# **Kyoto Encyclopedia of Genes and Genomes derived Suggestions**

## **Review Overview**

These suggestions 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 almost 2 million facts with backward chaining to sources of information. The typical sources are studies published on the US National Library of Medicine. Note: That many of the bacteria species used are NOT reported on many tests.

These are suggestions that are predicted to independently Decreasing histidine decarboxylase by impacting the bacteria listed on <u>KEGG: Kyoto Encyclopedia of Genes and Genomes</u>. 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

This is an experimental feature – manual validations is recommended. For background, see this post

There is a separate report for probiotics. That report use the enzymes in probiotic species.

### **Analysis Provided by Microbiome Prescription**

A Microbiome Analysis Company

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**Our Facebook Discussion Page** 

# **Bacteria being targeted by suggestions.**

These bacteria levels were deemed atypical

Bacteria Name	Rank	Shitt	onomy ID	Bacteria Name	Rank	Shift Taxonomy ID
Sorangium cellulosum	species		56	Kitasatospora setae	species	2066
Francisella tularensis subsp.	subspecies		264	Halomonas elongata	species	2746
novicida	Subspecies		_0.	Francisella philomiragia	species	<b>2811</b> 0
Pseudomonas fluorescens	species		294	Vibrio nigripulchritudo	species	28173
Methylococcus capsulatus	species		414	Arcobacter nitrofiglis	species	28199
Acinetobacter baumannii	species		470	Ornithobacterium rhinotracheale	species	28251
lodobacter fluviatilis	species		537	Streptantibioticus cattleyicolor	species	29303
Klebsiella aerogenes	species		548	Aeromonas salmonicida subsp.	subspecies	29491
Dickeya chrysanthemi	species		556	salmonicida	Subspecies	23431
Klebsiella pneumoniae	species		573	Vibrio tubiashii	species	29498
Aeromonas hydrophila	species		644	Staphylococcus saccharolyticus	species	33028
Aeromonas salmonicida	species		645	Gloeobacter violaceus	species	33072
Vibrio vulnificus	species		672	Streptomyces subrutilus	species	36818
Vibrio gazogenes	species		687	Xenorhabdus poinarii	species	40577
Plesiomonas shigelloides	species		703	Micromonospora aurantiaca	species	47850
Haemophilus influenzae	species		727	Acinetobacter pittii	species	48296
Bacteroides fragilis	species		817	Tatumella citrea	species	53336
Fusobacterium varium	species		856	Dactylosporangium vinaceum	species	53362
Fusobacterium ulcerans	species		861	Nocardiopsis alba	species	53437
Cellulophaga lytica	species		979	Fructilactobacillus lindneri	species	53444
Solitalea canadensis	species		995	Raoultella omithinolytica	species	54291
Leptolyngbya boryana	species		1184	Erwinia persicina	species	55 <b>211</b>
Clostridium perfringens	species		1502	Vibrio anguillarum	species	55601
Paraclostridium sordellii	species		<b>1505</b>	Streptomyces platensis	species	58346
Clostridium tetani	species		<b>1513</b>	Desulfobacca acetoxidans	species	60893
Acetivibrio thermocellus	species		<b>1515</b>	Shewanella woodyi	species	60961
Clostridium baratii	species		1561	Serratia rubidaea	species	61652
				Zobellia galactanivorans	species	63186
Limosilactobacillus reuteri	species		1598			00000
Fructilactobacillus fructivorans	species		1614	Streptomyces fungicidicus	species	68203
Limosilactobacillus vaginalis	species			Musicola paradisiaca	species	69223
Streptomyces davuligerus	species		1901	Aliivibrio wodanis	species	80852

### 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.

The recommended process to obtain a persistent shift of the microbiome is:

Generate 4 lists from the suggestions with nothing repeated on another list

Emphasize one list each week

After 8 weeks (2 cycles), retest the microbiome to obtains the next set of course corrections

This approach allows the microbiome to stablize towards normal.

Pick only as many suggestions that suits you; there is no need to do all of them. Suggestions are based on your specific bacteria and not marketing concepts such as 'healthy choices'.

Allium cepa (Onion)

Azadirachta indica {Neem} 120 mg/day

Cinnamomum zeylanicum (Ceylon Cinnamon) 6 gram/day

Citrus limon {Lemon}

Coriandrum sativum (Coriander)

**DiferuloyImethane {Curcumin}** 3 gram/day

foeniculum vulgare, fennel

Metha family (Mint)

Nigella sativa {black cumin} 1000 mg/day

Ocimum basilicum (Basil)

origanum vulgare (oregano)

rosmarinus officinalis (rosemary)

**ß-(1?4)-linked D-glucosamine and N-acetyl-D-glucosamine** 

{Chitosan} 3 gram/day

syzygium aromaticum (clove)

tea

Thymus vulgaris (thyme)

Zingiber officinale Roscoe (ginger)

### **Substance to Consider Reducing or Eliminating**

These are the most signigicant 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}
arabinogalactan {arabinogalactan}
bacillus,lactobacillus,streptococcus,saccharomyces probiotic
Ferrum {Iron Supplements}
fructo-oligosaccharides
Grape Polyphenols {Grape Flavonoids}
Helianthus tuberosus {jerusalem artichoke}
Hordeum vulgare {Barley}
Outer Layers of Triticum aestivum {Wheat Bran}

Phaseolus vulgaris {Boston bean}
Pulses, Beans
raffinose {sugar beet}
resistant starch
resveratrol-pterostilbene {grapes, blueberries}
Saccharomyœs cerevisiae var boulardii {S. boulardii}
Sesamum indicum {Sesame}
synthetic disaccharide derivative of lactose {Lactulose}
Ulmus rubra {slippery elm}
walnuts

#### Sample of Literature Used

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

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