

Dear Richard Sprague,

The optimal way to experience Viome is through our state-of-the-art, Al-driven app and our web interface. We are an exclusively online service, and our goal is to provide the benefit of ongoing results and recommendations, rather than information from one specific point in time.

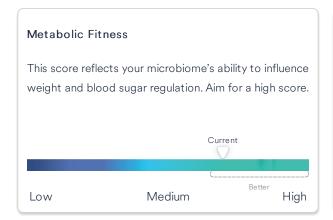
However, based on customer feedback, we've decided to offer a PDF version of your results and recommendations. We hope that this will enable you to share these valuable insights with your physician or loved ones.

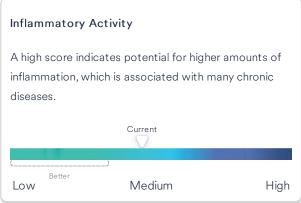
Viome recommendations are not evaluated or approved by FDA and are not required to be approved by FDA. The recommended supplements are intended to support general well-being and are not intended to treat, diagnose, mitigate, prevent, or cure any condition or disease. If conditions persist, please seek advice from your medical doctor.

Wellness Overview

Your microbiome is not in balance.

- You have relatively low beneficial microbial activity.
- You have relatively low unhealthy microbial activity.
- You have an average variety of microbial activities.
- Microbial butyrate pathway analysis indicates: butyrate production is high and potential to produce is low.

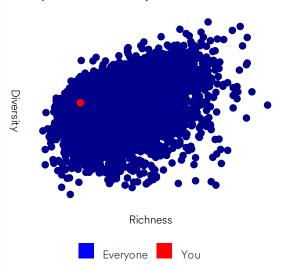




Diversity Metric



Compared to Everyone

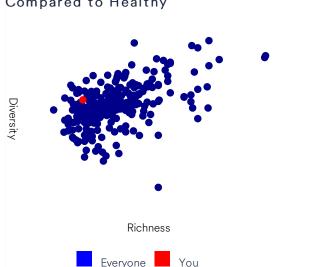


What does this mean? Richness is the total number of species of bacteria in your gut microbiome.

Diversity is the amount of individual bacteria from each of the bacterial species present in your gut microbiome.

This graph shows how you compare to the Everyone population in both the richness and diversity of your microbiome. A healthy microbiome is associated with a higher amount of both richness and diversity, and tends to be associated with overall wellness.

Compared to Healthy



What does this mean? Richness is the total number of species of bacteria in your gut microbiome.

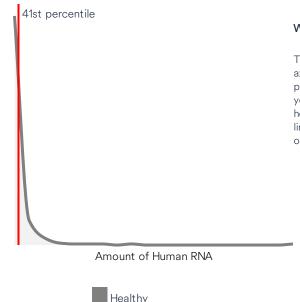
Diversity is the amount of individual bacteria from each of the bacterial species present in your gut microbiome.

This graph shows how you compare to the Healthy population in both the richness and diversity of your microbiome. A healthy microbiome is associated with a higher amount of both richness and diversity, and tends to be associated with overall wellness.

My RNA

01____

Compared to Healthy

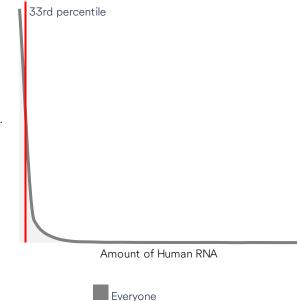


What does this mean?

The x-axis shows the proportion of human RNA in your samples, and the y-axis shows the number of people in the selected population with that proportion of human RNA. The red line shows the amount of human RNA in your sample. Typically, small amounts of human RNA are detected in healthy individuals. When an infection or inflammation occurs, the cells lining the gut are shed in higher amounts and can lead to a greater amount of human RNA in your stool.

02____

Compared to Everyone



What does this mean?

The x-axis shows the proportion of human RNA in your samples, and the y-axis shows the number of people in the selected population with that proportion of human RNA. The red line shows the amount of human RNA in your sample. Typically, small amounts of human RNA are detected in healthy individuals. When an infection or inflammation occurs, the cells lining the gut are shed in higher amounts and can lead to a greater amount of human RNA in your stool.

My Gut

Strain - Archaea

| Name | Activity | Everyone | Healthy |
|-----------------------------------------------------------|----------|----------|---------|
| Methano brevibacter smithii TS94C | 6.59% | 8.79% | 6.74% |
| Candidatus Methanomassiliicoccus intestinalis Issoire-Mx1 | 0.16% | 0.20% | 0.37% |

Species - Archaea

| I | Name | Activity | Everyone | Healthy |
|---|--------------------------------------------------|----------|----------|---------|
| ı | Methano brevibacter smithii | 6.77% | 11.04% | 9.14% |
| | Candidatus Methanomassiliicoccus intestinalis | 0.16% | 0.29% | 0.46% |

Genus - Archaea

| Name | Activity | Everyone | Healthy |
|-----------------------|----------|----------|---------|
| Methano brevibacter | 6.90% | 7.86% | 5.48% |
| Methanomassiliicoccus | 0.16% | 0.18% | 0.25% |

Phylum - Archaea

| Name | Activity | Everyone | Healthy |
|---------------|----------|----------|---------|
| Euryarchaeota | 1.20% | 21.32% | 24.44% |

Strain - Bacteria

| Name | Activity | Everyone | Healthy |
|-------------------------------------------------|----------|----------|---------|
| [Eubacterium] rectale strain T1-815 | 21.75% | 5.20% | 5.17% |
| [Eubacterium] hallii DSM 3353 | 19.63% | 7.34% | 8.44% |
| Parabacteroides distasonis str. 3999B T(B) 6 | 8.98% | 6.98% | 5.72% |
| Butyricimonas synergistica DSM 23225 | 7.95% | 4.41% | 3.43% |
| Acineto bacter baumannii 855125 | 4.28% | 7.26% | 5.82% |
| Parabacteroides merdae ATCC 43184 | 3.59% | 13.78% | 13.06% |
| Blautia obeum ATCC 29174 | 3.52% | 6.25% | 7.78% |
| Enterococcus faecium isolate Hp_74-d6 | 3.42% | 2.94% | 3.55% |
| [Eubacterium] eligens strain 2789STDY5834875 | 3.06% | 4.31% | 4.68% |
| Blautia massiliensis sp. GD8 | 1.62% | 0.67% | 0.67% |
| Synergistes sp. 3_1_syn1 | 1.38% | 0.55% | 0.48% |
| Bilophila wadsworthia ATCC 49260 | 1.20% | 3.15% | 2.61% |
| Ruminococcus bicirculans | 0.98% | 0.15% | 0.15% |
| Eggerthella lenta 1_1_60AFAA | 0.92% | 1.84% | 1.31% |
| | | | |

| Nama | A ativity | Everyene | Hoolthy. |
|--------------------------------------------------------------------------|-------------------|-------------------|----------|
| Name Clostridium sp. ATCC BAA-442 | Activity 0.92% | Everyone 0.15% | 0.36% |
| Clostridium baratii strain | 0.90% | 0.79% | 0.76% |
| 2789STDY5834907 Faecalibacterium prausnitzii A2-165 | 0.90% | 2.24% | 2.45% |
| · | | | |
| Acinetobacter baumannii 24975_5 | 0.89% | 15.88% | 13.59% |
| Clostridiales bacterium KLE1615 | 0.78% | 0.50% | 0.55% |
| Jonquetella anthropi E3_33 E1 | 0.74% | 0.37% | 0.17% |
| Anaerotruncus colihominis DSM 17241 | 0.61% | 0.85% | 0.74% |
| Oscillibacter sp. ER4 | 0.59% | 0.28% | 0.26% |
| Enorma massiliensis phl | 0.49% | 0.17% | 0.16% |
| Leuconostoc mesenteroides subsp. cremoris TIFN8 | 0.44% | 0.71% | 0.66% |
| Barnesiella intestinihominis YIT 11860 | 0.39% | 0.14% | 0.14% |
| Veillonella dispar ATCC 17748 | 0.34% | 1.60% | 0.98% |
| Faecalibacterium prausnitzii M21/2 | 0.23% | 0.40% | 0.45% |
| Eubacterium ramulus strain 2789STDY5608891 | 0.20% | 1.26% | 1.48% |
| Alistipes indistinctus YIT 12060 | 0.18% | 0.11% | 0.10% |
| Butyricimonas virosa DSM 23226 | 0.18% | 0.10% | 0.10% |
| Clostridium phoceensis strain GD3 | 0.16% | 0.26% | 0.22% |
| Turicibacter sanguinis strain 2789STDY5608865 | 0.15% | 0.19% | 0.09% |
| Alistipes obesi | 0.13% | 0.08% | 0.08% |
| Bacteroides massiliensis B84634 = Timor 84634 = DSM 17679 = JCM 13223 | ne 0.13% | 0.13% | 0.11% |
| Alistipes ihumii AP11 | 0.11% | 0.18% | 0.18% |
| Odoribacter laneus YIT 12061 | 0.11% | 0.11% | 0.11% |
| Intestinimonas massiliensis sp. GD2 | 0.10% | 0.08% | 0.09% |
| Lachnospiraceae bacterium TF01-11 | 0.09% | 0.25% | 0.20% |
| Sutterellaceae bacterium ND3 | 0.09% | 0.12% | 0.09% |
| Faecalibacterium prausnitzii strain 2789STDY5834930 | 0.09% | 0.10% | 0.08% |
| Coprococcus eutactus strain 2789STDY5608888 | 0.08% | 0.13% | 0.17% |
| Adlercreutzia equolifaciens DSM 19450 | 0.08% | 0.08% | 0.07% |
| Intestinimonas butyriciproducens strain 27 5-10 | 0.08% | 0.08% | 0.04% |
| Culturomica massiliensis strain Marseille- P2698 | 0.07% | 0.10% | 0.16% |
| Alistipes shahii WAL 8301 | 0.07% | 0.08% | 0.08% |
| Bacteroides cellulosilyticus DSM 14838 | 0.07% | 0.09% | 0.07% |
| | | | |

| Name | Activity | Everyone | Healthy |
|-----------------------------------------------|----------|----------|---------|
| Odoribacter splanchnicus DSM 20712 | 0.07% | 0.07% | 0.07% |
| Mageeibacillus indolicus UPII9-5 | 0.06% | 0.05% | |
| Gordonibacter pamelaeae 7-10-1-b | 0.06% | 0.05% | 0.05% |
| Parabacteroides sp. SN4 strain SN4, sp. SB4 | 0.05% | 0.08% | 0.09% |
| Bacteroides plebeius DSM 17135 | 0.05% | 0.10% | 0.09% |
| Emergencia timonensis strain SN18 | 0.05% | 0.05% | 0.04% |
| Variovorax paradoxus 110B | 0.05% | 0.04% | 0.04% |
| Alistipes finegoldii DSM 17242 | 0.05% | 0.07% | 0.07% |
| Anaerostipes hadrus strain 2789STDY5608830 | 0.04% | 0.06% | 0.06% |

Species - Bacteria

| N. | A | _ | |
|---------------------------------|----------|----------|---------|
| Name | Activity | Everyone | Healthy |
| [Eubacterium] rectale | 22.33% | 3.36% | 3.32% |
| [Eubacterium] hallii | 20.15% | 4.92% | 6.19% |
| Parabacteroides distasonis | 9.22% | 1.64% | 2.02% |
| Butyricimonas synergistica | 8.16% | 7.70% | 5.95% |
| Parabacteroides merdae | 3.69% | 5.84% | 6.02% |
| Blautia o beum | 3.62% | 4.49% | 6.66% |
| Acineto bacter baumannii | 3.51% | 19.49% | 16.21% |
| Enterococcus faecium | 3.51% | 2.05% | 2.35% |
| [Eubacterium] eligens | 3.14% | 4.36% | 4.74% |
| Blautia massiliensis | 1.66% | 1.20% | 1.25% |
| Synergistes sp. 3_1_syn1 | 1.42% | 0.86% | 0.76% |
| Bilophila wadsworthia | 1.23% | 4.60% | 4.02% |
| Ruminococcus bicirculans | 1.01% | 0.25% | 0.23% |
| Eggerthella lenta | 0.95% | 2.81% | 2.18% |
| Clostridium sp. ATCC BAA-442 | 0.95% | 0.23% | 0.48% |
| Clostridium baratii | 0.92% | 1.37% | 1.30% |
| Clostridiales bacterium KLE1615 | 0.80% | 0.84% | 0.94% |
| Jonquetella anthropi | 0.75% | 0.72% | 0.34% |
| Anaerotruncus colihominis | 0.63% | 1.25% | 1.09% |
| Oscillibacter sp. ER4 | 0.61% | 0.46% | 0.43% |
| Enorma massiliensis | 0.50% | 0.30% | 0.36% |
| Leuconostoc mesenteroides | 0.45% | 1.16% | 1.19% |
| Faecalibacterium prausnitzii | 0.43% | 1.69% | 1.68% |
| Barnesiella intestinihominis | 0.40% | 0.24% | 0.23% |
| Veillonella dispar | 0.35% | 2.57% | 1.75% |

| Name | Activity | Everyone | Healthy |
|-----------------------------------|----------|----------|---------|
| Eubacterium ramulus | 0.20% | 1.99% | 2.48% |
| Alistipes indistinctus | 0.19% | 0.19% | 0.17% |
| Butyricimonas virosa | 0.18% | 0.17% | 0.17% |
| Clostridium phoceensis | 0.16% | 0.43% | 0.36% |
| Turicibacter sanguinis | 0.15% | 4.78% | 4.47% |
| Alistipes obesi | 0.14% | 0.13% | 0.12% |
| Bacteroides massiliensis | 0.13% | 0.22% | 0.20% |
| Alistipes ihumii | 0.11% | 0.30% | 0.32% |
| Odoribacter laneus | 0.11% | 0.17% | 0.19% |
| Intestinimonas massiliensis | 0.11% | 0.13% | 0.14% |
| Lachnospiraceae bacterium TF01-11 | 0.10% | 0.44% | 0.34% |
| Sutterellaceae bacterium ND3 | 0.09% | 0.20% | 0.15% |
| Coprococcus eutactus | 0.09% | 1.27% | 2.25% |
| Adlercreutzia equolifaciens | 0.08% | 0.14% | 0.13% |
| Culturomica massiliensis | 0.07% | 0.15% | 0.22% |
| Alistipes shahii | 0.07% | 0.13% | 0.13% |
| Bacteroides cellulosilyticus | 0.07% | 0.15% | 0.13% |
| Odoribacter splanchnicus | 0.07% | 0.12% | 0.12% |
| Mageeibacillus indolicus | 0.06% | 0.07% | |
| Intestinimonas butyriciproducens | 0.06% | 0.13% | 0.10% |
| Gordonibacter pamelaeae | 0.06% | 0.10% | 0.09% |
| Parabacteroides sp. SN4 | 0.05% | 0.15% | 0.16% |
| Bacteroides plebeius | 0.05% | 0.18% | 0.15% |
| Emergencia timonensis | 0.05% | 0.08% | 0.08% |
| Variovorax paradoxus | 0.05% | 0.07% | 0.06% |
| Alistipes finegoldii | 0.05% | 0.13% | 0.12% |
| Anaerostipes hadrus | 0.05% | 0.13% | 0.13% |
| | | | |

Genus - Bacteria

| Name | Activity | Everyone | Healthy |
|----------------|----------|----------|---------|
| Parasutterella | 19.36% | 3.07% | 2.95% |
| Serratia | 11.79% | 12.23% | 12.48% |
| Romboutsia | 9.86% | 4.00% | 3.93% |
| Eubacterium | 8.41% | 2.54% | 3.59% |
| Akkermansia | 8.31% | 6.52% | 6.25% |
| Lactococcus | 5.23% | 16.08% | 15.89% |
| Turicibacter | 3.65% | 7.24% | 7.73% |
| Acinetobacter | 3.58% | 15.18% | 12.54% |
| Enterococcus | 3.58% | 1.43% | 1.80% |

| Name | Activity | Everyone | Healthy |
|------------------|----------|----------|---------|
| Blautia | 2.37% | 1.37% | 2.15% |
| Escherichia | 2.31% | 11.19% | 9.51% |
| Bifido bacterium | 2.18% | 8.33% | 5.98% |
| Synergistes | 1.32% | 0.84% | 0.52% |
| Bilophila | 1.25% | 2.53% | 2.30% |
| Ruminococcus | 1.03% | 0.24% | 0.23% |
| Eggerthella | 0.97% | 1.68% | 1.35% |
| Parabacteroides | 0.93% | 1.87% | 2.04% |
| Jonquetella | 0.77% | 3.42% | 0.13% |
| Butyricimonas | 0.75% | 0.20% | 0.25% |
| Anaerotruncus | 0.64% | 0.74% | 0.70% |
| Oscillibacter | 0.62% | 0.36% | 0.35% |
| Enorma | 0.51% | 0.24% | 0.30% |
| Leuconostoc | 0.46% | 0.97% | 1.05% |
| Faecalibacterium | 0.44% | 1.33% | 1.32% |
| Barnesiella | 0.41% | 0.18% | 0.18% |
| Veillonella | 0.36% | 1.81% | 1.14% |
| Morococcus | 0.30% | 1.42% | 1.51% |
| Tannerella | 0.27% | 0.75% | 0.69% |
| Clostridium | 0.24% | 0.67% | 0.80% |
| Alistipes | 0.13% | 0.13% | 0.13% |
| Bacteroides | 0.11% | 0.27% | 0.23% |
| Odoribacter | 0.11% | 0.09% | 0.09% |
| Intestinimo nas | 0.09% | 0.10% | 0.09% |
| Coprococcus | 0.09% | 0.49% | 0.94% |
| Adlercreutzia | 0.08% | 0.10% | 0.10% |
| Culturomica | 0.08% | 0.09% | 0.12% |
| Mageeibacillus | 0.06% | 0.05% | |
| Gordonibacter | 0.06% | 0.07% | 0.07% |
| Emergencia | 0.05% | 0.06% | 0.06% |
| Variovorax | 0.05% | 0.05% | 0.05% |
| Anaerostipes | 0.05% | 0.37% | 0.36% |
| | | | |

Phylum - Bacteria

| Name | Activity | Everyone | Healthy |
|-----------------|----------|----------|---------|
| Verrucomicrobia | 53.15% | 45.74% | 43.61% |
| Proteobacteria | 25.58% | 34.35% | 35.30% |
| Firmicutes | 8.88% | 9.68% | 10.85% |
| Synergistetes | 8.12% | 4.61% | 3.18% |
| | | | |

| Name | Activity | Everyone | Healthy |
|----------------|----------|----------|---------|
| Actinobacteria | 1.63% | 10.14% | 7.53% |
| Bacteroidetes | 0.81% | 3.97% | 3.67% |

Strain - Viruses

| Name | Activity | Everyone | Healthy |
|--------------------------------------------------------|----------|----------|---------|
| Beet cryptic virus 1 | 0.05% | 0.07% | 0.04% |
| Tropical soda apple mosaic virus isolate Okeechobee | 0.04% | 0.36% | 0.10% |

Species - Viruses

| Name | Activity | Everyone | Healthy |
|----------------------------------|----------|----------|---------|
| Beet cryptic virus 1 | 0.05% | 0.11% | 0.09% |
| Tropical soda apple mosaic virus | 0.04% | 0.60% | 0.21% |

Genus - Viruses

| Name | Activity | Everyone | Healthy |
|-------------------|----------|----------|---------|
| Alphapartitivirus | 0.06% | 0.09% | 0.05% |
| Tobamovirus | 0.04% | 0.90% | 0.26% |

Phylum - Viruses

| Name | Activity | Everyone | Healthy |
|---------------|----------|----------|---------|
| dsRNA viruses | 0.36% | 3.00% | 3.92% |
| ssRNA viruses | 0.26% | 4.05% | 3.87% |

Good Guys

Archaea - Species

| Name | Activity | Everyone | Healthy |
|-----------------------------|----------|----------|---------|
| Methano brevibacter smithii | 6.77% | 11.04% | 9.14% |

Bacteria - Species

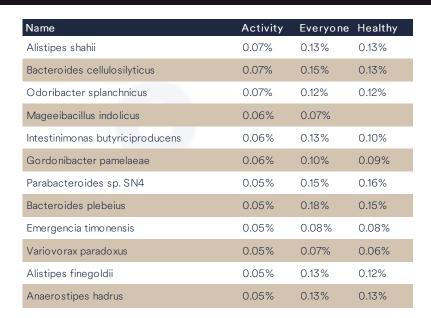
| Name | Activity | Everyone | Healthy |
|------------------------------|----------|----------|---------|
| Faecalibacterium prausnitzii | 0.43% | 1.69% | 1.68% |

Rare Bacteria

Bacteria - Species

| Name | Activity | Everyone | Healthy |
|-----------------------|----------|----------|---------|
| [Eubacterium] rectale | 22.33% | 3.36% | 3.32% |
| [Eubacterium] hallii | 20.15% | 4.92% | 6.19% |

| Name | Activity | Everyone | Healthy |
|-----------------------------------|----------|----------|---------|
| Parabacteroides distasonis | 9.22% | 1.64% | 2.02% |
| Butyricimonas synergistica | 8.16% | 7.70% | 5.95% |
| Parabacteroides merdae | 3.69% | 5.84% | 6.02% |
| Blautia obeum | 3.62% | 4.49% | 6.66% |
| Acinetobacter baumannii | 3.51% | 19.49% | 16.21% |
| Enterococcus faecium | 3.51% | 2.05% | 2.35% |
| [Eubacterium] eligens | 3.14% | 4.36% | 4.74% |
| Blautia massiliensis | 1.66% | 1.20% | 1.25% |
| Synergistes sp. 3_1_syn1 | 1.42% | 0.86% | 0.76% |
| Bilophila wadsworthia | 1.23% | 4.60% | 4.02% |
| Ruminococcus bicirculans | 1.01% | 0.25% | 0.23% |
| Eggerthella lenta | 0.95% | 2.81% | 2.18% |
| Clostridium sp. ATCC BAA-442 | 0.95% | 0.23% | 0.48% |
| Clostridium baratii | 0.92% | 1.37% | 1.30% |
| Clostridiales bacterium KLE1615 | 0.80% | 0.84% | 0.94% |
| Jonquetella anthropi | 0.75% | 0.72% | 0.34% |
| Anaerotruncus colihominis | 0.63% | 1.25% | 1.09% |
| Oscillibacter sp. ER4 | 0.61% | 0.46% | 0.43% |
| Enorma massiliensis | 0.50% | 0.30% | 0.36% |
| Leuconostoc mesenteroides | 0.45% | 1.16% | 1.19% |
| Faecalibacterium prausnitzii | 0.43% | 1.69% | 1.68% |
| Barnesiella intestinihominis | 0.40% | 0.24% | 0.23% |
| Veillonella dispar | 0.35% | 2.57% | 1.75% |
| Eubacterium ramulus | 0.20% | 1.99% | 2.48% |
| Alistipes indistinctus | 0.19% | 0.19% | 0.17% |
| Butyricimonas virosa | 0.18% | 0.17% | 0.17% |
| Clostridium phoceensis | 0.16% | 0.43% | 0.36% |
| Turicibacter sanguinis | 0.15% | 4.78% | 4.47% |
| Alistipes obesi | 0.14% | 0.13% | 0.12% |
| Bacteroides massiliensis | 0.13% | 0.22% | 0.20% |
| Alistipes ihumii | 0.11% | 0.30% | 0.32% |
| Odoribacter laneus | 0.11% | 0.17% | 0.19% |
| Intestinimo nas massiliensis | 0.11% | 0.13% | 0.14% |
| Lachnospiraceae bacterium TF01-11 | 0.10% | 0.44% | 0.34% |
| Sutterellaceae bacterium ND3 | 0.09% | 0.20% | 0.15% |
| Coprococcus eutactus | 0.09% | 1.27% | 2.25% |
| Adlercreutzia equolifaciens | 0.08% | 0.14% | 0.13% |
| Culturomica massiliensis | 0.07% | 0.15% | 0.22% |
| | | | |



Probiotics

Bacteria - Species

| Name | Activity | Everyone | Healthy |
|---------------------------|----------|----------|---------|
| Enterococcus faecium | 3.51% | 2.05% | 2.35% |
| Leuconostoc mesenteroides | 0.45% | 1.16% | 1.19% |

Recommendations

It's here! Your personalized Viome recommendations.

Your recommendations

Your personalized recommendations are based on the activity of microbes in your gut and the information you've provided. Your recommendations are aimed at balancing your overall microbiome. Let's put it this way: Your food list highlights foods that will be transformed by your microbes into beneficial substances while limiting foods that will be transformed into harmful metabolites.

Remember, you and your microbiome are unique, and no single recommendation applies to everyone. The same foods can be beneficial for one person, neutral for another, and harmful for others. Ready to dig in?

Your foods

Your food recommendations have been classified into 5 rankings to help you achieve optimum health and well-being. These are:

- 1. Superfoods. Meet your food destiny. These are your top 5 foods.
- 2. Indulge. After superfoods, these are the next most important foods to add to your diet.
- 3. Enjoy. Build a strong foundation with these foods.
- 4. Minimize. You can still eat these (but within limits).
- 5. Avoid. Nothing beats a helpful reminder. These foods are universally known to damage the gut microbiome.

Your recommended servings

We all struggle to figure out serving sizes on food labels because they only act as measurement tools. They are not

personalized for you.
With your food list, you get personalized servings to inform you on how much you should eat from each food category in a given day. And under each food, you'll find Viome's serving size, so you know the exact amount of that food to eat. Tip: If you are very active in a day, you can increase your servings from each food category proportionally for that day. Once you master your total servings per day, you can aim to achieve diversity by eating your recommended servings for each food rank.

Before you get started

Your success means a lot to us. Read our tips below before you begin.

What About Allergies?

You may notice some foods that you are allergic or sensitive to in your recommended food lists. Err on the side of caution. If you know you have a reaction or dislike to a recommended food, please do not consume it.

Foods are specifically chosen based on your unique microbiome rather than on allergies.

When is it best to eat?

Aim to eat 3 meals a day, and you may also need to snack in between meals.

Avoid eating 1 hour before you go to bed.

Go for variety

Explore foods that you haven't tried and since we're at it, alternate choices instead of eating the same food every day. Choose different foods from each of your indulge, enjoy, and minimize food categories based on your recommended amounts.

\'IOME

Listen to your body

Your recommended amounts are a guideline on the quantity of foods you should aim for. Stop eating once you are comfortably satiated or 80% full. Monitor how you feel, including your hunger, energy level, and mood or other forms of discomfort 1-3 hours after eating. If you consistently feel worse in any of these areas, you may need to adjust your food choices.

What else?

In addition to your food plan, your microbiome and your metabolism will gain an extra benefit from interval training at least 3 times per week.

Caloric restriction may provide more benefit than intermittent fasting.

Your Food List



Indulge

Eat some of these foods every day and vary which ones you eat so that you get a variety. Your Superfoods are foods from your indulge list that will be most beneficial for you. It is best to rotate your Indulge and Superfoods so you don't eat the same foods everyday. These should be your "go-to foods", as your microbes thrive on them. To help balance your body so it can function optimally, replace your Avoid foods with these Indulge foods.

Carbohydrates Grains Brown Rice Oats Vegetables Arugula Bok Choy Broccoli Capers Celery (organic) Cilantro Endive Garlic Jicama Kale Sauerkraut Seaweed Watercress **Fruits** Avocado Guava Papaya **Beverages** Green Tea (brewed) Extras Dill (fresh) Ginger Turmeric **Fats** Nuts and Seeds Brazil Nuts Filberts or Hazelnuts Pumpkin Seeds Oils Flax Oil Olive Oil **Proteins** Meats Bone Broth (mammal) Chicken (dark) Lamb Turkey (white) Seafood Salmon, Pacific (wild-caught) Legumes Black Beans Other Proteins Egg (large) Your microbiome does well with these foods so they should form the foundation of your Enjoy diet. Eat at least 4 servings of your Enjoy foods each day, but remember to rotate them so you don't eat the same foods everyday. Carbohydrates Grains Amaranth Buckwheat Bulgar Quinoa Rye (sprouted bread) Wheat (sprouted bread) Wild Rice Vegetables Alfalfa Sprouts Artichoke Asparagus Bamboo Shoots Basil Bean Sprouts Beet Greens

Bell Pepper (organic) Brussels Sprouts Burdock Root Cabbage Cardoon (thistle stem) Carrot Cauliflower Celeriac Chard Chicory (root) Collard Greens Cucumber Dandelion Greens Eggplant Escarole Fennel Bulb Gourd Grape Leaves Green Beans Hot Pepper (organic) Jerusalem Artichoke Kimchi Kohlrabi Leek Lettuce Mushrooms Mustard Greens Okra Onion Parsley Peas Pickle (unsweetened) Potato (small, organic) Pumpkin Radicchio Radish Rhubarb Snap Peas | Spinach (organic) | Sprouted Radish Seeds | Summer Squash | Taro | Tomato (organic) Water Chestnuts Wheatgrass Zucchini Squash **Fruits** Apple (medium, organic) Apricot Blackberry Blueberry Boysenberry Breadfruit Cranberry Currant Elderberry Fig Goji Berry Gooseberry Grapefruit Grapes (organic) Huckleberry Jackfruit Kiwi Kumquat Lemon Lime Loganberries Lychee Marionberry Melon Mulberries Olives Orange Passionfruit Pear (organic) Persimmon Plantain Plum Pomegranate Pummelo Raspberry Salmonberry Sour Cherries Star Fruit Strawberry (organic) **Beverages** Almond Milk (unsweetened) Black Tea (brewed) Coconut Milk (unsweetened) Coconut Water Coffee (brewed, organic) Herbal Tea (brewed) Soy Milk (unsweetened) White Tea (brewed) Sweeteners Cocoa (unsweetened) Extras Allspice Bay Leaf Black Pepper Caraway Seed Cardamom Carob Cayenne Pepper Celery Seed Chervil Chili Powder Cinnamon Cloves Coriander Cumin Fennel Seed Fenugreek Seed Horseradish Lotus Seeds Mace Marjoram Mustard Seed Nutmeg Oregano Paprika Peppermint (fresh) Poppy Seed Rosemary (fresh) Saffron Sage Savoury Spearmint (fresh) Spirulina Tarragon Vanilla Extract Vinegar Vinegar Apple Cider **Fats Nuts and Seeds** Almonds Cashews Chestnuts Chia Seeds Coconut Meat Flax Seeds Hemp Hearts Hickory Nuts Macadamia Nuts Pecans Pine Nuts Pistachios Sesame Seeds Sunflower Seeds Walnuts Oils Avocado Oil Butter | Coconut MCT Oil | Coconut Oil | Ghee | Grape Seed Oil | Safflower Oil **Proteins** Meats Beef (fatty, grass-fed) Beef (lean, grass-fed) Bone Broth (poultry) Buffalo Chicken (white) Cornish Game Hen Duck Emu Game Meat (venison, elk) Goat Goose Ostrich Pheasant Quail Turkey (dark) Veal Seafood Abalone Anchovy Bone Broth (fish) Catfish Caviar or Roe Cod, Alaskan Crayfish Dungeness Crab, Pacific Eel Haddock Halibut, Pacific Herring Lobster Mackerel Mussel Perch Rainbow Trout Sardine Scallops Shellfish Clam Shellfish Oyster Shrimp (domestic) Squid Tuna (pole caught) Turbot Legumes Chickpeas Fava Beans Miso Natto Peanuts Red Beans Soybeans (non-GMO) White Beans

Y I O M E

Dairy Cheese Goat Cheese Goat Milk Heavy Cream (33% fat) Kefir Ricotta or Cottage Cheese (2% fat)

Sheep Cheese Sheep Milk Whole Milk Yogurt (plain)

Other Proteins Egg White Egg Yolk Tempeh Tofu

Minimize

Minimize foods can be eaten once in a while, but they will not promote optimal wellness. Try to limit your intake of each Minimize food to a maximum of 3 servings per week.

Carbohydrates

Grains Barley Couscous Kamut Millet Triticale White Rice

Vegetables Cassava Corn Tortilla (organic, non-GMO) Hard Squash Parsnip Rutabaga Turnip

Yam or Sweet Potato

Fruits Banana (small) Cherry (organic) Dates Mango Mangosteen Nectarine (organic) Peach Pineapple

Prunes

Beverages Rice Milk

Salt Sea Salt or Himalayan Salt

Sweeteners Cane Sugar Honey Maple Syrup Molasses Stevia

Extras Rice Noodles

Proteins

Meats Pork (lean)

Legumes Adzuki Beans Black Eyed Peas Lentils Lima Beans Pinto Beans

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Avoid

Stay away from these foods. Avoid foods are harmful to your overall wellness and should be removed from your diet altogether. Your microbes will thank you for it.

Carbohydrates

Fruits Raisins

Sweeteners Agave Nectar Beet Sugar Dextrose Maltose

French Fries Fruit Juices Granola Bars Hydrogenated Vegetable Oil lodized Salt Oatmeal (flavored)

Processed Cheese Processed Meat Rice Cakes (flavored) Saccharin Soda (regular or diet)

Sucralose Sugar (white) White Flour Yogurt (flavored)

\'IOME

Fats

Oils

Canola Oil Lard Margarine Shortening

Supplements

Supplement recommendations are designed to promote healthy activities of your gut microbiome. These supplements are available for purchase through online retailers.



ABx Support

Probiotics by Klaire Labs

1 capsule 2x daily



PhytoGanix

Polyphenols by Metagenics

1 tsp daily



Cal-Mag Butyrate

Gut Support by BodyBio

1 capsule 3x daily for 90 days



PaleoFiber

Prebiotic by Designs for Health

Build up to 2 tsps daily

Viome recommendations are not evaluated or approved by FDA and are not required to be approved by FDA. The recommended supplements are intended to support general wellbeing and are not intended to treat, diagnose, mitigate, prevent, or cure any condition or disease. Please seek advice from your medical doctor and check all ingredients for a known allergy or sensitivity prior to taking.