus plantarum (Lp-115)
 Lactobacillus rhamnosus (GG)
 Saccharomyces boulardii ... 3 Billion CFU



Repair

- GI-Revive L-Glutamine: N-Acetyl Glucosamine: Citrus Pectin: Deglycyrrhizinated Licorice: Aloe Vera: Slippery Elm: Mucin: Marshmallow: Chamomile: Okra: Cat's Claw: Methylsulfonylmethane: Quercetin: Prunus: Zinc-Carnosine
- Fish oil, Mg

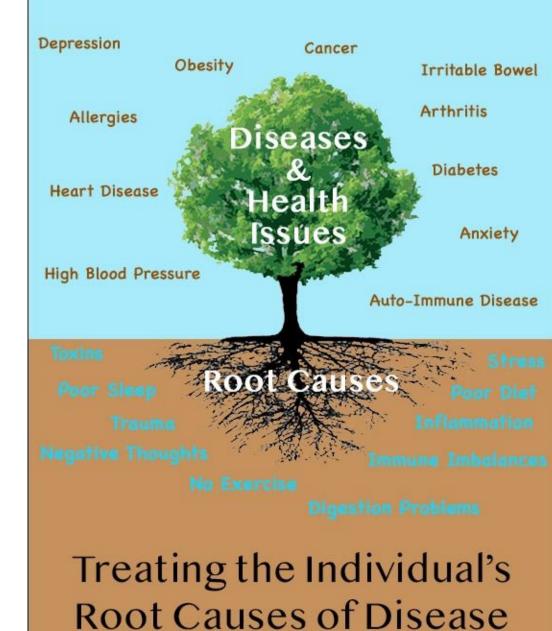
Rebalance

- Diet: Increase phytonutrients 9 servings of fruits and vegetables a day, eliminate alcohol and increase water intake (double)
- Physical activity: 3-4 times a week and add weight training. Walk the dog after meals
- Stress management and sleep: 10- minutes of meditation or journaling a day. Prioritizing 7-8 hours of sleep a day

Follow up

- Patient reports
 - Improved GI symptoms
 - Less food intolerances
 - Rarely uses Prilosec
 - Reduced allergy symptoms
 - Improved skin
 - Less rosacea and dry eye flareups

Functional Medicine



Thank you

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