

## Microbiome Information for: Lung Cancer

### 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 *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 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](mailto:Research@MicrobiomePrescription.com)

[Our Facebook Discussion Page](#)

## Bacteria being reported because of atypical values.

These bacteria were reported atypical in studies of Lung Cancer

*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      |
|------------------|---------------|-------|----------|---------|------------------------------|----------------|-------|----------|---------|
| Clostridia       | <i>class</i>  | Low   |          | 186801  | Hungatella                   | <i>genus</i>   | High  |          | 1649459 |
| Barnesiellaceae  | <i>family</i> | Low   |          | 2005519 | Ruminococcus                 | <i>genus</i>   | Low   |          | 1263    |
| Lachnospiraceae  | <i>family</i> | Low   |          | 186803  | Shigella                     | <i>genus</i>   | High  |          | 620     |
| Oscillospiraceae | <i>family</i> | Low   |          | 216572  | Eubacteriales                | <i>order</i>   | Low   |          | 186802  |
| Tannerellaceae   | <i>family</i> | Low   |          | 2005525 | [Eubacterium] siraeum        | <i>species</i> | Low   |          | 39492   |
| Anaerotruncus    | <i>genus</i>  | High  |          | 244127  | Akkermansia muciniphila      | <i>species</i> | Low   |          | 239935  |
| Blautia          | <i>genus</i>  | Low   |          | 572511  | Alistipes onderdonkii        | <i>species</i> | Low   |          | 328813  |
| Dorea            | <i>genus</i>  | Low   |          | 189330  | Anaerobutyricum hallii       | <i>species</i> | Low   |          | 39488   |
| Escherichia      | <i>genus</i>  | High  |          | 561     | Bacteroides caccae           | <i>species</i> | High  |          | 47678   |
|                  |               |       |          |         | Faecalibacterium prausnitzii | <i>species</i> | Low   |          | 853     |

## 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.

alcoholic beverages

Apple skin

**bacillus amyloliquefaciens (probiotic)** 1 BCFU/day

bentonite

**bifidobacterium pseudocatenulatum li09,bifidobacterium catenulatum li10 (probiotics)**

**brassica juncea**

**camelina seed**

**candida albicans (prescription)**

**carboxymethyl cellulose (prebiotic)**

**colostrum**

**lactobacillus delbrueckii bulgaricus,bifidobacterium**

**bifidum,enterococcus faecium,candida pintolopesii,aspergillus oryzae (probiotics)**

**lactobacillus salivarius (probiotics)** 9 BCFU/day

**lactulose**

**lard**

**l-citrulline**

**l-glutamine** 5 gram/day

**linseed(flaxseed)** 30 mg/day

**lupin kernel fiber (legume) (anaphylaxis risk, toxic if not prepared properly)**

**mannooligosaccharide (prebiotic)** 8 gram/day

**Methionine** 5 gram/day

**Psyllium (Plantago Ovata Husk)** 6.8 gram/day

**quercetin,resveratrol**

**rare meat**

**red alga Laurencia tristicha**

**smoking**

**sodium stearoyl lactylate**

**sorghum**

**stevia** 800 mg/day

**symbioflor 2 e.coli probiotics**

**Tributylin**

**Vitamin B9,folic acid** 5 mg/day

## **Retail Probiotics**

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

symbiopharm / symbioflo 2  
Bromatech (IT) / Acronelle  
klair labs / target gb-x  
Sash Vitality / Bio-Cultures Probiotics for Adults  
SuperSmart / Vaginal Health  
CustomProbiotics.com / L. Salivarius Probiotic Powder  
Nu U (uk) / Bio-Cultures Complex  
Maple Life Science™ / Lactobacillus salivarius  
wakamoto (jp) / strong wakamoto w  
HLH BIOPHARMA(DE) / LACTOBACT ® OMNI FOS  
Metabolics / Lactobacillus Salivarius Powder  
ecology\_allergycare

**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.

arabinogalactan (prebiotic)

bacillus subtilis (probiotics)

barley

bifidobacterium longum (probiotics)

fasting

garlic (allium sativum)

inulin (prebiotic)

lactobacillus paracasei (probiotics)

lactobacillus plantarum (probiotics)

lactobacillus reuteri (probiotics)

lactobacillus rhamnosus gg (probiotics)

quebracho

resistant starch

saccharomyces boulardii (probiotics)

soy

vitamin d

wheat

## Sample of Literature Used

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

### [Gut microbial signature in lung cancer patients highlights specific taxa as predictors for durable clinical benefit.](#)

**Scientific reports** , Volume: 13 Issue: 1 2023 Feb 3

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### [Attempting to Identify Bacterial Allies in Immunotherapy of NSCLC Patients.](#)

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### [Impact of Gut Dysbiosis on the Risk of Non-Small-Cell Lung Cancer.](#)

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gallstone disease (gsd)  
Gastroesophageal reflux disease (Gerd) including Barrett's esophagus  
Generalized anxiety disorder  
giant cell arteritis  
Glioblastoma  
Gout  
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Halitosis

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Heart Failure  
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hyperglycemia  
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Multiple system atrophy (MSA)  
myasthenia gravis  
neuropathic pain  
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Nonalcoholic Fatty Liver Disease (nafld) Nonalcoholic  
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Osteoarthritis  
Osteoporosis  
pancreatic cancer  
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Premenstrual dysphoric disorder  
primary biliary cholangitis  
Psoriasis  
rheumatoid arthritis (RA), Spondyloarthritis (SpA)  
Rosacea  
Schizophrenia  
scoliosis  
sensorineural hearing loss  
Sjögren syndrome  
Sleep Apnea  
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**