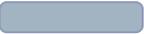


Microbiological test report for :

Full gut microbiome profiling

Shotgun Metagenomic Sequencing - 20M reads - PE150

Sample ID: RG-XXXX-XXX

Improvable : 

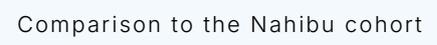
Correct : 

Optimal : 

Sample position : 

This report presents your results, comparing them with those obtained from the Nahibu cohort.

MAIN CHARACTERISTICS

| Indicator | Result | Comparison to the Nahibu cohort |
|--------------------|-------------|---|
| Enterotype | Bacteroides |  |
| Microbial richness | 124 |  |
| Balance | Unbalanced |  |

BACTERIA OF INTEREST

Description of these bacteria is provided in the appendix.

POSITIVE IMPACT BACTERIA

| Indicator | Result | Comparison to the Nahibu cohort |
|-------------------------------------|---------------------------|---|
| <i>Faecalibacterium prausnitzii</i> | Correct |  |
| <i>Eubacterium hallii</i> | Improvable (Not detected) |  |
| <i>Bifidobacterium longum</i> | Improvable (Not detected) |  |
| <i>Roseburia intestinalis</i> | Correct |  |
| <i>Akkermansia muciniphila</i> | Correct |  |
| <i>Veillonella atypica</i> | Correct (Not detected) |  |

NEGATIVE IMPACT BACTERIA

| Indicator | Result | Comparison to the Nahibu cohort |
|---------------------------------|------------------------|---|
| <i>Bilophila wadsworthia</i> | Correct |  |
| <i>Clostridioides difficile</i> | Optimal (Not detected) |  |
| <i>Ruminococcus gnavus</i> | Optimal (Not detected) |  |



FUNCTIONAL POTENTIAL

Description of each compound is provided in the appendix.

DIGESTION AND INTESTINAL COMFORT

| Indicator | Result | Comparison to the Nahibu cohort |
|---|------------|---|
| BLOATING AND GAS | | |
| Hydrogen sulfide | Optimal |  |
| ACIDS AND REFLUX | | |
| Lactate | Improvable |  |
| SATIETY | | |
| Acetate, lactate, propionate | Improvable |  |
| FATTY ACID METABOLISM REGULATION | | |
| Spermidine | Optimal |  |
| VITAMIN PRODUCTION | | |
| Adénosylcobalamine | Improvable |  |
| Menaquinone | Improvable |  |

IMMUNITY

| Indicator | Result | Comparison to the Nahibu cohort |
|---|------------|---|
| DEVELOPMENT AND MAINTENANCE OF GUT TISSUES | | |
| Butyrate | Improvable |  |
| Hydrogen sulfide | Optimal |  |
| Putrescine, spermidine | Correct |  |
| INFLAMMATION | | |
| Butyrate | Improvable |  |
| Acetate | Improvable |  |
| Lactate | Improvable |  |
| Histidine | Improvable |  |



PHYSICAL ABILITIES

| Indicator | Result | Comparison to the Nahibu cohort |
|--------------|------------|---|
| Acetate | Improvable |  |
| Histidine | Improvable |  |
| Pantothenate | Optimal |  |

NEUROPSYCHOLOGICAL ABILITIES

| Indicator | Result | Comparison to the Nahibu cohort |
|-----------|--------|---------------------------------|
|-----------|--------|---------------------------------|

GENERAL BRAIN FUNCTION

| | | |
|-----------|------------|---|
| Histidine | Improvable |  |
|-----------|------------|---|

COGNITIVE ABILITIES AND MEMORY

| | | |
|------------------|------------|---|
| Tetrahydrofolate | Optimal |  |
| Butyrate | Improvable |  |

SLEEP

| | | |
|------------|------------|---|
| Butyrate | Improvable |  |
| GABA | Optimal |  |
| Tryptophan | Optimal |  |

DEPRESSION AND MOOD DISORDERS

| | | |
|----------|---------|---|
| GABA | Optimal |  |
| Tyrosine | Optimal |  |

STRESS AND ANXIETY

| | | |
|--------------|---------|---|
| Pantothenate | Optimal |  |
| GABA | Optimal |  |
| Tryptophan | Optimal |  |
| Tyrosine | Optimal |  |



NUTRITIONAL ADVICE

Generic Advice

Fiber plays a key role in ensuring proper intestinal transit, by facilitating digestive movements and supplying nutrients to the microbiota. They are essential to the overall balance of the intestinal microbiota, contributing to its health benefits.

FODMAPs are sugars naturally present in our diet. They are barely digested before reaching the colon, where bacteria ferment them very quickly. Some people may be sensitive to FODMAPs (e.g. FOS and GOS), sometimes causing bloating and digestive discomfort. It is then recommended to introduce these foods gradually to help restore the balance of the microbiota.

For a balanced microbiota, it's advisable to eat a varied diet and limit processed foods rich in fat or salt. Adequate hydration, a high-fiber diet and regular physical activity remain the basic tips for maintaining a healthy intestinal balance.

Recommendations adapted to your microbiotic data

| Indicator | Recommended foods or supplements |
|--|---|
| Development and maintenance of gut tissues | Almond • Aloe • Asparagus • Avocado • Banana • Basil • Blackcurrant • Broccoli • Cashew nut • Cauliflower • Cheese • Chestnut • Chickpea • Chinese cabbage • Cinnamon • Coconut • Corn • Dried bean • Fava bean • Fresh bean (green, wax) • Garlic • Hazelnut • Kale • Lemon • Millet • Mustard • Oat • Oyster • Passion fruit • Pecan nut • Pistachio • Potato • Prickly pear • Quince • Red cabbage • Sesame • Shallot • Sorrel • Squash • Sunflower seed • Swiss chard • Wheat germ • White cabbage • Whole wheat • Yogurt • Zucchini |
| Inflammation | Acai • Acerola • Almond • Aloe • Amla • Apricot • Aronia • Artichoke • Asparagus • Astaxanthin • Avocado • Baobab • Basil • Bell pepper • Black pepper • Blackberry • Blackcurrant • Brussels cabbage • Butternut squash • Camu-camu • Chamomile • Cheese • Cherry • Chlorella • Cinnamon • Cocoa • Cranberry • Cucumber • Dandelion • Dill • Eggplant • Fennel • Fermented cabbage • Garlic • Ginger • Goji berry • Grapefruit • Hemp seed • Honey • Kale • Lemon • Lemongrass • Licorice • Moringa • Oat • Olive • Olive oil • Physalis • Pineapple • Pomegranate • Rapeseed oil • Red cabbage • Royal jelly • Sage • Sea buckthorn berry • Sesame oil • Sorrel • Soy • Spinach • Squash • Starfruit • Turmeric • Violet • Watermelon • Wheatgrass • Yogurt |
| Bloating and gas | Acai • Aloe • Artichoke • Blueberry • Brown rice • Celery • Chamomile • Coriander • Dill • Eggplant • Fennel • Ginger • Grape • Kiwi • Lemon • Lemongrass • Mandarin • Milk kefir • Mint • Orange • Papaya • Pineapple • Quince • Quinoa • Spinach • Squash • Sweet potato • Thyme • Tomato • Turmeric • Water kefir • Yogurt |
| Acids and reflux | Almond • Aloe • Amla • Apple • Asparagus • Avocado • Banana • Basil • Broccoli • Brown rice • Cauliflower • Celery • Chamomile • Chia seed • Cucumber • Egg yolk • Eggplant • Flaxseed • Fresh bean (green, wax) • Ginger • Lemon • Melon • Oat • Olive oil • Pear • Sesame oil • Sunflower oil • Turnip • Walnut • Wheatgrass |

This analysis report is provided for informational purposes only and is not intended to diagnose, treat, cure, or prevent any disease. The results presented are based on data related to microbiome diversity and trends and should be interpreted as part of a comprehensive health assessment by a qualified professional. The information contained in this report should not be considered as medical advice or treatment recommendations. It is recommended to consult a doctor or healthcare professional before making decisions based on these results.

| Indicator | Recommended foods or supplements |
|----------------------------------|--|
| Satiety | Acai • Acerola • Agar-agar • Almond • Aloe • Apple • Aronia • Artichoke • Asparagus • Avocado • Banana • Baobab • Barley • Beetroot • Blackcurrant • Broccoli • Brown rice • Brussels cabbage • Carrot • Cauliflower • Celery • Cheese • Chestnut • Chia seed • Chickpea • Chicory • Chili pepper • Cocoa • Corn • Date • Egg white • Egg yolk • Eggplant • Endive • Flaxseed • Fresh bean (green, wax) • Goji berry • Hemp seed • Konjac • Leek • Lemon • Lentils • Lettuce • Lucuma • Lytchee • Mandarin • Mango • Melon • Mushroom • Nectarine • Oat • Oily fish • Olive • Onion • Orange • Papaya • Parsnip • Passion fruit • Pea • Peanut butter • Peanut • Persimmon • Quince • Quinoa • Radish • Raspberry • Red cabbage • Red meat • Redcurrant • Seafood • Spinach • Squash • Sweet potato • Swiss chard • Teff • Tomato • Walnut • Wheatgrass • White fish • White rice • Whole grain • Zucchini |
| Fatty acid metabolism regulation | Almond • Aloe • Amla • Apple • Apricot • Aronia • Artichoke • Astaxanthin • Broccoli • Carrot • Chestnut • Chia seed • Eggplant • Garlic • Goji berry • Grape • Hazelnut • Hemp seed • Konjac • Milk kefir • Mint • Oat • Pecan nut • Psyllium • Quince • Rhubarb • Sea buckthorn berry • Soy • Spirulina • Sunflower seed • Walnut • Wheat germ • Wheatgrass • Whole grain |
| Vitamin production | Acai • Acerola • Almond • Aloe • Amla • Apricot • Aronia • Artichoke • Asparagus • Avocado • Basil • Beetroot • Bell pepper • Blackberry • Blackcurrant • Blueberry • Broccoli • Brussels cabbage • Butter • Camu-camu • Carrot • Cauliflower • Chestnut • Chickpea • Chicory • Chili pepper • Chinese cabbage • Coriander • Corn • Cranberry • Dandelion • Date • Egg white • Egg yolk • Eggplant • Endive • Fava bean • Fermented cabbage • Flaxseed • Fresh bean (green, wax) • Garlic • Goji berry • Hemp seed • Kale • Kiwi • Lemon • Lentils • Lettuce • Lucuma • Mandarin • Milk kefir • Mushroom • Mustard • Oily fish • Olive • Orange • Oyster • Papaya • Parsley • Parsnip • Passion fruit • Pea • Peanut butter • Pear • Pineapple • Potato • Prickly pear • Rapeseed oil • Red cabbage • Redcurrant • Sea buckthorn berry • Sesame • Spinach • Squash • Starfruit • Sunflower seed • Swiss chard • Turnip • Watermelon • Wheatgrass • White cabbage • Whole wheat • Yogurt • Zucchini |
| Physical abilities | Acai • Acerola • Agave syrup • Alfalfa • Almond • Apple • Apricot • Aronia • Artichoke • Asparagus • Astaxanthin • Avocado • Banana • Baobab • Basil • Beetroot • Blackcurrant • Brown rice • Brussels cabbage • Butternut squash • Cauliflower • Cheese • Cherry • Chestnut • Chia seed • Chickpea • Chili pepper • Chinese cabbage • Cocoa • Coffee • Coriander • Cucumber • Date • Dried bean • Durian • Fava bean • Fermented cabbage • Fresh bean (green, wax) • Garlic • Gelatin • Goji berry • Green tea • Guarana seed • Guava • Hazelnut • Hemp seed • Kale • Lemon • Lemongrass • Lentils • Liver • Lucuma • Lytchee • Mandarin • Milk • Moringa • Mustard • Oat • Oily fish • Olive oil • Orange • Parsnip • Passionflower • Pea • Peanut butter • Peanut • Pineapple • Quince • Quinoa • Rapeseed oil • Raspberry • Red cabbage • Red meat • Redcurrant • Royal jelly • Seafood • Sesame • Shallot • Sorrel • Soy • Squash • Starfruit • Sunflower seed • Sweet potato • Swiss chard • Teff • Turmeric • Walnut • Watermelon • Wheat germ • White cabbage • White fish • White meat • Whole grain • Whole wheat • Yogurt |



| Indicator | Recommended foods or supplements |
|--------------------------------|--|
| General brain function | Almond • Amla • Apple • Aronia • Artichoke • Asparagus • Astaxanthin • Avocado • Basil • Beetroot • Bell pepper • Blackberry • Blueberry • Broccoli • Carrot • Celery • Chia seed • Chickpea • Chicory • Cocoa • Coconut oil • Coconut • Cranberry • Dried bean • Egg yolk • Eggplant • Endive • Fermented cabbage • Flaxseed • Fresh bean (green, wax) • Garlic • Goji berry • Grape • Grapefruit • Green tea • Hazelnut • Kale • Lemon • Lemongrass • Lentils • Lettuce • Liver • Mandarin • Maple syrup • Moringa • Oat • Oily fish • Olive • Olive oil • Orange • Pea • Peanut • Pomegranate • Potato • Rapeseed oil • Raspberry • Rosemary • Royal jelly • Saffron • Sea buckthorn berry • Seafood • Sesame • Sesame oil • Shallot • Spinach • Squash • Strawberry • Sweet potato • Teff • Tomato • Turmeric • Walnut • Wheat germ • Whole grain • Whole wheat • Zucchini |
| Cognitive abilities and memory | Almond • Aloe • Amla • Apple • Apricot • Aronia • Artichoke • Asparagus • Astaxanthin • Avocado • Banana • Baobab • Basil • Blackberry • Blueberry • Broccoli • Carrot • Cashew nut • Celery • Chicory • Cinnamon • Dried bean • Egg yolk • Eggplant • Fermented cabbage • Garlic • Gelatin • Grape • Green tea • Guarana seed • Hazelnut • Honey • Lemon • Moringa • Oat • Oily fish • Olive • Olive oil • Pistachio • Radish • Rosemary • Royal jelly • Sage • Spinach • Strawberry • Thyme • Turmeric • Walnut • Wheat germ • Wheatgrass |
| Sleep | Almond • Apple • Apricot • Asparagus • Avocado • Banana • Basil • Black pepper • Chamomile • Cherry • Chia seed • Dill • Dried bean • Fermented cabbage • Garlic • Gelatin • Honey • Lemongrass • Onion • Passionflower • Pineapple • Saffron • Valerian |
| Depression and mood disorders | Avocado • Banana • Black pepper • Blackcurrant • Camu-camu • Eggplant • Fermented cabbage • Garlic • Oily fish • Olive oil • Oyster • Saffron • Walnut |
| Stress, and anxiety | Amla • Apple • Asparagus • Avocado • Banana • Basil • Blackcurrant • Chamomile • Cocoa • Cucumber • Eggplant • Fermented cabbage • Garlic • Goji berry • Lemongrass • Passionflower • Saffron • Valerian • Wheat germ |

| Indicator | Foods or supplements to limit |
|--|--|
| Development and maintenance of gut tissues | Chili pepper |
| Inflammation | Red meat |
| Bloating and gas | Broccoli • Brussels cabbage • Cauliflower • Cherry • Chickpea • Chinese cabbage • Fava bean • Fermented cabbage • Kale • Milk • Onion • Red cabbage • Watermelon • White cabbage |
| Acids and reflux | Aronia • Baobab • Chili pepper • Cocoa • Coffee • Dandelion • Egg white • Garlic • Grape • Mandarin • Mint • Onion • Orange • Pineapple • Tomato |
| Satiety | Butter • Cashew nut • Grape • Hazelnut • Honey • Olive oil |



| Indicator | Foods or supplements to limit |
|----------------------------------|--|
| Fatty acid metabolism regulation | Cold cuts • Liver |
| Vitamin production | Konjac |
| Sleep | Baobab • Camu-camu • Cocoa • Coffee • Eggplant • Goji berry • Rosemary • Sea buckthorn berry • Spirulina • Thyme |
| Depression and mood disorders | Bitter orange • Passionflower |
| Stress, and anxiety | Bitter orange |



APPENDICES

BACTERIA OF INTEREST

- *Faecalibacterium prausnitzii*

F. prausnitzii is one of the most abundant species in the intestinal microbiota. A butyrate producer, an anti-inflammatory compound, it contributes to good intestinal health by nourishing colon cells and reinforcing the intestinal barrier.

- *Eubacterium halii*

E. halii is a butyrate- and propionate-producing species, two compounds with health benefits.

- *Bifidobacterium longum*

B. longum is a commercially available probiotic. It has anti-inflammatory effects and appears to improve symptoms in people suffering from constipation, celiac disease or ulcerative colitis.

- *Roseburia intestinalis*

R. intestinalis is one of the dominant species in the gut microbiota. As a butyrate producer, it contributes to the proper functioning of the intestinal barrier and has anti-inflammatory properties.

- *Akkermansia muciniphila*

A. muciniphila helps strengthen the intestinal barrier, supports weight management, has anti-inflammatory properties, and is associated with overall good health.

- *Veillonella atypica*

V. atypica seems to promote the growth of many other microorganisms, including beneficial bacteria for the host. It also helps to eliminate potentially harmful compounds produced by bacteria living in the digestive system. Mainly found in the microbiota of athletes, it appears to enhance physical performance by transforming the lactic acid produced during exercise.

- *Bilophila wadsworthia*

B. wadsworthia has pro-inflammatory properties and appears to worsen metabolic disorders in people with a high-fat diet. Its abundance may be increased in high saturated fat diets, or as part of a low FODMAP diet.

- *Clostridioides difficile*

C. difficile is a pathogenic bacterium responsible for the majority of nosocomial infections worldwide. Its effects range from mild diarrhea to severe intestinal pathologies.

- *Ruminococcus gnavus*

R. gnavus is a commensal (common) species of the intestine. It has a high inflammatory potential.



COMPOUNDS OF INTEREST

- **Acetate**

This volatile fatty acid, produced by bacteria and transported to organs by the blood, is a prime source of energy for muscles. Also involved in immune cell expression, its concentration increases during infection to support the adaptive immune response.

- **Acetate, Lactate, Propionate**

These volatile fatty acids are involved in satiety regulation by inhibiting hormone secretion from colon mucosal cells and promoting the release of peptides and hormones that act on the central nervous system. High concentrations lead to reduced appetite.

- **Adénosylcobalamine**

Adenosylcobalamin is one of the active forms of vitamin B12. It comes mainly from the diet, but can be produced by certain bacteria, reducing the risk of deficiency. It is involved in the synthesis of essential amino acids for both the microbiota and the host. Additionally, it plays a role in regulating fatty acid metabolism and cell development.

- **Butyrate**

Short-chain fatty acid produced by intestinal microbiota during fiber fermentation. As the main nutrient for intestinal mucosa cells, it stimulates mucus production, helping to maintain digestive tract tissue. Butyrate also has anti-inflammatory properties and promotes sleep onset and deep sleep phases.

- **GABA**

GABA (gamma-aminobutyric acid) inhibits neurons to prevent constant excitation. It must be balanced with glutamate to reduce the risk of epilepsy.

- **Histidine**

An essential compound in hemoglobin synthesis, it plays a crucial role in arterial health. It also contributes to neurological well-being by protecting nerve cells. Sufficient histidine levels help reduce mental fatigue, support memory, and promote quality sleep. However, an excess of histidine can lead to stress and anxiety.

- **Lactate**

The acidity from lactate inhibits the growth of potentially pathogenic bacteria, but maintaining balance is essential to prevent excessive acidification of the intestine. Its presence is directly linked to increased local acidity through a higher concentration of lactate-producing bacteria and a decrease in lactate-utilizing bacteria.

- **Menaquinone**

Menaquinone is the form of vitamin K produced by bacteria. It is involved in cardiovascular and bone metabolism, cell growth and immunity. Vitamin K also promotes the development of beneficial bacteria and limits the development of pathogens.

- **Pantothenate**

Pantothenate, or vitamin B5, is involved in the regulatory mechanisms of adrenalin, insulin and porphyrin (hemoglobin precursor). It has anti-stress properties and helps reduce fatigue.



- **Putrescine, spermidine**

Putrescine and spermidine are polyamines found in the intestine, part of which is produced by the intestinal microbiota. They help regulate cell growth and support intestinal maturation. Some of these polyamines are used by bacteria to form biofilms, which play a protective role in the intestinal lining. However, excessive concentrations may be linked to an increased risk of cancer and chronic inflammation.

- **Spermidine**

Spermidine is a polyamine produced by the gut microbiota. It helps reduce adiposity and fat accumulation in the liver. Spermidine thus plays a role in preventing diet-induced obesity.

- **Hydrogen sulfide**

At low concentrations, hydrogen sulfide reduces inflammation in the digestive tract. It stabilizes mucus and bacterial populations, reinforcing the intestinal microbiota's barrier effect. However, excessive bacterial production of hydrogen sulfide can lead to intestinal irritation and inflammation. When absorbed by the body, this gas may be linked to hypertension issues.

- **Tetrahydrofolate**

Tetrahydrofolate is the active form of vitamin B9 and can be synthesized by intestinal bacteria. It is involved in the development of the nervous system and the maintenance of cognitive functions.

- **Tryptophan**

Tryptophan is an essential amino acid precursor of serotonin and melatonin. It promotes sleep and helps to reduce stress.

- **Tyrosine**

Tyrosine is a precursor amino acid of dopamine, adrenalin and noradrenalin. It helps maintain balance within the nervous system, playing an important role in regulating motivation, mood and stress.



TAXONOMIC ABUNDANCE TABLE

| Phylum | Class | Order | Family | Gender | Species | Abundance |
|----------------|---------------------|--------------------|---------------------|-------------------------------|---|-----------|
| Bacteroidota | Bacteroidia | Bacteroidales | Prevotellaceae | Segatella | Segatella_sinensis | 18.342% |
| Bacteroidota | Bacteroidia | Bacteroidales | Prevotellaceae | Segatella | Segatella_copri | 17.308% |
| Bacteroidota | Bacteroidia | Bacteroidales | Prevotellaceae | Segatella | Segatella_hominis | 11.757% |
| Bacteroidota | Bacteroidia | Bacteroidales | Prevotellaceae | Segatella | Segatella_brunsvicensis | 8.361% |
| Bacteroidota | Bacteroidia | Bacteroidales | Prevotellaceae | Segatella | Segatella_sanihominis | 7.670% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Faecalibacterium | Faecalibacterium_prausnitzii | 3.211% |
| Bacteroidota | Bacteroidia | Bacteroidales | Rikenellaceae | Alistipes | Alistipes_SGB2313 | 2.736% |
| Bacteroidota | Bacteroidia | Bacteroidales | Rikenellaceae | Alistipes | Alistipes_putredinis | 2.070% |
| Firmicutes | CFGB2947 | OFGB2947 | FGB2947 | GGB9286 | GGB9286_SGB27113 | 1.856% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Oscillibacter | Oscillibacter_sp_ER4 | 1.383% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Roseburia | Roseburia_inulinivorans | 1.199% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Lachnospira | Lachnospira_pectinoschiza | 1.158% |
| Proteobacteria | Betaproteobacteria | Burkholderiales | Sutterellaceae | Dakarella | Dakarella_massiliensis | 1.136% |
| Bacteroidota | Bacteroidia | Bacteroidales | Tannerellaceae | Parabacteroides | Parabacteroides_distasonis | 0.980% |
| Bacteroidota | Bacteroidia | Bacteroidales | Bacteroidaceae | Phocaeicola | Phocaeicola_vulgatus | 0.941% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | GGB13489 | GGB13489_SGB15224 | 0.858% |
| Bacteroidota | Bacteroidia | Bacteroidales | Bacteroidaceae | Phocaeicola | Phocaeicola_massiliensis | 0.830% |
| Firmicutes | CFGB8331 | OFGB8331 | FGB8331 | GGB9067 | GGB9067_SGB13986 | 0.829% |
| Bacteroidota | Bacteroidia | Bacteroidales | Tannerellaceae | Parabacteroides | Parabacteroides_merdae | 0.771% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | GGB3740 | GGB3740_SGB5076 | 0.767% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Wujia | Wujia_chippingensis | 0.758% |
| Bacteroidota | Bacteroidia | Bacteroidales | Bacteroidaceae | Bacteroides | Bacteroides_thetaiotaomicron | 0.704% |
| Bacteroidota | Bacteroidia | Bacteroidales | Bacteroidaceae | Bacteroides | Bacteroides_stercoris | 0.678% |
| Bacteroidota | Bacteroidia | Bacteroidales | Bacteroidaceae | Bacteroides | Bacteroides_uniformis | 0.648% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Lachnospiraceae_unclassified | Eubacterium_rectale | 0.628% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Brotolimicola | Brotolimicola_acetigignens | 0.588% |
| Firmicutes | Clostridia | Eubacteriales | Clostridiaceae | Clostridium | Clostridium_fessum | 0.585% |
| Firmicutes | Negativicutes | Veillonellales | Veillonellaceae | GGB4266 | GGB4266_SGB5809 | 0.524% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Simiaoa | Simiaoa_sunii | 0.443% |
| Bacteroidota | Bacteroidia | Bacteroidales | Barnesiellaceae | Barnesiella | Barnesiella_intestinihominis | 0.390% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Lacrimispora | Lacrimispora_amygdalina | 0.381% |
| Bacteroidota | Bacteroidia | Bacteroidales | Odoribacteraceae | Odoribacter | Odoribacter_splanchnicus | 0.364% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Oscillibacter | Oscillibacter_valericigenes | 0.343% |
| Firmicutes | CFGB2984 | OFGB2984 | FGB2984 | GGB9347 | GGB9347_SGB14313 | 0.325% |
| Firmicutes | CFGB2932 | OFGB2932 | FGB2932 | GGB9261 | GGB9261_SGB14209 | 0.320% |
| Firmicutes | CFGB1354 | OFGB1354 | FGB1354 | GGB3304 | GGB3304_SGB4367 | 0.315% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Enterocloster | Enterocloster_hominis | 0.306% |
| Bacteroidota | Bacteroidia | Bacteroidales | Rikenellaceae | Alistipes | Alistipes_shahii | 0.282% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | GGB33469 | GGB33469_SGB15236 | 0.261% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Dysosmobacter | Dysosmobacter_welbionis | 0.258% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | GGB9715 | GGB9715_SGB15267 | 0.251% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Mediterraneibacter | Mediterraneibacter_faecis | 0.251% |
| Firmicutes | Clostridia | Eubacteriales | Clostridiaceae | Clostridiaceae_unclassified | Clostridiaceae_bacterium_Marselle_Q4149 | 0.242% |
| Bacteroidota | Bacteroidia | Bacteroidales | Bacteroidaceae | Bacteroides | Bacteroides_finegoldii | 0.232% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | GGB9634 | GGB9634_SGB15093 | 0.226% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | GGB9707 | GGB9707_SGB15229 | 0.223% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | GGB3619 | GGB3619_SGB4894 | 0.215% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Oscillospiraceae_unclassified | Eubacterium_siraeum | 0.209% |
| Proteobacteria | Deltaproteobacteria | Desulfovibrionales | Desulfovibrionaceae | GGB9819 | GGB9819_SGB15460 | 0.194% |
| Proteobacteria | Deltaproteobacteria | Desulfovibrionales | Desulfovibrionaceae | Desulfovibrio | Desulfovibrio_fairfieldensis | 0.194% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | GGB9708 | GGB9708_SGB15233 | 0.184% |
| Bacteroidota | Bacteroidia | Bacteroidales | Bacteroidaceae | Bacteroides | Bacteroides_ovatus | 0.183% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | GGB9602 | GGB9602_SGB15031 | 0.181% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Faecalibacterium | Faecalibacterium_SGB15346 | 0.180% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Oscillospiraceae_unclassified | Oscillospiraceae_unclassified_SGB15256 | 0.172% |
| Firmicutes | Clostridia | Eubacteriales | Clostridiaceae | Clostridium | Clostridium_sp_AM22_11AC | 0.170% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Roseburia | Roseburia_intestinalis | 0.168% |

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Microbiome profiling

6 rue Pierre Joseph Colin - 35000 RENNES France



| Phylum | Class | Order | Family | Gender | Species | Abundance |
|----------------|---------------------|--------------------|----------------------------|-------------------------------|--|-----------|
| Proteobacteria | Gammaproteobacteria | Enterobacterales | Enterobacteriaceae | Enterobacter | Enterobacter_kobei | 0.158% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | GGB9699 | GGB9699_SGB15216 | 0.143% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Lachnospiraceae_unclassified | Lachnospiraceae_bacterium_A M48_27BH | 0.134% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Agathobaculum | Agathobaculum_butyriciproducens | 0.133% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Lachnospira | Lachnospira_eligens | 0.126% |
| Bacteroidota | Bacteroidia | Bacteroidales | Rikenellaceae | Alistipes | Alistipes_communis | 0.124% |
| Proteobacteria | Deltaproteobacteria | Desulfovibrionales | Desulfovibrionaceae | Bilophila | Bilophila_SGB15451 | 0.122% |
| Bacteroidota | Bacteroidia | Bacteroidales | Bacteroidaceae | Bacteroides | Bacteroides_caccae | 0.116% |
| Firmicutes | Clostridia | Eubacteriales | Clostridiaceae | Clostridiaceae_unclassified | Clostridiaceae_bacterium_AF18_31LB | 0.115% |
| Firmicutes | CFGB83839 | OFGB83839 | FGB83839 | GGB3328 | GGB3328_SGB4406 | 0.108% |
| Bacteroidota | Bacteroidia | Bacteroidales | Prevotellaceae | Paraprevotella | Paraprevotella_clara | 0.106% |
| Bacteroidota | Bacteroidia | Bacteroidales | Rikenellaceae | Alistipes | Alistipes_finegoldii | 0.096% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Lawsonibacter | Lawsonibacter_SGB15145 | 0.088% |
| Firmicutes | CFGB1298 | OFGB1298 | FGB1298 | GGB3118 | GGB3118_SGB4130 | 0.084% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Anaerostipes | Anaerostipes_faenicola | 0.083% |
| Firmicutes | Clostridia | Eubacteriales | Clostridiaceae | Clostridium | Clostridium_sp_AF27_2AA | 0.080% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | GGB3167 | GGB3167_SGB4181 | 0.080% |
| Firmicutes | Clostridia | Eubacteriales | Clostridiaceae | Clostridiaceae_unclassified | Clostridiaceae_bacterium_Mars_eille_Q4143 | 0.069% |
| Firmicutes | Clostridia | Eubacteriales | Eubacteriaceae | GGB3277 | GGB3277_SGB4327 | 0.069% |
| Firmicutes | Negativicutes | Veillonellales | Veillonellaceae | Megasphaera | Megasphaera_sp_NM10 | 0.067% |
| Firmicutes | Clostridia | Eubacteriales | Clostridiaceae | Clostridium | Clostridium_SGB4750 | 0.063% |
| Firmicutes | CFGB1776 | OFGB1776 | FGB1776 | GGB4684 | GGB4684_SGB6478 | 0.062% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | GGB79104 | GGB79104_SGB105821 | 0.054% |
| Bacteroidota | Bacteroidia | Bacteroidales | Bacteroidaceae | Bacteroides | Bacteroides_xylanisolvens | 0.053% |
| Firmicutes | Clostridia | Eubacteriales | Eubacteriales_unclassified | Colidextribacter | Colidextribacter_sp_210702_DFI_3_9 | 0.051% |
| Firmicutes | Clostridia | Eubacteriales | Eubacteriales_unclassified | Gemmiger | Gemmiger_formicilis | 0.045% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Vescimonas | Vescimonas_coprocola | 0.045% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | GGB9708 | GGB9708_SGB15234 | 0.043% |
| Bacteroidota | Bacteroidia | Bacteroidales | Odoribacteraceae | Butyricimonas | Butyricimonas_virosa | 0.042% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Faecalibacterium | Faecalibacterium_sp_HFFF | 0.039% |
| Firmicutes | CFGB72924 | OFGB72924 | FGB72924 | GGB3612 | GGB3612_SGB4882 | 0.038% |
| Firmicutes | CFGB2954 | OFGB2954 | FGB2954 | GGB9297 | GGB9297_SGB14255 | 0.032% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Coprococcus | Coprococcus_comes | 0.030% |
| Firmicutes | CFGB9989 | OFGB9989 | FGB9989 | GGB9712 | GGB9712_SGB15244 | 0.030% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | GGB9614 | GGB9614_SGB15049 | 0.028% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Flavonifractor | Flavonifractor_plautii | 0.028% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Lachnospiraceae_unclassified | Lachnospiraceae_bacterium_CL_AA_H244 | 0.028% |
| Firmicutes | CFGB2918 | OFGB2918 | FGB2918 | GGB9240 | GGB9240_SGB14182 | 0.025% |
| Firmicutes | CFGB2834 | OFGB2834 | FGB2834 | GGB9064 | GGB9064_SGB13983 | 0.025% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Oscillospiraceae_unclassified | Oscillospiraceae_bacterium_Marseille_Q3528 | 0.024% |
| Firmicutes | CFGB1211 | OFGB1211 | FGB1211 | GGB2970 | GGB2970_SGB3952 | 0.024% |
| Firmicutes | CFGB36753 | OFGB36753 | FGB36753 | GGB2658 | GGB2658_SGB3579 | 0.023% |
| Firmicutes | CFGB10477 | OFGB10477 | FGB10477 | GGB9345 | GGB9345_SGB14311 | 0.022% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Faecalicatena | Faecalicatena_fissicatena | 0.022% |
| Firmicutes | Clostridia | Eubacteriales | Clostridiaceae | Clostridiaceae_unclassified | Clostridiaceae_bacterium | 0.021% |
| Bacteroidota | Bacteroidia | Bacteroidales | Rikenellaceae | Alistipes | Alistipes_underdonkii | 0.021% |
| Firmicutes | Erysipelotrichia | Erysipelotrichales | Erysipelotrichaceae | Holdemanella | Holdemanella_porci | 0.021% |
| Bacteroidota | Bacteroidia | Bacteroidales | Porphyromonadaceae | Sanguibacteroides | Sanguibacteroides_justesenii | 0.019% |
| Firmicutes | Clostridia | Eubacteriales | Clostridiaceae | Clostridiaceae_unclassified | Clostridiaceae_unclassified_SGB4771 | 0.019% |
| Firmicutes | Clostridia | Eubacteriales | Clostridiaceae | Clostridium | Clostridium_sp_AM33_3 | 0.018% |
| Bacteroidota | Bacteroidia | Bacteroidales | Odoribacteraceae | Butyricimonas | Butyricimonas_faechominis | 0.017% |
| Bacteroidota | Bacteroidia | Bacteroidales | Rikenellaceae | Alistipes | Alistipes_indistinctus | 0.017% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Youxingia | Youxingia_wuxianensis | 0.016% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Dorea | Dorea_formicigenerans | 0.015% |
| Firmicutes | Clostridia | Eubacteriales | Lachnospiraceae | Dorea | Dorea_longicatena | 0.015% |
| Firmicutes | CFGB2840 | OFGB2840 | FGB2840 | GGB9093 | GGB9093_SGB14023 | 0.015% |
| Firmicutes | CFGB3070 | OFGB3070 | FGB3070 | GGB9775 | GGB9775_SGB15395 | 0.013% |



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| Phylum | Class | Order | Family | Gender | Species | Abundance |
|-----------------|---------------------|--------------------|---------------------|-----------------------------|---|-----------|
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Faecalibacterium | Faecalibacterium_sp_CLA_AA_H233 | 0.010% |
| Firmicutes | CFGB38642 | OFGB38642 | FGB38642 | GGB9758 | GGB9758_SGB15368 | 0.010% |
| Firmicutes | CFGB3023 | OFGB3023 | FGB3023 | GGB9501 | GGB9501_SGB14898 | 0.009% |
| Firmicutes | CFGB1814 | OFGB1814 | FGB1814 | GGB4802 | GGB4802_SGB6641 | 0.009% |
| Verrucomicrobia | Verrucomicrobiae | Verrucomicrobiales | Akkermansiaceae | Akkermansia | Akkermansia_muciniphila | 0.006% |
| Proteobacteria | Gammaproteobacteria | Enterobacteriales | Enterobacteriaceae | Escherichia | Escherichia_coli | 0.004% |
| Proteobacteria | Deltaproteobacteria | Desulfovibrionales | Desulfovibrionaceae | Bilophila | Bilophila_wadsworthia | 0.004% |
| Firmicutes | Clostridia | Eubacteriales | Oscillospiraceae | Lawsonibacter | Lawsonibacter_asaccharolyticus | 0.004% |
| Firmicutes | Clostridia | Eubacteriales | Clostridiaceae | Clostridiaceae_unclassified | Clostridiaceae_bacterium_Marselle_Q4145 | 0.001% |
| Firmicutes | CFGB4621 | OFGB4621 | FGB4621 | GGB18336 | GGB18336_SGB20700 | 0.001% |



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