

Microbiome Information for: Autism

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 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

[Our Facebook Discussion Page](#)

Bacteria being reported because of atypical values.

These bacteria were reported atypical in studies of Autism

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
Alphaproteobacteria	class	High	28211	Leptotrichia	genus	High	32067
Bacilli	class	High	91061	Megamonas	genus	High	158846
Bacteroidia	class	High	200643	Megasphaera	genus	High	906
Clostridia	class	High	186801	Moryella	genus	Low	437755
Deinococci	class	Low	188787	Odoribacter	genus	High	283168
Fusobacteriia	class	Low	203490	Oscillospira	genus	High	119852
Holophagae	class	Low	533205	Paraprevotella	genus	Low	577309
Mollicutes	class	Low	31969	Parasutterella	genus	Low	577310
Negativicutes	class	Low	909932	Peptostreptococcus	genus	High	1257
Terriglobia	class	High	204432	Phascolarctobacterium	genus	High	33024
Acidaminococcaceae	family	Low	909930	Phyllobacterium	genus	High	28100
Alcaligenaceae	family	High	506	Prevotella	genus	Low	838
Bacteroidaceae	family	High	815	Pseudomonas	genus	Low	286
Bifidobacteriaceae	family	Low	31953	Ralstonia	genus	High	48736
Eggerthellaceae	family	Low	1643826	Romboutsia	genus	Low	1501226
Enterobacteriaceae	family	High	543	Rothia	genus	High	508215
Erysipelotrichaceae	family	High	128827	Ruminiclostridium	genus	High	1508657
Eubacteriaceae	family	High	186806	Ruminococcus	genus	High	1263
Lachnospiraceae	family	Low	186803	Saccharomyces	genus	High	4930
Lactobacillaceae	family	High	33958	Sarcina	genus	High	1266
Methylobacteriaceae	family	High	119045	Staphylococcus	genus	High	1279
Moraxellaceae	family	High	468	Streptococcus	genus	Low	1301
Oscillospiraceae	family	High	216572	Sutterella	genus	High	40544
Pasteurellaceae	family	High	712	Tannerella	genus	Low	195950
Rikenellaceae	family	High	171550	Turicibacter	genus	High	191303
Sutterellaceae	family	High	995019	Tyzzereella	genus	Low	1506577
Veillonellaceae	family	Low	31977	Veillonella	genus	Low	29465
Acinetobacter	genus	High	469	Weissella	genus	Low	46255
Actinobacillus	genus	High	713	Lachnospiraceae incertae sedis	no rank	High	2840493
Actinomyces	genus	High	1654	Bacteroidales	order	Low	171549
Aggregatibacter	genus	High	416916	Bifidobacteriales	order	Low	85004
Akkermansia	genus	Low	239934	Eubacteriales	order	High	186802
Alistipes	genus	Low	239759	Selenomonadales	order	Low	909929
Amphibacillus	genus	High	29331	Victivallales	order	High	278082
Anaerococcus	genus	High	165779	[Clostridium] leptum	species	Low	1535
Anaerostipes	genus	Low	207244	[Eubacterium] siraeum	species	Low	39492
Anaerotruncus	genus	Low	244127	[Ruminococcus] lactaris	species	Low	46228
Aspergillus	genus	High	5052	[Ruminococcus] torques	species	High	33039
Bacillus	genus	High	1386	Acinetobacter johnsonii	species	High	40214
Bacillus	genus	High	55087	Acinetobacter rhizosphaerae	species	High	243922
Bacteroides	genus	High	816				

Bacteria Name	Rank	Shift	Taxonomy ID	Bacteria Name	Rank	Shift	Taxonomy ID
Barnesiella	genus	High	397864	Adlercreutzia equolifaciens	species	High	446660
Bifidobacterium	genus	Low	1678	Agathobacter rectalis	species	Low	39491
Bilophila	genus	Low	35832	Akkermansia muciniphila	species	Low	239935
Blautia	genus	Low	572511	Alistipes putredinis	species	Low	28117
Borrelia	genus	High	138	Anaerostipes caccae	species	High	105841
Burkholderia	genus	High	32008	Bacteroides sp.	species	Low	29523
Butyricicoccus	genus	Low	580596	Bifidobacterium adolescentis	species	Low	1680
Butyricimonas	genus	High	574697	Bifidobacterium longum	species	Low	216816
Caloramator	genus	High	44258	Butyricicoccus pullicaecorum	species	Low	501571
Candida	genus	High	5475	Campylobacter jejuni	species	Low	197
Candida	genus	High	1535326	Candida albicans	species	High	5476
Catenibacterium	genus	High	135858	Chloracidobacterium thermophilum	species	Low	458033
Cetobacterium	genus	High	180162	Clostridioides difficile	species	High	1496
Chlamydia	genus	High	810	Clostridium botulinum	species	High	1491
Chloracidobacterium	genus	Low	458032	Clostridium perfringens	species	High	1502
Chryseobacterium	genus	High	59732	Coralimargarita akajimensis	species	Low	395922
Clostridium	genus	High	1485	Desulfovibrio intestinalis	species	High	58621
Collinsella	genus	High	102106	Desulfovibrio piger	species	High	901
Coprobacter	genus	High	1348911	Dialister invisus	species	Low	218538
Coprococcus	genus	Low	33042	Eggerthella lenta	species	High	84112
Corynebacterium	genus	High	1716	Enterobacter cloacae	species	Low	550
Desulfovibrio	genus	High	872	Enterocloster bolteae	species	High	208479
Devosia	genus	Low	46913	Enterocloster clostridioformis	species	High	1531
Dialister	genus	Low	39948	Enterococcus faecalis	species	Low	1351
Dorea	genus	High	189330	Enterococcus gallinarum	species	Low	1353
Eggerthella	genus	High	84111	Escherichia coli	species	Low	562
Eisenbergiella	genus	Low	1432051	Faecalibacterium prausnitzii	species	Low	853
Enhydrobacter	genus	High	212791	Haemophilus parainfluenzae	species	Low	729
Enterobacter	genus	High	547	Hathewayia histolytica	species	High	1498
Enterococcus	genus	Low	1350	Intestinibacter bartlettii	species	High	261299
Escherichia	genus	Low	561	Klebsiella pneumoniae	species	High	573
Eubacterium	genus	Low	1730	Lachnospira eligens	species	Low	39485
Ezakiella	genus	Low	1582879	Lactacaseibacillus rhamnosus	species	Low	47715
Faecalibacterium	genus	High	216851	Lactiplantibacillus plantarum	species	Low	1590
Faecalibaculum	genus	Low	1729679	Limosilactobacillus reuteri	species	Low	1598
Filifactor	genus	High	44259	Mediterraneibacter gnavus	species	High	33038
Flavonifractor	genus	Low	946234	Mycoplasmoides pneumoniae	species	High	2104
Fusobacterium	genus	High	848	Parabacteroides distasonis	species	Low	823
Haemophilus	genus	Low	724	Phocaeicola vulgatus	species	High	821
Holdemanella	genus	High	1573535	Prevotella sp.	species	Low	59823
Intestinibacter	genus	High	1505657	Proteus mirabilis	species	Low	584
Klebsiella	genus	High	570	Romboutsia ilealis	species	High	1115758
Lachnospira	genus	Low	28050	Ruminiclostridium cellulolyticum	species	High	1521
Lactobacillus	genus	High	1578	Spirochaeta thermophila	species	Low	154
				Streptococcus pyogenes	species	High	1314
				Yersinia enterocolitica	species	High	630

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.

Ethyl alcohol {Grain alcohol}
Ferrum {Iron Supplements} 400 mg/day

high red meat
vegetarians

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.

(2->1)-beta-D-fructofuranan {Inulin}
fruit

Lactobacillus plantarum {L. plantarum}
oligosaccharides {oligosaccharides}
synthetic disaccharide derivative of lactose {Lactulose}

Sample of Literature Used

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Abdominal Aortic Aneurysm

Acne

Addison's Disease (hypocortisolism)

ADHD

Age-Related Macular Degeneration and Glaucoma

Allergic Rhinitis (Hay Fever)

Allergies

Allergy to milk products

Alopecia (Hair Loss)

Alzheimer's disease

Amyotrophic lateral sclerosis (ALS) Motor Neuron
Ankylosing spondylitis
Anorexia Nervosa
Antiphospholipid syndrome (APS)
Asthma
Atherosclerosis
Atrial fibrillation
Autism
Autoimmune Disease
Barrett esophagus cancer
benign prostatic hyperplasia
Biofilm
Bipolar Disorder
Brain Trauma
Breast Cancer
Cancer (General)
Carcinoma
cdkl5 deficiency disorder
Celiac Disease
Cerebral Palsy
Chronic Fatigue Syndrome
Chronic Kidney Disease
Chronic Lyme
Chronic Obstructive Pulmonary Disease (COPD)
Chronic Urticaria (Hives)
Coagulation / Micro clot triggering bacteria
Cognitive Function
Colorectal Cancer
Constipation
Coronary artery disease
COVID-19
Crohn's Disease
Cushing's Syndrome (hypercortisolism)
cystic fibrosis
d-lactic acidosis (one form of brain fog)
deep vein thrombosis
Denture Wearers Oral Shifts
Depression
Dermatomyositis
Eczema
Endometriosis
Eosinophilic Esophagitis
Epilepsy
erectile dysfunction
Fibromyalgia
Food Allergy
Functional constipation / chronic idiopathic constipation
gallstone disease (gsd)
Gastroesophageal reflux disease (Gerd) including Barrett's esophagus
Generalized anxiety disorder
giant cell arteritis
Glioblastoma
Gout
Graves' disease
Gulf War Syndrome
Halitosis
Hashimoto's thyroiditis
Heart Failure

hemorrhagic stroke
Hemorrhoidal disease, Hemorrhoids, Piles
Hidradenitis Suppurativa
High Histamine/low DAO
hypercholesterolemia (High Cholesterol)
hyperglycemia
Hyperlipidemia (High Blood Fats)
hypersomnia
hypertension (High Blood Pressure)
Hypothyroidism
Hypoxia
IgA nephropathy (IgAN)
Inflammatory Bowel Disease
Insomnia
Intelligence
Intracranial aneurysms
Irritable Bowel Syndrome
ischemic stroke
Juvenile idiopathic arthritis
Liver Cirrhosis
Long COVID
Low bone mineral density
Lung Cancer
Lymphoma
Mast Cell Issues/ mastitis
ME/CFS with IBS
ME/CFS without IBS
membranous nephropathy
Menopause
Metabolic Syndrome
Mood Disorders
multiple chemical sensitivity [MCS]
Multiple Sclerosis
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
Peanut Allergy
Polycystic ovary syndrome
Postural orthostatic tachycardia syndrome
Premenstrual dysphoric disorder
primary biliary cholangitis
Primary sclerosing cholangitis
Psoriasis
rheumatoid arthritis (RA), Spondyloarthritis (SpA)
Rosacea
Schizophrenia
scoliosis
sensorineural hearing loss

Sjögren syndrome
Sleep Apnea
Slow gastric motility / Gastroparesis
Small Intestinal Bacterial Overgrowth (SIBO)
Stress / posttraumatic stress disorder
Systemic Lupus Erythematosus
Tic Disorder
Tourette syndrome
Type 1 Diabetes
Type 2 Diabetes
Ulcerative colitis
Unhealthy Ageing
Vitiligo