

Microbiome Information for: Inflammatory Bowel Disease

For non-prescribing Medical professionals Review

The suggestions below are based on an Expert System (Artificial Intelligence) modelled after the MYCIN Expert System produced at Stanford University School of Medicine in 1972. The system uses over 1,800,000 facts with backward chaining to sources of information. The typical sources are studies published on the US National Library of Medicine.

Many recent studies has found that symptoms and symptom severity has strong associations to the microbiome for many conditions. Correcting the microbiome dysfunction is beleived to reduce the severity of symptoms. In some cases, this correction may cause symptoms to disappear.

These are a *a priori suggestions* that are predicted to independently reduce microbiome dysfunction. Suggestions should *only be done after a review* by a medical professional factoring in patient's conditions, allergies and other issues.

This report may be freely shared by a patient to their medical professionals

Best practise for making microbiome adjustments is to obtain the individuals microbiome. The following are the best microbiome to use with this expert system model. The suggestions below are intended as temporary suggestions until a test result in received.

In the USA

Ombre (<https://www.ombrelab.com/>)

Thome (<https://www.thome.com/products/dp/gut-health-test>)

Worldwide: BiomeSight (<https://biomesight.com>) - Discount Code 'MICRO'

Analysis Provided by Microbiome Prescription

A Microbiome Analysis Company

892 Lake Samish Rd, Bellingham WA 98229

Email: Research@MicrobiomePrescription.com

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Bacteria being reported because of atypical values.

These bacteria were reported atypical in studies of Inflammatory Bowel Disease

Nota Bena: Many studies are done with a small sample size or mixtures of condition subsets which can greatly diminish the ability to detect bacteria shifts.

Bacteria Name	Rank	Shift	Taxonomy ID	Bacteria Name	Rank	Shift	Taxonomy ID
Betaproteobacteria	class	High	28216	Salmonella	genus	High	590
Chloroflexia	class	High	32061	Sellimonas	genus	High	1769710
Christensenellaceae	family	Low	990719	Shigella	genus	High	620
Actinomyces	genus	High	1654	Staphylococcus	genus	High	1279
Agathobacter	genus	Low	1766253	Streptococcus	genus	High	1301
Akkermansia	genus	Low	239934	Sutterella	genus	High	40544
Alistipes	genus	Low	239759	Cytophagales	order	High	768507
Anaerostipes	genus	Low	207244	Eubacteriales	order	Low	186802
Bacteroides	genus	High	816	[Clostridium] colinum	species	Low	36835
Barnesiella	genus	Low	397864	[Clostridium] innocuum	species	High	1522
Blautia	genus	Low	572511	[Clostridium] leptum	species	Low	1535
Campylobacter	genus	High	194	[Eubacterium] brachy	species	High	35517
Clostridium	genus	Low	1485	[Eubacterium] siraeum	species	Low	39492
Collinsella	genus	High	102106	[Ruminococcus] gnavus	species	High	33038
Coprococcus	genus	Low	33042	Actinomyces graevenitzii	species	High	55565
Eggerthella	genus	High	84111	Adlercreutzia equolifaciens	species	Low	446660
Enterobacter	genus	Low	547	Agathobacter rectalis	species	Low	39491
Enterococcus	genus	High	1350	Akkermansia muciniphila	species	Low	239935
Erysipelatoclostridium	genus	High	1505663	Alistipes putredinis	species	Low	28117
Escherichia	genus	High	561	Anaerobutyricum hallii	species	Low	39488
Eubacterium	genus	Low	1730	Bifidobacterium longum	species	Low	216816
Faecalibacterium	genus	Low	216851	Butyricicoccus pullicaecorum	species	Low	501571
Faecalitalea	genus	High	1573534	Clostridium butyricum	species	Low	1492
Fenollaria	genus	Low	1686313	Collinsella aerofaciens	species	Low	74426
Flavobacterium	genus	Low	237	Enterococcus faecalis	species	High	1351
Flavonifractor	genus	High	946234	Escherichia coli	species	High	562
Fusicatenibacter	genus	Low	1407607	Eubacterium ruminantium	species	Low	42322
Gemella	genus	High	1378	Eubacterium ventriosum	species	Low	39496
Haemophilus	genus	High	724	Eubacterium xylanophilum	species	Low	39497
Intestinibacter	genus	High	1505657	Faecalibacterium prausnitzii	species	Low	853
Intestinimonas	genus	Low	1392389	Gemmiger fornicilis	species	Low	745368
Klebsiella	genus	High	570	Lactocaseibacillus rhamnosus	species	Low	47715
Lachnospira	genus	Low	28050	Methanobrevibacter smithii	species	Low	2173
Lactobacillus	genus	High	1578	Methanosphaera stadtmanae	species	High	2317
Lactococcus	genus	High	1357	Parabacteroides distasonis	species	Low	823
Oscillospira	genus	Low	119852	Roseburia intestinalis	species	Low	166486
Phascolarctobacterium	genus	Low	33024	Ruminococcus bromii	species	Low	40518
Prevotella	genus	High	838	Segatella copri	species	Low	165179
Proteus	genus	High	583	Streptococcus salivarius	species	High	1304
Pseudomonas	genus	High	286	Thomasclavelia ramosa	species	High	1547
Roseburia	genus	Low	841	Veillonella atypica	species	High	39777
Ruminococcus	genus	Low	1263	Campylobacter concisus	strain	High	360104
				Adlercreutzia equolifaciens subsp. celatus	subspecies	Low	394340

Substance to Consider Adding or Taking

These are the most significant substances that are likely to improve the microbiome dysfunction. Dosages are based on the dosages used in clinical studies. For more information see: <https://microbiomeprescription.com/library/dosages>. These are provided as examples only

Colors indicates the type of substance: i.e. probiotics and prebiotics, herbs and spices, etc. There is no further meaning to them.

dairy

d-ribose 10 gram/day

fat

fluorine

green-lipped mussel

lactulose

lard

linseed(flaxseed) 30 mg/day

omega-3 fatty acids 4 gram/day

raffinose(sugar beet)

smoking

symbioflor 2 e.coli probiotics

Vitamin C (ascorbic acid) 30 g/day

Retail Probiotics

Over 260 retail probiotics were evaluated with the following deemed beneficial with no known adverse risks.

symbiopharm / symbioflo 2

Note: Some of these are only available regionally – search the web for sources.

Substance to Consider Reducing or Eliminating

These are the most significant substances have been identified as probably contributing to the microbiome dysfunction.

In some cases blood work may show low levels of some vitamins, etc. listed below. This may be due to *greedy* bacteria reported at a high level above. Viewing bacteria data on the Kyoto Encyclopedia of Genes and Genomes (<https://www.kegg.jp/>) may provide better insight on the course of action to take.

arabinogalactan (prebiotic)	oregano (organum vulgare, oil)
barley	resistant starch
cinnamon (oil, spice)	rosmarinus officinalis, rosemary
clostridium butyricum (probiotics), Miya, Miyarisan	Slippery Elm
Curcumin	soy
fasting	syzygium aromaticum (clove)
foeniculum vulgare, fennel	thyme (thymol, thyme oil)
garlic (allium sativum)	triphala
inulin (prebiotic)	vegetarians
lactobacillus plantarum (probiotics)	vitamin d
lactobacillus reuteri (probiotics)	walnuts
lactobacillus rhamnosus gg (probiotics)	wheat

Sample of Literature Used

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Asthma

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Autism

Autoimmune Disease

Barrett esophagus cancer

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Bipolar Disorder

Brain Trauma

Breast Cancer

Cancer (General)

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Epilepsy
erectile dysfunction
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gallstone disease (gsd)
Gastroesophageal reflux disease (Gerd) including Barrett's esophagus
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giant cell arteritis
Glioblastoma
Gout
Graves' disease
Halitosis
Hashimoto's thyroiditis
Heart Failure
Hemorrhoidal disease, Hemorrhoids, Piles
Hidradenitis Suppurativa
Histamine Issues
hypercholesterolemia (High Cholesterol)
hyperglycemia
Hyperlipidemia (High Blood Fats)
hypersomnia
hypertension (High Blood Pressure)
Hypothyroidism
Hypoxia
IgA nephropathy (IgAN)
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ME/CFS without IBS
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Mood Disorders
multiple chemical sensitivity [MCS]
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Multiple system atrophy (MSA)
myasthenia gravis
neuropathic pain
Neuropathy (all types)
neuropsychiatric disorders (PANDAS, PANS)
Nonalcoholic Fatty Liver Disease (nafld) Nonalcoholic
NonCeliac Gluten Sensitivity
Obesity
obsessive-compulsive disorder

Osteoarthritis
Osteoporosis
pancreatic cancer
Parkinson's Disease
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Postural orthostatic tachycardia syndrome
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