

Actinobacteria

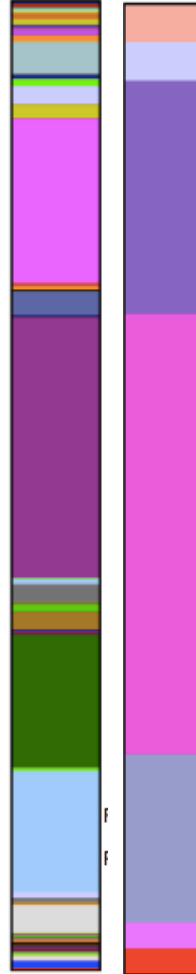
0.00%	Actinomyces species	0,001-1%
0.00%	Atopobium rimae	0,1-20%
0.01%	Bifidobacterium bifidum	0,1-20%
0.03%	Bifidobacterium longum	0,01-20%
0.02%	Collinsella aerofaciens	0,1-40%
0.28%	Collinsella stercoris	0,01-5%
0.01%	Coriobacteriaceae	0,01-1%
0.35%	Eggerthella lenta	0,01-1%
0.15%	Kiloniella species	0,1-20%

Bacteroidetes 71%

0.25%	Alistipes putredinis	0,01-20%
0.03%	Alistipes species	0,01-20%
0.01%	Bacteroides caccae	0,01-20%
0.15%	Bacteroides cellulosilyticus	0,001-10%
0.67%	Bacteroides dorei	0,001-2%
0.00%	Bacteroides eggerthii	0,01-1%
0.57%	Bacteroides fragilis	0,1-5%
0.24%	Bacteroides fragilis	0,1-3%
0.10%	Bacteroides massiliensis	0,001-0,5%
0.00%	Bacteroides ovatus	0,01-10%
0.71%	Bacteroides species	0,01-5%
0.68%	Bacteroides stercoris	0,01-10%
3.38%	Bacteroides vulgatus	0,1-10%
0.00%	Barnesiella species	0,01-5%
0.42%	Barnesiellaceae	0,1-5%
0.00%	Butyricimonas species	0,01-2%
0.01%	Odoribacter species	0,001-10%
0.03%	Odoribacter spianchnicus	0,01-1%
0.01%	Parabacteroides distasonis	0,01-1%
0.71%	Parabacteroides merdae	0,01-1%
0.02%	Paraprevotella species	0,01-0,1%
1.86%	Prevotella copri	0,001-0,5%
0.05%	Prevotella stercorea	0,001-5%
0.05%	Rikenellaceae	0,001-5%

Firmicutes 16%

0.03%	Acidaminococcus	0,01-0,1%
0.26%	Blautia luti	0,001-5%
0.01%	Blautia obeum	0,001-5%
0.01%	Blautia producta	0,001-0,5%
0.41%	Blautia species	0,001-0,5%
0.14%	Bulleidia moorei	0,01-3%
2.48%	Butyrivibrio spezieis	0,001-0,5%
0.26%	Clostridiales	0,0001-5%
0.00%	Clostridium species	0,01-1%
0.02%	Coprococcus species	0,01-1%
26.82%	Dialister invisus	0,01-0,1%
0.02%	Dialister species	0,001-0,5%
0.07%	Dorea formicigenerans	0,001-5%



0.06%	Dorea species	0,01-5%
0.48%	Dorea longicatena	0,01-1%
1.91%	Eisenbergiella species	0,001-1%
0.01%	Eisenbergiella tayi	0,001-1%
0.00%	Enterococcus species	0,001-0,1%
0.01%	Erysipelotrichaceae	0,001-0,1%
0.04%	Eubacterium coprostanoligenes	0,001-0,1%
0.04%	Eubacterium ramulus	0,001-0,1%
0.82%	Faecalibacterium prausnitzii	0,001-0,1%
0.02%	Fusicatenib. saccharivorans	0,01-5%
0.00%	Gemella species	0,01-1%
0.00%	Granulicatella species	0,001-1%
1.84%	Hungatella species	0,001-1%
0.13%	Lachnospra species	0,001-0,1%
0.34%	Lachnospiraceae	0,001-0,1%
0.02%	Lactobacillales	0,001-0,1%
13.80%	Lactobacillus mucosae	0,001-0,1%
0.00%	Lactobacillus species	0,001-0,1%
0.28%	Leuconostocaceae	0,001-0,1%
12.45%	Megamonas species	0,001-0,1%
0.64%	Micrococcus species	0,001-0,1%
0.50%	Oscillospira species	0,001-0,1%
0.01%	Peptostreptococcaceae	0,01-5%
0.21%	Phascolarctobacterium species	0,01-1%
2.90%	Roseburia species	0,001-1%
0.15%	Roseburia faecis	0,001-1%
0.06%	Roseburia finulinivorans	0,001-0,1%
0.04%	Roseburia inulinivorans	0,001-0,1%
0.29%	Ruminococcaceae	0,001-0,1%
0.15%	Ruminococcus bromii	0,001-0,1%
0.26%	Ruminococcus faecis	0,001-0,1%
0.31%	Ruminococcus gnavus	0,001-0,1%
0.15%	Ruminococcus species	0,001-0,1%
0.01%	Salinibacter ruber	0,001-0,1%
0.42%	Staphylococcus species	0,001-0,1%
0.01%	Streptococcus species	0,01-5%
0.14%	Turcibacter species	0,01-1%
0.14%	Veillonella dispar	0,001-0,1%
0.14%	Veillonella species	0,001-0,1%

Fusobacteria 0%

0.00%	Fusobacterium species	0,001-0,1%
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Proteobacteria 1.7%

0.01%	Bilophila species	0,001-0,1%
0.00%	Enterobacteriaceae	0,001-0,1%
0.03%	Erwinia species	0,001-0,1%
0.00%	Klebsiella species	0,001-0,1%
0.57%	Sutterella species	0,01-5%

Tenericutes 0.1%

0.71%	Mollicutes	0,001-1%
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Verrucomicrobia 0.0%

0.21%	Akkermansia muciniphila	0,001-0,1%
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