

## Microbiome Information for: Chronic Kidney 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 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 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 Chronic Kidney 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		
Enterobacteriaceae	family	High	543	Parabacteroides	genus	High	375288
Erysipelotrichaceae	family	High	128827	Prevotella	genus	High	838
Halomonadaceae	family	High	28256	Romboutsia	genus	Low	1501226
Lachnospiraceae	family	Low	186803	Roseburia	genus	Low	841
Lactobacillaceae	family	Low	33958	Ruminiclostridium	genus	High	1508657
Moraxellaceae	family	High	468	Scardovia	genus	High	196081
Porphyromonadaceae	family	High	171551	Shigella	genus	High	620
Prevotellaceae	family	Low	171552	Subdoligranulum	genus	Low	292632
Pseudomonadaceae	family	High	135621	Thiothrix	genus	High	1030
Akkermansia	genus	Low	239934	Tyzzerella	genus	Low	1506577
Bacteroides	genus	High	816	Veillonella	genus	High	29465
Bilophila	genus	Low	35832	Weissella	genus	High	46255
Brachybacterium	genus	High	43668	[Clostridium] innocuum	species	High	1522
Butyricoccus	genus	Low	580596	[Ruminococcus] lactaris	species	Low	46228
Carnobacterium	genus	Low	2747	[Ruminococcus] torques	species	High	33039
Catenibacterium	genus	High	135858	Agathobacter rectalis	species	Low	39491
Coprococcus	genus	Low	33042	Bacteroides caccae	species	High	47678
Dialister	genus	Low	39948	Bacteroides eggerthii	species	High	28111
Dielma	genus	High	1472649	Bacteroides stercoris	species	Low	46506
Dorea	genus	High	189330	Butyrivibrio crossotus	species	Low	45851
Eggerthella	genus	High	84111	Clostridioides difficile	species	High	1496
Erysipelatoclostridium	genus	High	1505663	Coprococcus catus	species	High	116085
Escherichia	genus	High	561	Eubacterium coprostanoligenes	species	High	290054
Flavonifractor	genus	High	946234	Faecalibacterium prausnitzii	species	Low	853
Holdemanella	genus	High	1573535	Flavobacteriaceae bacterium	species	High	1871037
Klebsiella	genus	High	570	Francisella tularensis	species	Low	263
Lachnolostridium	genus	Low	1506553	Lachnospiraceae bacterium	species	Low	1898203
Lachnospira	genus	Low	28050	Listeria monocytogenes	species	High	1639
Lactobacillus	genus	Low	1578	Mediterraneibacter gnavus	species	High	33038
Megamonas	genus	High	158846	Oxalobacter formigenes	species	High	847
Nesterenkonia	genus	High	57494	Prevotella sp.	species	Low	59823
Oscillibacter	genus	High	459786	Roseburia inulinivorans	species	Low	360807

## 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.

D-glucose {Glucose}

Ethyl alcohol {Grain alcohol}

Ferrum {Iron Supplements} 400 mg/day

N(phosphonomethyl)glycine {glyphosate}

Sleep apnea {partial sleep deprivation}

Sucratose {Splenda} 340 mg/day

Tobacco consumption {Smoking}

## 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}	<i>lactobacillus acidophilus</i> {L. acidophilus}
3,3',4',5,7-pentahydroxyflavone {Quercetin}	<i>Lactobacillus plantarum</i> {L. plantarum}
Avena sativa x Hordeum vulgare {barley,oat}	<i>lactobacillus rhamnosus</i> gg <i>bifidobacterium animalis lactis</i>
bacillus	, <i>lactobacillus paracasei</i> {cvs maximum strength probiotic}
bacillus subtilis {B.Subtilis }	<i>Limosilactobacillus fermentum</i> {L. fermentum}
bacillus,lactobacillus,streptococcus,saccharomyces probiotic	<i>Limosilactobacillus reuteri</i> {L. Reuteri}
bifidobacterium longum {B.Longum }	long-term, moderate-intensity exercise {exercise}
Bovine Milk Products {Dairy}	oligosaccharides {oligosaccharides}
Clostridium butyricum MIYAIRI 588 {Miyarisan}	<i>origanum vulgare</i> {oregano}
Diferuloylmethane {Curcumin}	<i>Panax</i> ... {Ginseng}
enterococcus faecium {E. faecium}	pectin {pectin}
fruit	polyphenols
fruit/legume fibre	Probiotic Mixture 1 {Japanese Vet Probiotic}
Hordeum vulgare {Barley}	Probiotic Mixture 2 {Vetafarm Probiotic}
Lacticaseibacillus casei {L. casei}	<i>Saccharomyces cerevisiae</i> var <i>boulardii</i> {S. boulardii}
Lacticaseibacillus paracasei {L.paracasei}	<i>Thymus vulgaris</i> {thyme}
Lacticaseibacillus rhamnosus {l. rhamnosus}	walnuts
	wheat
	yogurt

## Sample of Literature Used

The following are the most significant of the studies used to generate these suggestions.

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Anorexia Nervosa

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Chronic Lyme  
Chronic Obstructive Pulmonary Disease (COPD)  
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Cognitive Function  
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Constipation  
Coronary artery disease  
COVID-19  
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Cushing's Syndrome (hypercortisolism)  
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deep vein thrombosis  
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Depression  
Dermatomyositis  
Eczema  
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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  
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High Histamine/low DAO  
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hyperglycemia  
Hyperlipidemia (High Blood Fats)  
hypersomnia  
hypertension (High Blood Pressure)  
Hypothyroidism  
Hypoxia

IgA nephropathy (IgAN)  
Inflammatory Bowel Disease  
Insomnia  
Intelligence  
Intracranial aneurysms  
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myasthenia gravis  
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NonCeliac Gluten Sensitivity  
Obesity  
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Peanut Allergy  
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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**