

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 Hydrogen | H2 H2 by impacting the bacteria listed on [KEGG: Kyoto Encyclopedia of Genes and Genomes](#). 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

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Bacteria being targeted by suggestions.

These bacteria levels were deemed atypical

Bacteria Name	Rank	Shift	Taxonomy ID	Bacteria Name	Rank	Shift	Taxonomy ID
				Legionella israelensis	species		454
Persephonella marina	species		309805	Legionella jordanis	species		456
Sulfurihydrogenibium azureum	species		309806	Desulfovibrio piger	species		901
Desulforapulum autotrophicum	species		2296	Hydrogenobacter thermophilus	species		940
Streptomyces nodosus	species		40318	Allochromatium vinosum	species		1049
Desulfurobacterium thermolithotrophum	species		64160	Prosthecochloris aestuarii	species		1102
Desulfocapsa sulfexigens	species		65555	Mycobacterium kansasii	species		1768
Streptomyces globosus	species		68209	Mycolicibacterium smegmatis	species		1772
Thioalkalivibrio paradoxus	species		108010	Mycobacterium gordonae	species		1778
Caldilinea aerophila	species		133453	Mycobacterium marinum	species		1781
Thermovibrio ammonificans	species		228745	Mycolicibacterium thermoresistibile	species		1797
Desulfovibrio ferrophilus	species		241368	Mycolicibacterium aichiense	species		1799
Adlercreutzia equolifaciens	species		446660	Mycolicibacterium chubuense	species		1800
Gordonibacter pamelaee	species		471189	Mycolicibacterium diernhoferi	species		1801
Thiohalobacter thiocyanaticus	species		585455	Streptomyces reticuli	species		1926
Neptunomonas concharum	species		1031538	Streptomyces rimosus	species		1927
Magnetospira sp. QH-2	species		1288970	Streptomyces scabiei	species		1930
Salinivirga cyanobacteriivorans	species		1307839	Streptosporangium roseum	species		2001
Labilithrix luteola	species		1391654	Thermobispora bispora	species		2006
Desulfocurvibacter africanus subsp. africanus	subspecies		1511600	Desulfomonile tiedjei	species		2358
Limnochorda pilosa	species		1555112	Legionella cherrii	species		28084
Sulfurifustis variabilis	species		1675686	Legionella sainthelensi	species		28087
Candidatus Velamenicoccus archaeovorus	species		1930593	Thermodesulfovibrio yellowstonii	species		28262
Thiocapsa sp.	species		2024551	Legionella oakridgensis	species		29423
Streptomyces tirandamycinicus	species		2174846	Rhodothermus marinus	species		29549
Labrenzia sp. PHM005	species		2590016	Halothermothrix orenii	species		31909
Dictyoglomus thermophilum	species		14	Streptomyces laurentii	species		39478
Legionella pneumophila	species		446	Mycolicibacterium duvalii	species		39688
Legionella longbeachae	species		450	Mycobacterium branderi	species		43348
Legionella micdadei	species		451	Kutzneria albida	species		43357
Legionella spiritensis	species		452				

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 stabilize 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'.

berberine 1.5 gram/day

iron 400 mg/day

Prescript Assist (2018 Formula)

Slippery Elm

triphala 9000 mg/day

vegetarians

walnuts 75 gram/day

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.

Arbutin (polyphenol)	melatonin supplement
Cacao	N-Acetyl Cysteine (NAC),
chitosan,(sugar)	peppermint (spice, oil)
cinnamon (oil. spice)	retinoic acid,(Vitamin A derivative)
Curcumin	rosmarinus officinalis,rosemary
diosmin,(polyphenol)	syzygium aromaticum (clove)
Hesperidin (polyphenol)	thyme (thymol, thyme oil)
Human milk oligosaccharides (prebiotic, Holigos, Stachyose)	Vitamin B1,thiamine hydrochloride
lactobacillus casei (probiotics)	Vitamin B-12
lactobacillus paracasei (probiotics)	vitamin B3,niacin
luteolin (flavonoid)	Vitamin B6,pyridoxine hydrochloride
	Vitamin C (ascorbic acid)

Sample of Literature Used

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

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