

Microbiome Information for: benign prostatic hyperplasia

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

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

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 benign prostatic hyperplasia

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

Oscillospiraceae	family	Low	216572
Bacteroides	genus	Low	816
Corynebacterium	genus	High	1716
Eisenbergiella	genus	Low	1432051

Bacteria Name Rank Shift Taxonomy ID

Escherichia	genus	High	561
Faecalibacterium	genus	Low	216851
Shigella	genus	High	620
Staphylococcus	genus	High	1279

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.

2-Amino-4-(methylthio)butanoic acid {Methionine} 5 gram/day

Arthrosira platensis {Spirulina}

Chamaemelum nobile {Camomile}

Citrus polymethoxyflavone {Nobiletin (oranges and lemons)}

d-galactose {milk sugar}

Escherichia coli cryodesiccata {colinfant probiotics}

Escherichia coli:DSM 16441-16448 {symbioflor-2}

Glycine max {Doenjang}

humic substances {Humic,Fulvic acids}

Lentilactobacillus kefiri {Kefibios}

Monascus purpureus x Oryza sativa {Red yeast rice}

Morus {Mulberry }

Nicotine, Nicotine Patch

N-L-a-asparty-L-phenylalanine methyl ester {aspartame}

Sodium 2-stearoyllactate {sodium stearoyl lactylate}

Sucralose {Splenda} 340 mg/day

Tobacco consumption {Smoking}

Under cooked animal protein {Rare meat}

Retail Probiotics

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

symbiopharm / symbioflo 2
PIANETA FARMA/KefiBios

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.

5,6-dihydro-9,10-dimethoxybenzo[<i>g</i>]-1,3-benzodioxolo[5,6- <i>a</i>]quinolizinium {Berberine}	Honey {Honey }
(2->1)-beta-D-fructofuranan {Inulin}	Lacticaseibacillus casei {L casei}
2,3-dihydroxypropyl dodecanoate {Monolaurin}	Lacticaseibacillus paracasei {L.paracasei}
arabinogalactan {arabinogalactan}	lactobacillus acidophilus {L acidophilus}
bacillus	Lactobacillus plantarum {L plantarum}
bacillus subtilis {B.Subtilis }	lactobacillus rhamnosus gg bifidobacterium animalis lactis
bacillus,lactobacillus,streptococcus,saccharomyces probiotic	,lactobacillus paracasei {cvs maximum strength probiotic}
bifidobacterium bifidum {B. bifidum}	laminaria digitata {Oarweed}
bifidobacterium longum {B.Longum }	Malus domestica {apple}
Capsicum annuum {Peppers} {Cayenne Pepper, Hot Pepper}	Moringa Oleifera
Cinnamomum zeylanicum {Ceylon Cinnamon}	oligosaccharides {oligosaccharides}
Curcuma longa {Turmeric}	origanum vulgare {oregano}
Diferuloylmethane {Curcumin}	Panax ... {Ginseng}
fruit	pectin {pectin}
fruit/legume fibre	Propolis {Bee glue}
fucoidan {Brown Algae Extract}	resistant starch
Ganoderma sichuanense {Reishi Mushroom}	Saccharomyces cerevisiae var boulardii {S. boulardii}
Hericium erinaceus {Lion's Mane Mushroom }	β -glucan {Beta-Glucan}
Heyndrickxia coagulans {B. coagulans}	Sulforaphane {Dark Greens}
	Thymus vulgaris {thyme}
	yogurt

Sample of Literature Used

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

[Metagenomics of Parkinson`s disease implicates the gut microbiome in multiple disease mechanisms.](#)

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membranous nephropathy
Menopause
Metabolic Syndrome
Mood Disorders
multiple chemical sensitivity [MCS]
Multiple Sclerosis
Multiple system atrophy (MSA)
myasthenia gravis
neuropathic pain
Neuropathy (all types)
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