



HOMEOSTASIS DIET

The Gut Microbiome



You Are What You Eat

I am so grateful that you have taken the time to read this message. Before you continue, I urge you to take a moment to deeply inhale and exhale. Allow yourself to relax and let go of any stress you may be holding onto. You are on the brink of a truly transformative discovery. We all yearn for something that can improve our lives in a meaningful way, and I believe that you have the potential to find just that here. Trust in yourself and keep moving forward and the answer you seek will find you.

The gut microbiome is a crucial aspect of overall health and well-being. The old adage "you are what you eat" has never been more true than when it comes to the health of the gut. In fact, we have coined a new phrase, "mind your gut and you'll keep your mind," to emphasize the importance of gut health in maintaining a healthy mind and body. By understanding the gut microbiome and taking steps to maintain a healthy balance of bacteria, we can improve our digestion, boost our immune system, and even prevent chronic diseases. Don't neglect your gut health any longer, start paying attention to what you eat and how it affects your gut microbiome today.

I urge you to take advantage of the valuable resources we have made available in our PDFs across this website. They are specifically designed to educate and provide complete transparency on various processes. By reading and utilizing the information provided, you will not only empower yourself with knowledge, but also contribute to building a brighter future for all of us. Together, we can achieve a world where all the answers you seek are within your reach. So, take the time to read and learn from these resources, and be a part of the positive change we are striving for.

If you belly is your brain, what are you feeding it?

Our Process



Genomic Nutritional Data

Our Gut Microbiome Test package includes the following:

- Review from a naturopathic doctor who specializes in gut microbiome health
- Three one-hour consultations with a gut restoration dietitian via zoom
- Recommendations for pre and probiotic supplements
- Recommendations for foods to reinoculate the gut microbiome
- A personalized diet plan with meals included
- A grocery shopping list
- An invitation to our "reset button" program ([more information can be found here](#))

The cost of this package is \$1,200.

If you want to add genomics testing, the total cost is \$1,800.

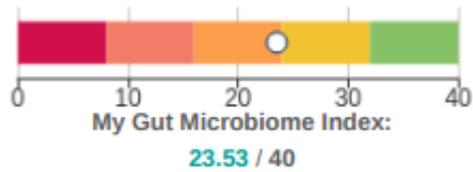
Alternatively, you can purchase just the tests without professional review for \$600.

If you are interested in this package or have any additional questions, you can click [here](#) to book a call.

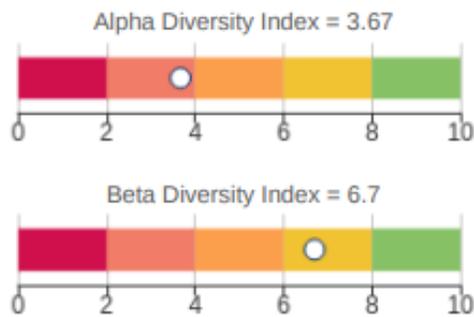
<https://calendly.com/semc3/microbiome>

Report Summary

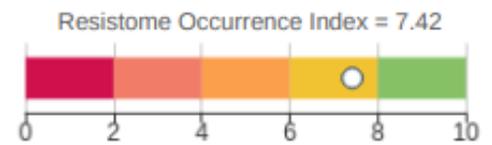
Gut Microbiome Index



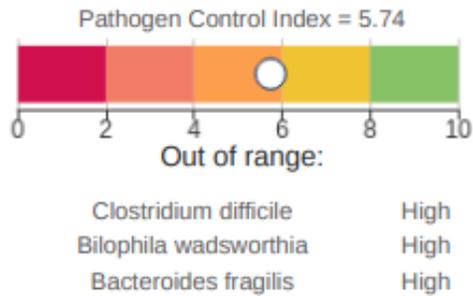
Diversity



AMR Richness



Pathogens



Keystone Species

Out of range:

Akkermansia muciniphila	High
Ruminococcus bromii	Low
Ruminococcus flavefaciens	Low
Roseburia intestinalis	Low
Bifidobacterium longum	High
Butyrivibrio pullicaecorum	Low
Lactobacillus rhamnosus	Low
Lactobacillus reuteri	Low
Total Lactobacillus species	Low

Functions

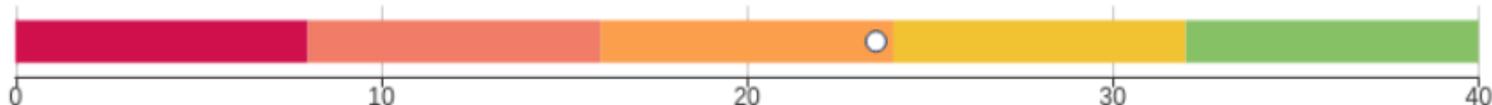
Out of range:

Propionate production	High
Indole production	High
Vit B9 - Folate	High
Vit B12 - Cobalamin	High
Vitamin biosynthesis	High

Dysbiosis

Out of range

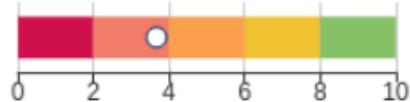
Proteobacteria: Actinobacteria Low



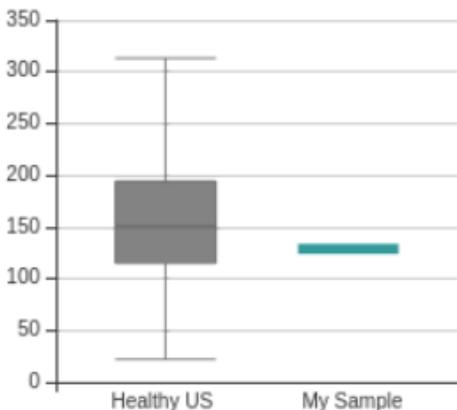
The Gut Microbiome Index (GMI) is an overall score for gut microbiome health. A score above 30 is considered excellent. It is calculated by assessing four key indicators of microbiome health for your gut microbiome and comparing them to the typical healthy gut microbiome. The four key indicators include **Alpha Diversity** (species richness), **Beta Diversity** (composition), **Pathogen Occurrence** (population of pathogens) and **Resistome Occurrence** (population of antibiotic resistance genes).

My Gut Microbiome Alpha- and Beta-Diversity

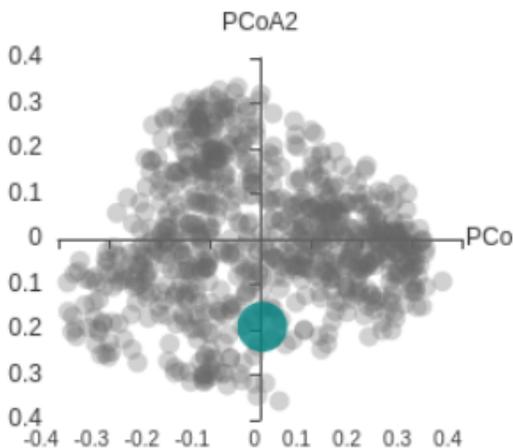
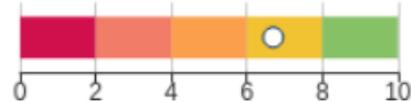
Alpha Diversity Index = 3.67



Healthy US My Sample



Beta Diversity Index = 6.7



Number of species in gut microbiome: **129**

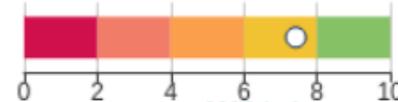
In ecology, **Alpha-Diversity** refers to the average diversity, or the richness of species, in a particular ecosystem. This marker is looking at your own personal species richness within your gut microbiome.

A Low Alpha-Diversity Index suggests that your gut microbiome was recently damaged by antibiotics, environmental toxins, stress, diet, or other factors.

Beta-diversity is the variation of species when comparing the composition of two separate ecosystems. This marker compares the composition of your gut microbiome to healthy populations in order to illustrate notable differences. The green dot for your sample not falling within the clusters of grey dots (healthy) leads to a low Beta-Diversity Index suggesting that your gut microbiome composition is trending away from a healthy gut to an imbalanced, dysbiotic gut.

My Gut Stability and Uniformity

Resistome Occurrence Index = 7.42



AMR index

AMR index indicates distance between my sample and the average Healthy microbiome



This section explores the richness and stability of your gut microbiome by comparing the resilience of your gut microbiome to Healthy populations. A low index suggests that you have low richness and resilience in your gut.



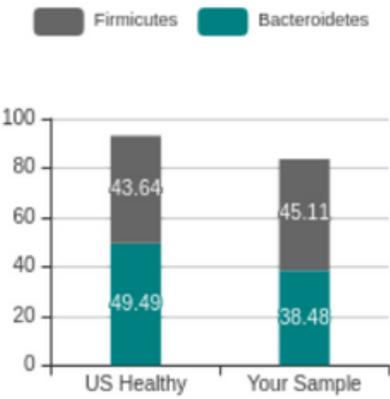
Dysbiosis Ratios

This section compares the abundances of important groups of gut bacteria between your gut and the typical healthy gut microbiome. Elevated dysbiosis ratios for these bacterial phyla or genera point to imbalances in abundance (dysbiosis) which are associated with a range of health conditions.

Description	Healthy Ratio IQR	My Ratio
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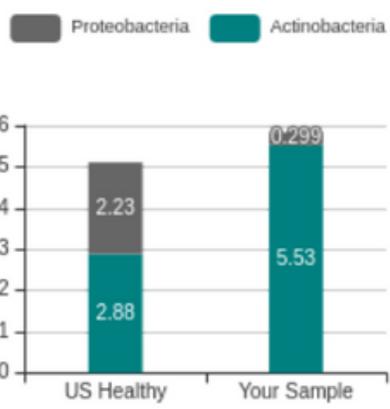
Firmicutes:Bacteroidetes (F/B) Ratio
 In adults, Firmicutes and Bacteroidetes are the most abundant bacterial phyla in the gut. The Firmicutes to Bacteroidetes (F/B) abundance ratio was shown to increase from infancy to adulthood and subsequently to decrease again in the elderly. Healthy F/B ratios differ significantly between studies and can exceed 0.25 (shown here), in some studies even 1.

0.42 ~ 1.73	1.17
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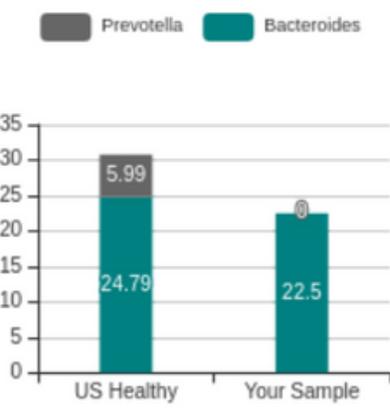
Proteobacteria:Actinobacteria (P/A) Ratio
 Together these phyla comprise about 10% of total gut microbes. Healthy adults tend to have no more than 4.5% Proteobacteria. Proteobacteria:Actinobacteria (P/A) ratios less than 1.0 are associated with healthy metabolism and cell turnover. Increasing Actinobacteria can be accomplished by consuming plant-derived carbohydrate starch and polysaccharides, such as FOS, GOS, XOS, inulin or arabinoxylan.

0.28 ~ 5.64	0.05
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Prevotella:Bacteroides (P/B) Ratio
 High Prevotella:Bacteroides ratios are associated with lower BMI and reduced incidence of chronic, inflammatory disease.
 Low Prevotella:Bacteroides ratios (smaller numbers) are associated with metabolic imbalances and are positively correlated with high intake of protein and animal fat as typical for a Western diet. Higher abundance of Prevotella is observed in individuals that consume diets rich in carbohydrates and fiber. Bacteroides is increased by sugar and saturated fat intake, while Prevotella generally thrives on fiber rich foods, like fruit, vegetables, beans and whole grains. Levels of Prevotella tend to decrease with age, particularly among centenarian populations.

0.0 ~ 0.52	0
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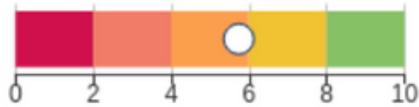




Pathobiome

Low levels of pathogens can be normal and characteristic of a healthy, diverse gut microbiome. Increased levels of pathogens however could indicate that a pathogen is playing a role in symptoms you are experiencing. This section compares the relative abundances (RA) of specific pathogens to normal levels present in the healthy gut.

Pathogen Control Index = 5.74



BiomeFX is NOT a diagnostic test. If your Pathogen levels are abnormally high consult your physician who can make a diagnosis and provide treatment if needed.

Pathogen Species	Healthy RA IQR %	My Sample RA %
Clostridium difficile	0.02 - 0.17	0.3376
Bilophila wadsworthia	0.05 - 0.18	0.2193
Bacteroides fragilis	0.07 - 0.79	2.3875

Virulence Factors Tested in Your Sample

Virulence factors are genes that when transcribed enable a microorganism to establish itself on or within a host and enhance its potential to cause disease (they help bacteria invade the host, cause disease, and evade host defenses), and include bacterial toxins, cell surface proteins that mediate bacterial attachment, cell surface carbohydrates and proteins that protect a bacterium, and hydrolytic enzymes that may contribute to the pathogenicity of the bacterium.

Virulence Factors	My Sample
Clostridiodes difficile cdtA	Not Detected
Clostridiodes difficile cdtB	Not Detected
Clostridiodes difficile tcdA	Not Detected
Clostridiodes difficile tcdB	Not Detected
Bacteroides fragilis BFT	Not Detected



Full List of Pathogen Species Tested

Bacteria

Bacteroides fragilis	Bilophila wadsworthia
Campylobacter	Campylobacter jejuni
Citrobacter freundii	Clostridiodes difficile
Clostridiodes perfringens	Enterococcus gallinarum
Escherichia coli	Hafnia alvei
Helicobacter pylori	Klebsiella pneumoniae
Porphyromonas gingivalis	Proteus mirabilis
Proteus vulgaris	Pseudomonas aeruginosa
Raoultella ornithinolytica	Salmonella enterica
Sutterella stercoricanis	Sutterella wadsworthensis
Vibrio cholerae	Yersinia enterocolitica

Protists

Blastocystis hominis	Cryptosporidium
Cyclospora cayetanensis	Entamoeba histolytica
Giardia lamblia	

Fungi

Candida	Geotrichum spp
Microsporidia spp	Rhodotorula spp

Viruses

Adenovirus	Cytomegalovirus
Epstein Barr Virus	



Functional Keystone Species in My Gut

Keystone species are beneficial bacteria that have a disproportionately large effect on both their habitat and the status of other microbial communities of the gut. Keystone species create an environment that is unfriendly to pathogens yet allows good gut microbes (commensal) to thrive. This section compares relative abundances between your gut and the healthy gut microbiome.

Keystone Species	Function	Healthy Relative Abundance IQR Range[%]	My Sample Relative Abundance
Phylum: Verrucomicrobia			
Akkermansia muciniphila	Acetate Producer	0.15 - 2.295	8.337
Phylum: Actinobacteria			
Bifidobacterium longum	Acetate Producer	0.127 - 1.268	1.467
Bifidobacterium adolescentis	GABA Producer	0.077 - 1.536	0.092
Phylum: Firmicutes			
Faecalibacterium prausnitzii	Butyrate Producer	0.675 - 2.032	0.927
Ruminococcus bromii	Cellulose Degradar	0.155 - 1.391	Not Detected
Ruminococcus flavefaciens	Cellulose Degradar	0.007 - 0.014	Not Detected
Roseburia intestinalis	Butyrate Producer	0.062 - 1.116	Not Detected
Eubacterium rectale	Butyrate Producer	0.665 - 2.238	0.765
Butyricoccus pullicaecorum	Butyrate Producer	0.011 - 0.04	Not Detected
Lactobacillus rhamnosus	Lactate Producer	0.016 - 0.051	Not Detected
Lactobacillus reuteri	Lactate Producer	0.011 - 0.016	Not Detected



Short Chain Fatty Acid (SCFA) Producers

In the functional analysis section we measure the genes and pathways identified for SCFA production. In this section we are measuring the abundance of certain bacteria that are known to produce SCFAs. Frequently, the keystone species known to produce these SCFAs are found at low abundance in the microbiome. This means that the entire genome may not be available, including the genes and pathways for SCFA production. This table provides the aggregated relative abundance of the SCFA producing bacteria.

SCFA Name	Healthy IQR	My Sample	My Sample Percentile [%]
Acetate Producers	5.23 - 13.99	5.78	29
Butyrate Producers	2.67 - 5.72	3.28	34
D-Lactate Producers	3.30 - 8.79	1.95	12
L-Lactate Producers	1.10 - 6.46	7.41	79
Propionate Producers	6.68 - 15.16	9.77	43

This section explores your gut microbiome for genes known to contribute metabolically important functions. A higher value means that more microbial genes contributing to a function have been identified. A low value in your gut (or the typical healthy gut) microbiome does not mean that your metabolite levels are low. It only reflects the extent to which your gut microbiome can contribute to your levels. The report shows absolute values and your functional gut microbiome composition as a percentile relative to the typical healthy microbiome.

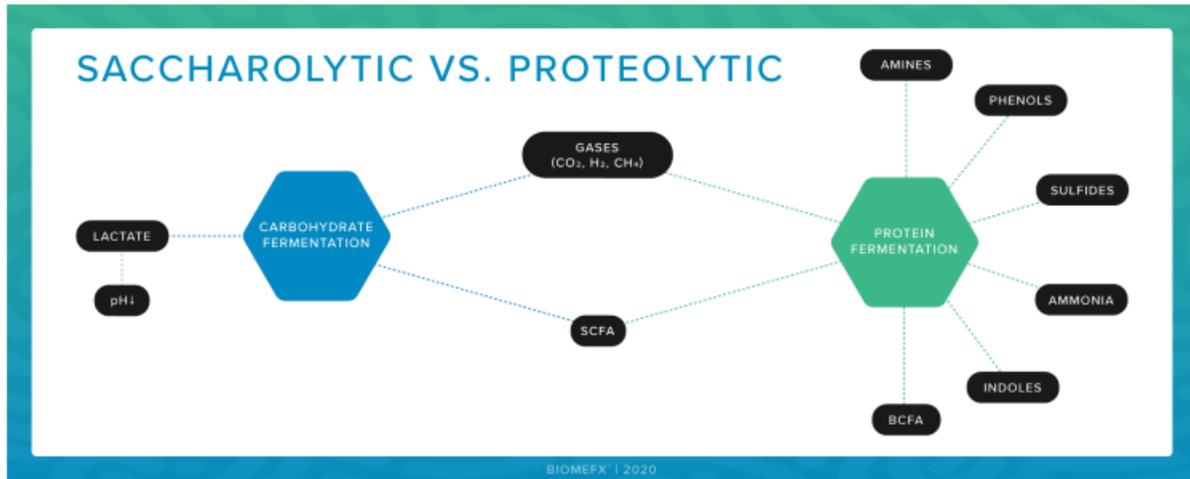
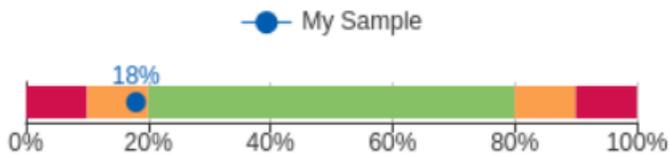


Figure 1. This image shows a comparison of the byproducts that result from carbohydrate and protein fermentation in the gut microbiome.

Saccharolytic fermentation

Summary:



US Healthy Population Percentiles

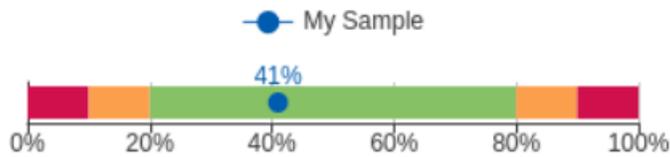
Report Descriptions

Gut bacteria prefer to ferment carbohydrates rather than protein. Saccharolytic fermentation produces short-chain fatty acids (SCFAs), like butyrate, acetate, and propionate, as by-products. These SCFAs are the preferred energy source of intestinal cells and, as a result, can support healthy gut barrier function.

Higher levels of saccharolytic fermentation are associated with healthy metabolism and reduced intestinal pH, which supports a healthy gut environment. Saccharolytic fermentation levels may be low as a result of a high-protein diet, low-fiber diet, or insufficient keystone species.

My Gut	US Healthy Range
630	535 to 1573

Butyrate production

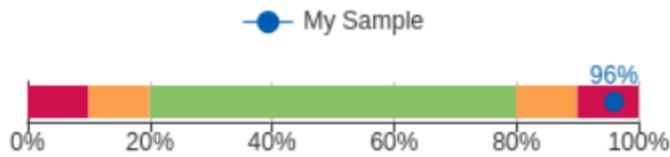


My Gut	US Healthy Range
496	256 to 1090

Report Descriptions

Butyrate is arguably the most important SCFA, yet it comprises only 15-20% of total SCFA production. Butyrate enhances intestinal barrier function, acts as a fuel source for enterocytes, scavenges ammonia, regulates the immune system, reduces oxidative stress, and much more. Butyrate production is mostly associated with microbial fermentation of fibers such as bran, oligosaccharides, arabinoxylan, resistant starches, and others. Furthermore, butyrate production requires an acidic environment in the gut.

Propionate production

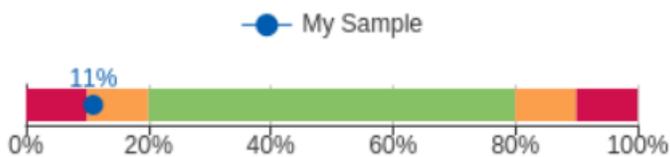


My Gut	US Healthy Range
1932	132 to 1523

Report Descriptions

Propionate is a short-chain fatty acid that can be produced by gut bacteria through the fermentation of key fibers or the metabolism of lactate. Propionate supports a healthy immune system by encouraging regulatory T cell differentiation in gut associated lymphoid tissues (GALT), and it also promotes gluconeogenesis in the liver, supports insulin sensitivity and improves gut hormone production. Propionate and butyrate both work together to support healthy inflammatory responses by inhibiting histone deacetylases (HDACs) in macrophages and dendritic cells.

Acetate production

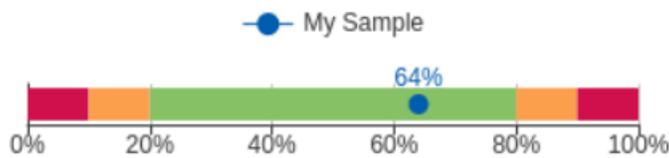


My Gut	US Healthy Range
630	616 to 1870

Report Descriptions

Acetate is another short-chain fatty acid produced by gut bacteria through the fermentation of prebiotic fibers like inulin and GOS or unabsorbed peptides and fats. Gut-derived acetate production is tightly regulated within the microbiome and determined by the presence of prebiotic fiber and the balance between saccharolytic and proteolytic fermentation. Acetate is used for cholesterol synthesis and lipogenesis but can also be utilized by muscle tissue. Additionally, some gut bacteria like *Roseburia* spp and *Faecalibacterium prausnitzii* can convert acetate into butyrate. Excessive acetate production combined with insufficient butyrate production can lead to fat gain, particularly around the liver.

Lactate production



US Healthy Population Percentiles

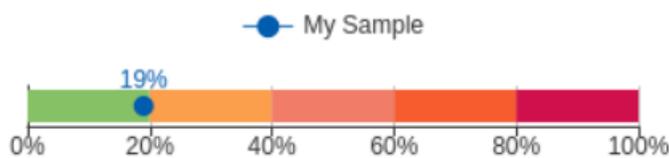
Report Descriptions

Lactate is an intermediate of carbohydrate metabolism, produced from pyruvate during lactic acid fermentation. Lactate also plays important roles in immunomodulation and inflammation modulation. These species use lactate as a substrate for short-chain fatty acid production. However, if there is an overabundance of lactate producers paired with low abundance of lactate utilizers (SCFA producers) this will cause a surge of lactate in the gut which can be toxic and harmful to host tissues.

My Gut	US Healthy Range
3011	1641 to 3913

Proteolytic fermentation

Summary



US Healthy Population Percentiles

Report Descriptions

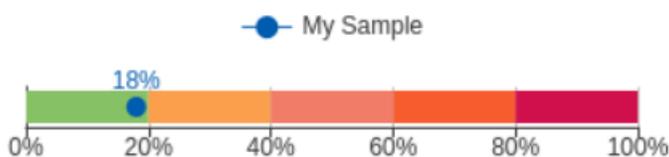
While both saccharolytic and proteolytic fermentation can yield beneficial SCFAs, studies show that protein fermentation also produces unfavorable metabolites like p-cresol, phenol, ammonia, and H₂S that can increase inflammation in the body. Protein-degrading microbes generally only ferment protein after all dietary carbohydrates have been utilized.

Higher levels of proteolytic fermentation are associated with gut dysbiosis and inflammation. Proteolytic fermentation levels may be low when the gut microbiome is more diverse and utilizing more saccharolytic fermentation.

My Gut	US Healthy Range
1688	1358 to 3464

Amines

Polyamine production



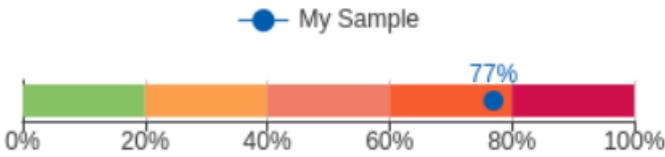
US Healthy Population Percentiles

Report Descriptions

Polyamines like putrescine, spermidine, and cadaverine are metabolites of arginine and tyrosine that have many important roles in the gut like stabilizing RNA and DNA structures, supporting protein synthesis, and scavenging free radicals. However, high amounts of polyamines can be toxic to the gut microbiome. Gut bacteria primarily synthesize amines from amino acids. Generally speaking, Gram-positive bacteria tend to reduce the concentration of amines, while Gram-negative species produce amines and increase their concentration.

My Gut	US Healthy Range
1688	1368 to 3480

Phenols



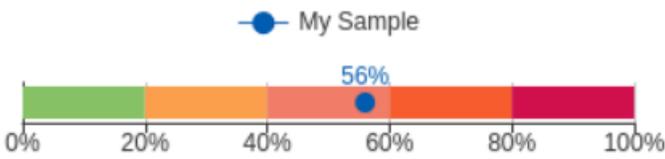
US Healthy Population Percentiles

Report Descriptions

P-cresol is a byproduct of tyrosine metabolism by gut bacteria that can be toxic to intestinal cells and impair intestinal barrier function. P-cresol is also toxic to a wide range of gut microbiota, particularly Gram-negative species.

My Gut	US Healthy Range
73	8 to 121

Ammonia production



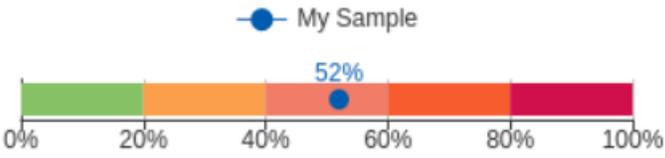
US Healthy Population Percentiles

Report Descriptions

Ammonia is a normal by-product of amino acid fermentation by gut microbes. Ammonia is also produced in the small intestine through the bacterial degradation of glutamine. Healthy liver and kidneys can filter and excrete ammonia through the urine.

My Gut	US Healthy Range
0.0787	0.01 to 0.26

Hydrogen Sulfide (H2S) production



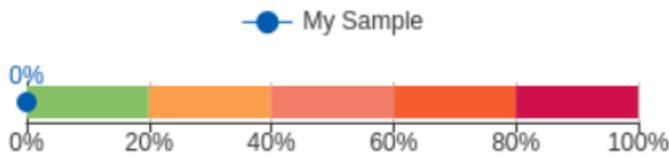
US Healthy Population Percentiles

Report Descriptions

Sulfate-reducing bacteria (SRB) convert dietary sulfur and taurine to H₂S, or hydrogen sulfide, a toxic compound that impairs intestinal detoxification pathways and can cause gas that smells like rotten eggs. H₂S production is associated with high-protein, low-fiber diets. Sulfate-reducing bacteria compete with methane-producers and acetate-producers for the same H₂ substrate; and as a result, it is important to maintain a delicate balance among all 3.

My Gut	US Healthy Range
0.0778	0.02 to 0.38

Methane production



US Healthy Population Percentiles

Report Descriptions

Methanogens convert acetate, ammonia, hydrogen gas, and trimethylamines (TMA) to methane gas. Methane gas also slows the intestinal transit and affects gut motility, which may also allow increased time for nutrient absorption. Furthermore, methane producers compete with acetate producers for substrate utilization, which may explain why methanogens are indirectly associated with digestive issues.

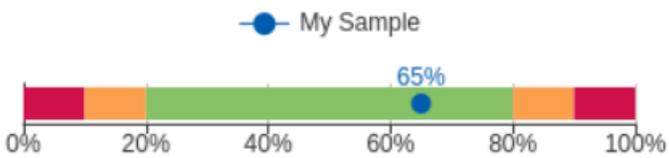
My Gut	US Healthy Range
Not Detected in my sample	0.01 to 0.91

Hormones & Neurotransmitters

Hormones are produced in endocrine glands and are transmitted through the blood stream. Neurotransmitters are released by presynaptic nerve terminal into the synapse and transmitted across the synaptic cleft.

Psychobiome

GABA production



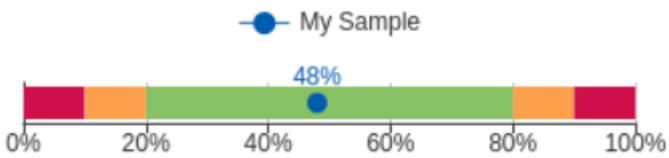
US Healthy Population Percentiles

Report Descriptions

Gamma-aminobutyric acid (GABA) is a neurotransmitter, or chemical messenger, in the brain that blocks specific signals in the central nervous system in order to slow down the brain. This provides a protective and calming effect on the brain and body. High fat diets are shown to reduce GABA levels in the prefrontal cortex by 40% which can result in various mood imbalances and difficulty sleeping.

My Gut	US Healthy Range
386	87 to 767

Glutathione production



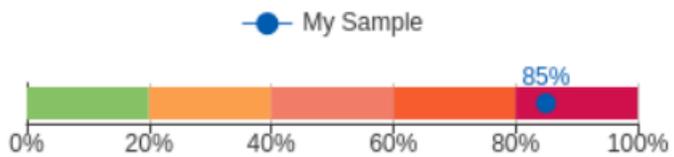
US Healthy Population Percentiles

Report Descriptions

Glutathione is the most powerful antioxidant in the human body. It is found in nearly every cell in the body and is the primary agent of detoxification in the liver. Glutathione can also act as a hormone, regulating the release of GABA and dopamine. Glutathione is produced from three amino acids glutamate, cysteine, and glycine which are obtained from food or supplementation. Deficiency in glutathione may lead to production of free radicals and oxidative damage throughout the body. Recent evidence suggests that the gut microbiome determines levels of glutathione throughout the body.

My Gut	US Healthy Range
116	42 to 232

Histamine production



My Gut

0.2135

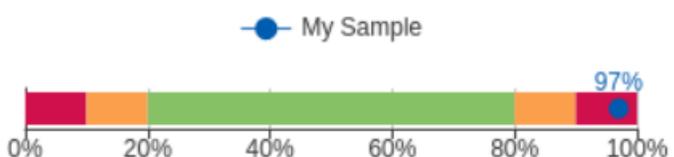
US Healthy Range

0 to 0.31

US Healthy Population Percentiles
[Report Descriptions](#)

Histamine can be produced in the gut and travel to distant areas of the body, where it may induce unfavorable symptoms. Patients with an overabundance of histamine-producing bacteria should focus on strengthening intestinal barrier function, as a leaky gut can allow gut-derived histamines to enter circulation and promote dietary intolerances or disruption in healthy allergic responses. High levels of gut-derived histamine are associated with high abundance of Proteobacteria, Roseburia, Morganii morganii, and Klebsiella pneumoniae and decreased abundance of Bifidobacterium

Indole production



My Gut

338

US Healthy Range

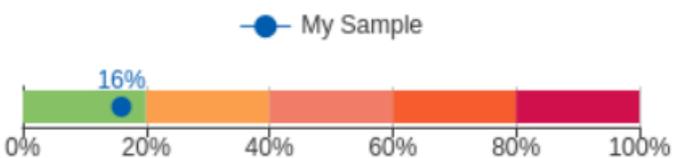
13 to 286

US Healthy Population Percentiles
[Report Descriptions](#)

Indole is a byproduct of the microbial degradation of tryptophan that can be utilized in a variety of ways in the gut microbiome. Indole can bind to serotonin receptors in order to regulate behavior, gut motility, and food intake, and it can support immune and intestinal health by interacting with gut microbes, scavenging free radicals, and increasing the expression of xenobiotic-metabolizing enzymes like cytochrome P450. Indole also functions as a signaling molecule that may be increased during latent infections. Indole production must be balanced, as too much indole may produce unwanted changes in mood or cognition, yet insufficient indole production may damage the gut barrier.

Sex Hormones

Estrogen recycling (Estrobolome)



My Gut

0.0667

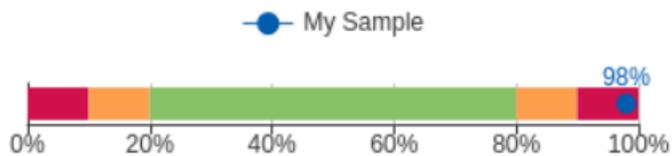
US Healthy Range

0.06 to 0.24

US Healthy Population Percentiles
[Report Descriptions](#)

The estrobolome is a network of over 60 genera of bacteria that can recycle or deconjugate inactivated estrogens for reabsorption into circulation by producing very powerful enzymes. This recycling process is handled by gut bacteria with β -glucuronidase and β -glucosidase activity. When the estrobolome is too abundant, the body is unable to efficiently eliminate estrogen, causing estrogens to build up and ultimately leading to estrogen dominance. On the other hand, if estrogen recycling (estrobolome) is too low, then this may lead to insufficient levels of estrogen in circulation.

Vitamin Biosynthesis



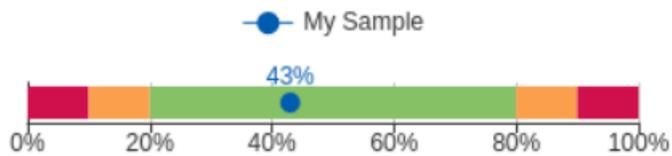
My Gut	US Healthy Range
6278	876 to 3843

US Healthy Population Percentiles

Report Descriptions

Gut bacteria synthesize vitamin K2 and many of the B vitamins including biotin (B7), cobalamin (B12), folates (B8), nicotinic acid (B3), pantothenic acid (B5), pyridoxine (B6), riboflavin (B2), and thiamine (B1). Vitamin production levels may be low as a result of low alpha- and beta-diversity in the gut.

Vit B1 Thiamin



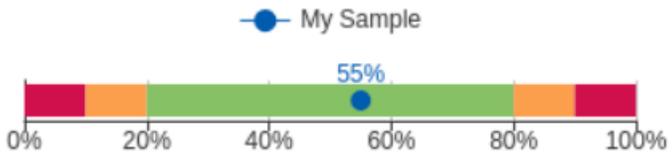
My Gut	US Healthy Range
3302	2000 to 5183

US Healthy Population Percentiles

Report Descriptions

Thiamin is a vitamin that plays a critical role in energy metabolism, especially in the brain and nervous system. Thiamin also plays an important role in muscle contraction and nerve conduction. Faecalibacterium spp utilize thiamine but do not produce it, indicating that there is a competition for vitamins within the gut microbiome.

Vit B2 Riboflavin



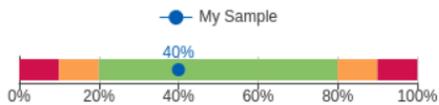
My Gut	US Healthy Range
3776	2048 to 5504

US Healthy Population Percentiles

Report Descriptions

Vitamin B2 (riboflavin) is a cofactor need for energy production and fat metabolism that also plays important roles in immune cell function.

Vit B5 - Pantothenic acid



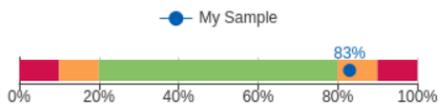
My Gut	US Healthy Range
10069	7372 to 13593

US Healthy Population Percentiles

Report Descriptions

Vitamin B5 (pantothenic acid) is essential for energy production and fat metabolism. Bacteroides fragilis, Prevotella copri, Ruminococcus spp, Salmonella enterica, and Helicobacter pylori can all produce vitamin B5 in the gut. However, there are many species that rely on vitamin B5 for growth but cannot synthesize it, like most Fusobacterium, Bifidobacterium spp, Faecalibacterium spp, Lactobacillus spp, and some strains of Clostridium difficile, suggesting that these bacteria may compete with the host for vitamin B5.

Vit B6 - Pyridoxine



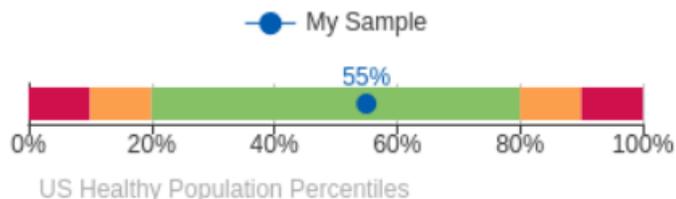
My Gut	US Healthy Range
6821	1051 to 8271

US Healthy Population Percentiles

Report Descriptions

Vitamin B6 (pyridoxine) is an incredibly versatile nutrient that supports immunity, brain function, and protein metabolism.

Vit B2 Riboflavin



Report Descriptions

Vitamin B2 (riboflavin) is a cofactor need for energy production and fat metabolism that also plays important roles in immune cell function.

My Gut

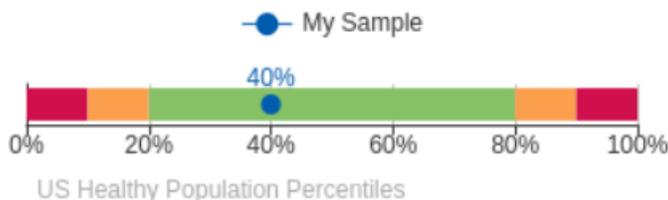
3776

US Healthy Range

2048 to 5504

Functional Analysis of Your Gut Microbiome

Vit B5 - Pantothenic acid



Report Descriptions

Vitamin B5 (pantothenic acid) is essential for energy production and fat metabolism. *Bacteroides fragilis*, *Prevotella copri*, *Ruminococcus* spp, *Salmonella enterica*, and *Helicobacter pylori* can all produce vitamin B5 in the gut. However, there are many species that rely on vitamin B5 for growth but cannot synthesize it, like most *Fusobacterium*, *Bifidobacterium* spp, *Faecalibacterium* spp, *Lactobacillus* spp, and some strains of *Clostridium difficile*, suggesting that these bacteria may compete with the host for vitamin B5.

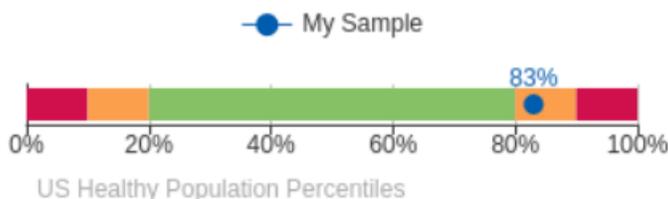
My Gut

10069

US Healthy Range

7372 to 13593

Vit B6 - Pyridoxine



Report Descriptions

Vitamin B6 (pyridoxine) is an incredibly versatile nutrient that supports immunity, brain function, and protein metabolism.

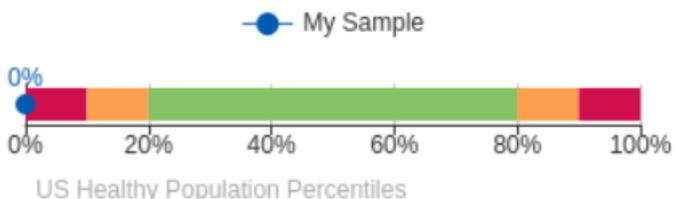
My Gut

6821

US Healthy Range

1051 to 8271

Vit B7 - Biotin



Report Descriptions

Biotin (also known as Vitamin H, Vitamin B7, or Vitamin B8) is a water soluble vitamin necessary for growth, development, and cellular energy production that can support healthy hair, skin, and nails and support healthy immune responses. Biotin is synthesized from tryptophan by intestinal bacteria like *Bacteroides fragilis*, *Prevotella copri*, *Ruminococcus lactaris*, *Clostridium difficile*, *Bifidobacterium infantis*, *Helicobacter pylori*, and *Fusobacterium varium*. In contrast, some species of *Prevotella*, *Bifidobacterium*, *Clostridium*, *Ruminococcus*, *Faecalibacterium*, and *Lactobacillus* may steal biotin from the host as they need it for survival.

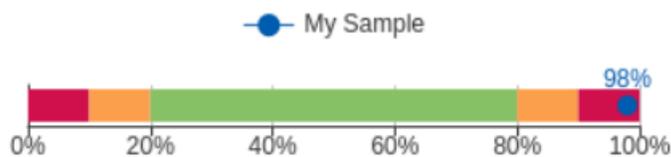
My Gut

Not Detected in my sample

US Healthy Range

291 to 1574

Vit B9 - Folate



US Healthy Population Percentiles

Report Descriptions

Vitamin B9 (folate/tetrahydrofolate), is essential for healthy blood cells. Gut-derived folate is directly absorbed into the colon, contributing up to 37% of the daily recommended intake. If vitamin B9 producers are low, there could be low levels of this nutrient available to the body.



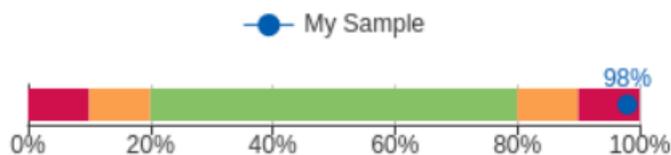
My Gut

US Healthy Range

9338

2499 to 5903

Vit B12 - Cobalamin



US Healthy Population Percentiles

Report Descriptions

Vitamin B12 (cobalamin) is crucial for healthy red blood cells, brain and nervous system function, DNA regulation, and metabolism. Production of B12 by gut bacteria contribute up to 31% of the daily recommended intake for this nutrient.



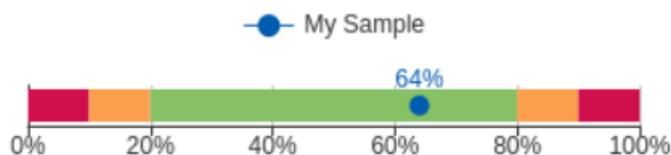
My Gut

US Healthy Range

9338

2499 to 5903

Vit K2 - Menaquinone



US Healthy Population Percentiles

Report Descriptions

Vitamin K2 is a fat soluble vitamin necessary for calcium metabolism and critical for the health of teeth, bones, nerves, and the cardiovascular system. Most K2 comes from dietary sources, however, gut bacteria including *Escherichia coli*, *Bacteroides vulgatus*, *Bacillus subtilis* and *Bacteroides fragilis* can also produce K2 endogenously. However, microbially derived K2 has protective role against oxidative tissue damage in the gut.



My Gut

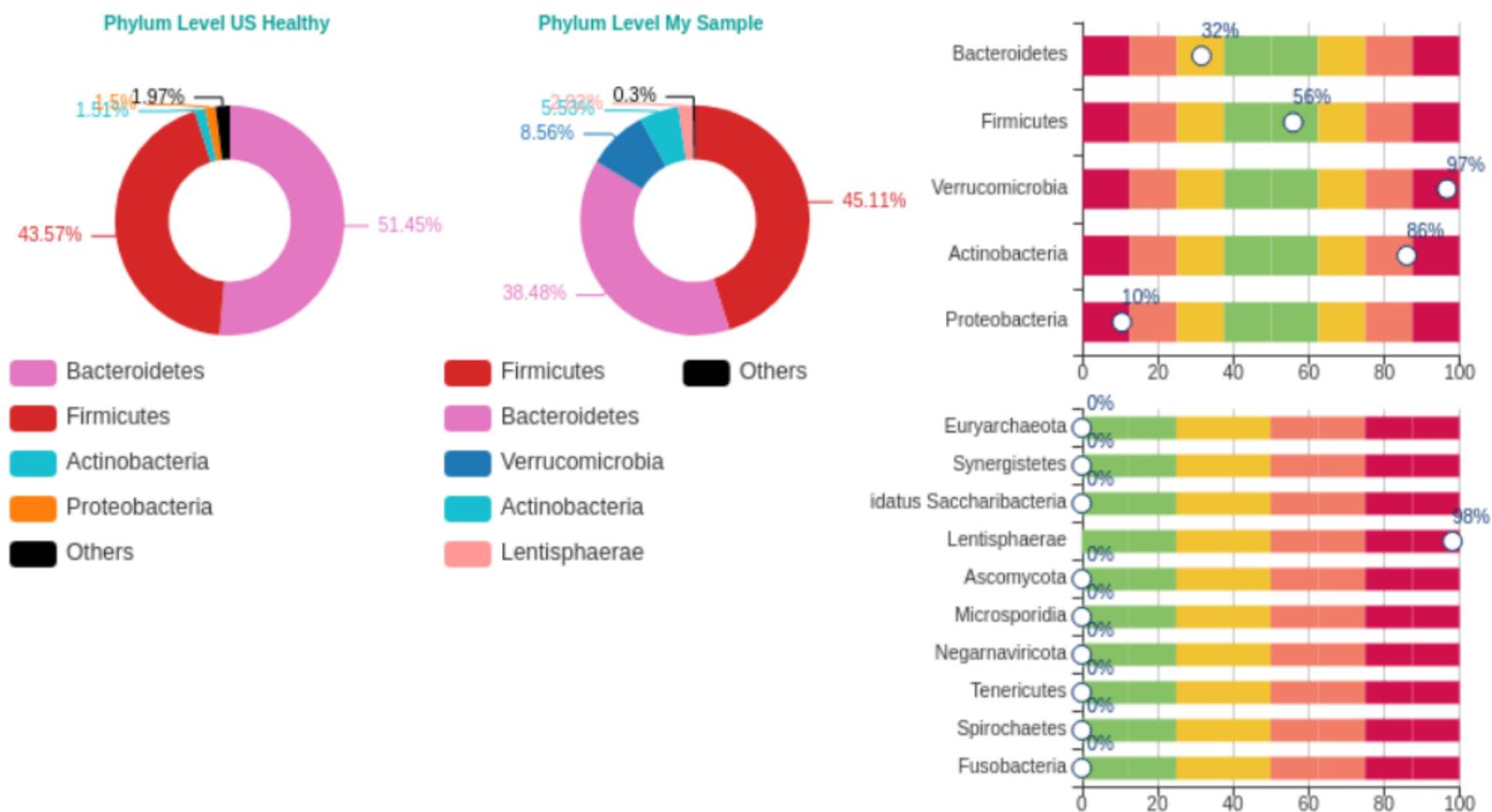
US Healthy Range

1034

361 to 1511

My Gut Microbiome Composition (Phylum level)

This section explores the composition of your gut microbiome at phylum level resolution.



The Donut Charts to the left visualizes the most abundant bacterial phyla in your gut. The Percentile Chart to the right compares the relative abundance (RA) for each bacterial phylum between your gut microbiome and the microbiomes typical for healthy populations. Percentile values between around 25% – 75% are typical, low values for a certain phylum suggest that in your case relative abundances are on the low side, high values suggest that your abundances are on the high side.

Phylum	Healthy Population Relative Abundance IQR Range [%]	My Sample Relative Abundance[%]	My Sample Percentile
Bacteroidetes	32.672 - 66.333	38.483	31.532
Firmicutes	27.629 - 57.18	45.111	55.856
Verrucomicrobia	0.22 - 1.794	8.564	96.68
Actinobacteria	0.25 - 3.382	5.527	86.001
Proteobacteria	0.731 - 2.739	0.299	10.403
Euryarchaeota	0.055 - 0.602	0	0
Synergistetes	0.022 - 0.277	0	0
Candidatus Saccharibacteria	0.007 - 0.016	0	0
Lentisphaerae	0.049 - 0.414	2.016	98.113
Ascomycota	0.001 - 0.002	0	0
Microsporidia	0.002 - 0.016	0	0
Negarnaviricota	0.001 - 0.006	0	0
Tenericutes	0.028 - 0.075	0	0
Spirochaetes	0.079 - 1.299	0	0
Fusobacteria	0.012 - 0.097	0	0

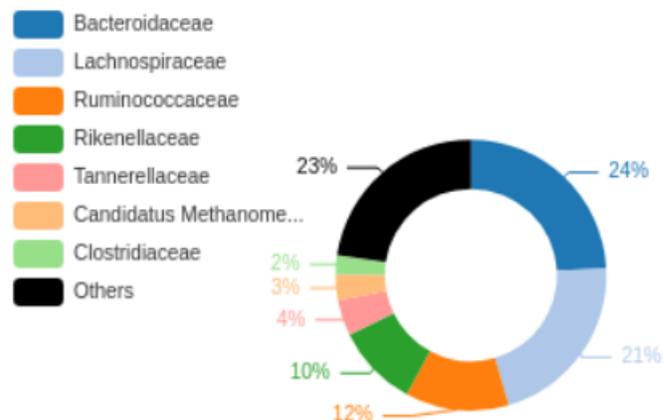
My Gut Microbiome Composition (Family level)

This section explores the composition of your gut microbiome at family level resolution.

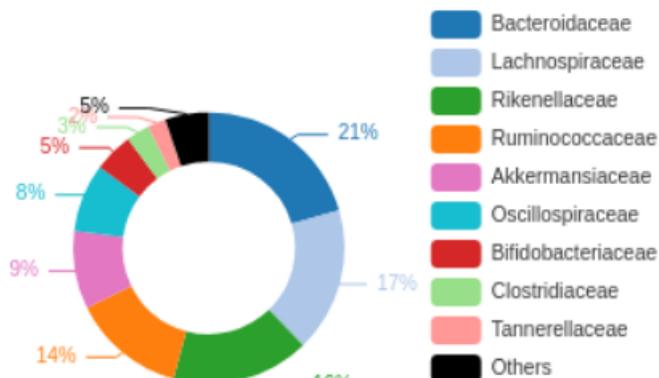
The **Donut Charts** visualize the most abundant bacterial genera in your gut.

The **Percentile Chart** below compares the relative abundance (RA) for each bacterial genera between your gut microbiome and the microbiomes typical for healthy populations. Percentile values between around 25% – 75% are typical, low values for a certain family suggest that in your case relative abundances are on the low side, high values suggest that your abundances are on the high side.

Family Level US Healthy



Family Level My Sample



Phylum	Family	Healthy Population Relative Abundance IQR Range [%]	My Sample Relative Abundance[%]	My Sample Percentile
Bacteroidetes	Bacteroidaceae	12.62 - 36.79	20.5	43.5
Firmicutes	Lachnospiraceae	11.77 - 30.91	17.26	42.21
Bacteroidetes	Rikenellaceae	5.37 - 14.22	16.44	82.22
Firmicutes	Ruminococcaceae	8.23 - 15.76	13.51	62.42
Verrucomicrobia	Akkermansiaceae	0.26 - 2.17	9.32	96.4
Firmicutes	Oscillospiraceae	0.8 - 2.07	8.14	99.87
Actinobacteria	Bifidobacteriaceae	0.31 - 3.3	4.67	82.49
Firmicutes	Clostridiaceae	1.4 - 3.14	2.79	66.84
Bacteroidetes	Tannerellaceae	2.23 - 6.37	2.18	24.43
Firmicutes	Acidaminococcaceae	0.52 - 1.99	0.67	29.81
Firmicutes	Eubacteriaceae	0.37 - 1.38	0.48	31.8
Firmicutes	Erysipelotrichacea...	0.14 - 1.51	0.17	28.86
Bacteroidetes	Odoribacteraceae	0.49 - 1.66	0	0
Bacteroidetes	Barnesiellaceae	0.69 - 2.31	0	0
Bacteroidetes	Porphyromonadaceae	0.09 - 1.28	0	0
Bacteroidetes	Prevotellaceae	0.06 - 5.91	0	0
Euryarchaeota	Candidatus Methano...	3.63 - 3.63	0	0
Firmicutes	Veillonellaceae	0.27 - 2.17	0	0
Proteobacteria	Sutterellaceae	0.3 - 1.68	0	0
Spirochaetes	Brachyspiraceae	0.08 - 1.4	0	0
Phylum unknown	Dysgonomonadaceae	1.15 - 4.8	0	0

Note: The organisms in the previous page typically occur in the healthy gut at Relative Abundances (RA) of > 0.5%. Organisms from the current page occur more rarely, but have been detected in your gut microbiome.

Phylum	Family	Healthy Population Relative Abundance IQR Range [%]	My Sample Relative Abundance[%]	My Sample Percentile
Actinobacteria	Eggerthellaceae	0.14 - 0.68	0.87	82.47
Phylum unknown	Victivallaceae	0.04 - 0.28	1.59	98.68



HOMEOSTASIS DIET

<https://calendly.com/semc3/microbiome>