

## Microbiome Information for: Crohn's 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 have found that symptoms and symptom severity has strong associations to the microbiome for many conditions. Correcting the microbiome dysfunction is believed to reduce the severity of symptoms. In some cases, this correction may cause symptoms to disappear.

These are *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**

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Best practise for making microbiome adjustments is to obtain the individual's 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 is received.

In the USA

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

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

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

### Analysis Provided by Microbiome Prescription

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## Bacteria being reported because of atypical values.

These bacteria were reported atypical in studies of Crohn's Disease

**Nota Benia:** 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
Clostridia	class	Low	186801	Klebsiella	genus	Low	570
Halobacteria	class	High	183963	Lachnospiraceae	genus	High	1506553
Thermoplasmata	class	High	183967	Lachnospira	genus	Low	28050
Thermoprotei	class	High	183924	Lactobacillus	genus	High	1578
Aerococcaceae	family	High	186827	Lactococcus	genus	Low	1357
Bifidobacteriaceae	family	Low	31953	Leuconostoc	genus	High	1243
Christensenellaceae	family	Low	990719	Malassezia	genus	Low	55193
Enterobacteriaceae	family	High	543	Marinibryantia	genus	Low	248744
Lachnospiraceae	family	Low	186803	Methanobrevibacter	genus	Low	2172
Ruminococcaceae	family	Low	541000	Methanospaera	genus	High	2316
Abiotrophia	genus	Low	46123	Mogibacterium	genus	High	86331
Acetobacter	genus	Low	434	Monoglobus	genus	Low	2039302
Acidaminococcus	genus	Low	904	Olsenella	genus	Low	133925
Actinobacillus	genus	High	713	Paenibacillus	genus	Low	44249
Actinomyces	genus	High	1654	Parimonas	genus	High	543311
Adlercreutzia	genus	High	447020	Peptostreptococcus	genus	High	1257
Alistipes	genus	High	239759	Phascolarctobacterium	genus	Low	33024
Anaerofustis	genus	Low	264995	Picrophilus	genus	High	46631
Anaerostipes	genus	Low	207244	Polynucleobacter	genus	High	44013
Anaerotruncus	genus	High	244127	Porphyromonas	genus	Low	836
Atopobium	genus	High	1380	Prevotella	genus	Low	838
Barnesiella	genus	High	397864	Proteus	genus	High	583
Bilophila	genus	High	35832	Proteus	genus	High	210425
Blautia	genus	Low	572511	Pseudomonas	genus	Low	286
Burkholderia	genus	High	32008	Romboutsia	genus	Low	1501226
Butyricicoccus	genus	Low	580596	Roseburia	genus	Low	841
Butyrivibrio	genus	Low	830	Ruminococcus	genus	Low	1263
Candida	genus	High	1535326	Shigella	genus	High	620
Candidatus Soleaferrea	genus	High	1470353	Slackia	genus	Low	84108
Catenibacterium	genus	Low	135858	Solobacterium	genus	High	123375
Ceduoivirus	genus	Low	186532	Succinatimonas	genus	High	674963
Cetobacterium	genus	High	180162	Sutterella	genus	Low	40544
Clostridium	genus	High	1485	Thermoanaerobacter	genus	High	1754
Colidextribacter	genus	Low	1980681	Treponema	genus	Low	157
Collinsella	genus	Low	102106	Turicibacter	genus	Low	191303
Coprococcus	genus	Low	33042	Tyzzerella	genus	High	1506577
Corynebacterium	genus	Low	1716	Veillonella	genus	High	29465
Dehalobacterium	genus	Low	51514	Vibrio	genus	High	662
Desulfovibrio	genus	Low	872	Eubacteriales	order	Low	186802
Dialister	genus	High	39948	Lactobacillales	order	High	186826
Dielma	genus	High	1472649	Verrucomicrobiales	order	High	48461

Bacteria Name	Rank Shift	Taxonomy ID	Bacteria Name	Rank Shift	Taxonomy ID
Dorea	genus Low	189330	[Clostridium] leptum	species Low	1535
Eggerthella	genus High	84111	[Ruminococcus] gnavus	species High	33038
Eisenbergiella	genus High	1432051	Agathobacter rectalis	species Low	39491
Enterobacter	genus High	547	Alistipes shahii	species Low	328814
Enterococcus	genus High	1350	Anaerobutyricum hallii	species Low	39488
Escherichia	genus High	561	Bacteroides uniformis	species High	820
Facklamia	genus Low	66831	Blautia coccoides	species Low	1532
Faecalibacterium	genus Low	216851	Blautia faecis	species High	871665
Fusicatenibacter	genus Low	1407607	Escherichia coli	species High	562
Fusobacterium	genus High	848	Faecalibacterium prausnitzii	species Low	853
Gemmiger	genus Low	204475	Faecilicatena fissicatena	species High	290055
Gordonibacter	genus High	644652	Francisella tularensis	species Low	263
Haemophilus	genus High	724	Fusobacterium nucleatum	species High	851
Halococcus	genus High	2249	Hoylesella oralis	species High	28134
Holdemanella	genus Low	1573535	Isoptericola variabilis	species High	139208
Hungatella	genus High	1649459	Lachnospira eligens	species Low	39485
Isoptericola	genus High	254250	Pseudodesulfovibrio aespoeensis	species Low	182210
Jonquetella	genus High	428711	Roseburia inulinivorans	species High	360807
			Roseburia sp.	species Low	2049040

## 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.

candida albicans (prescription)

carboxymethyl cellulose (prebiotic)

carob

d-ribose 10 gram/day

fluorine

GABA 6 gram/day

grape polyphenols

green-lipped mussel

lactulose

linseed(flaxseed) 30 mg/day

mannooligosaccharide (prebiotic) 8 gram/day

raffinose(sugar beet)

resveratrol (grape seed/polyphenols/red wine) 2 gram/day

Sauerkraut

sesame cake/meal

smoking

symbioflor 2 e.coli probiotics

## 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)	<i>lactobacillus plantarum</i> (probiotics)
<i>bacillus subtilis</i> (probiotics)	<i>lactobacillus reuteri</i> (probiotics)
<i>bifidobacterium longum</i> (probiotics)	<i>lactobacillus rhamnosus gg</i> (probiotics)
Cacao	<i>Limosilactobacillus fermentum</i> (probiotic)
cinnamon (oil, spice)	<i>oregano</i> ( <i>origanum vulgare</i> , oil)
<i>clostridium butyricum</i> (probiotics), Miya, Miyarisan	quebracho
Curcumin	<i>rosmarinus officinalis</i> , rosemary
<i>foeniculum vulgare</i> , fennel	soy
garlic ( <i>allium sativum</i> )	<i>syzygium aromaticum</i> (clove)
inulin (prebiotic)	thyme (thymol, thyme oil)
<i>lactobacillus casei</i> (probiotics)	triphala
<i>lactobacillus paracasei</i> (probiotics)	vitamin d
	wheat

## Sample of Literature Used

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