

Species SV group after experiment	Number of reads	Relative abundance
<i>Prevotella copri</i>	8047	20.23%
<i>Lactobacillus amylovorus</i>	6223	15.65%
<i>Barnesiella intestinihominis</i>	1835	4.61%
<i>Faecalibacterium prausnitzii</i>	1391	3.5%
<i>Megasphaera elsdenii</i>	1183	2.97%
<i>Collinsella aerofaciens</i>	1060	2.67%
<i>Prevotella brevis</i>	995	2.5%
<i>Clostridium cellulovorans</i>	969	2.44%
<i>Prevotella stercorea</i>	962	2.42%
Unclassified	938	2.36%
<i>Gemmiger formicilis</i>	897	2.26%
<i>Terrisporobacter glycolicus</i>	797	2%
<i>Olsenella scatoligenes</i>	524	1.32%
<i>Denitrobacterium detoxificans</i>	511	1.28%
<i>Eubacterium rectale</i>	492	1.24%
<i>Oscillospira guilliermondii</i>	460	1.16%
<i>Prevotella oris</i>	446	1.12%
<i>Roseburia faecis</i>	422	1.06%
<i>Oscillibacter ruminantium</i>	382	0.96%
<i>Prevotella maculosa</i>	345	0.87%
<i>Olsenella umbonata</i>	337	0.85%
<i>Enorma massiliensis</i>	318	0.8%
<i>Flintibacter butyricus</i>	298	0.75%
<i>Anaerovibrio lipolyticus</i>	287	0.72%
<i>Dialister succinatiphilus</i>	274	0.69%
<i>Enterorhabdus mucosicola</i>	231	0.58%
<i>Sporobacter termitidis</i>	230	0.58%
<i>Alloprevotella rava</i>	224	0.56%
<i>Olsenella uli</i>	214	0.54%
<i>Prevotella oralis</i>	212	0.53%
<i>Senegalimassilia anaerobia</i>	206	0.52%
<i>Butyricococcus pullicaecorum</i>	203	0.51%
<i>Fournierella massiliensis</i>	191	0.48%
<i>Intestinimonas butyriciproducens</i>	188	0.47%
<i>Blautia wexlerae</i>	180	0.45%
<i>Lactobacillus reuteri</i>	179	0.45%
<i>Catenibacterium mitsuokai</i>	168	0.42%
<i>Desulfovibrio piger</i>	157	0.39%
<i>Clostridium celatum</i>	150	0.38%
unclassified Planctomycetales	146	0.37%
<i>Oligosphaera ethanolica</i>	127	0.32%
<i>Prevotella dentalis</i>	123	0.31%
<i>Selenomonas bovis</i>	121	0.3%
<i>Lactobacillus pontis</i>	114	0.29%
<i>Olivibacter sitiensis</i>	105	0.26%
<i>Treponema succinifaciens</i>	103	0.26%
Bacteroidales oral	102	0.26%
unclassified Tannerella	99	0.25%
<i>Parvibacter caecicola</i>	95	0.24%

Gracilibacter thermotolerans	94 0.24%
Prevotella histicola	94 0.24%
Intestinibacter bartlettii	89 0.22%
Ruminiclostridium thermocellum	87 0.22%
Lactobacillus delbrueckii	87 0.22%
Succinivibrio dextrinosolvens	86 0.22%
Romboutsia sedimentorum	85 0.21%
unclassified Bacteroidales	82 0.21%
unclassified Erysipelotrichaceae	81 0.2%
Parabacteroides distasonis	80 0.2%
Lactobacillus kitasatonis	80 0.2%
unclassified Barnesiella	79 0.2%
Fibrobacter intestinalis	77 0.19%
Eubacterium ramulus	75 0.19%
Lactobacillus jensenii	73 0.18%
Intestinimonas timonensis	71 0.18%
Prevotella buccae	69 0.17%
Olsenella profusa	66 0.17%
Lactobacillus crispatus	65 0.16%
Eubacterium coprostanoligenes	62 0.16%
unclassified Prevotella	58 0.15%
Intestinimonas massiliensis	57 0.14%
Fusicatenibacter saccharivorans	57 0.14%
Acetanaerobacterium elongatum	54 0.14%
Prevotella salivae	53 0.13%
Holdemanella biformis	52 0.13%
Paraeggerthella hongkongensis	52 0.13%
Acidaminobacter hydrogenoformans	52 0.13%
Slackia isoflavoniconvertens	52 0.13%
Solobacterium moorei	50 0.13%
Paraprevotella clara	50 0.13%
Lactobacillus panis	50 0.13%
Clostridium phoceensis	47 0.12%
Sutterella stercoricanis	47 0.12%
Ruminococcus faecis	47 0.12%
Blautia obeum	47 0.12%
Coprococcus catus	46 0.12%
Turicibacter sanguinis	45 0.11%
Phascolarctobacterium succinatutens	44 0.11%
Eubacterium desmolans	43 0.11%
Coprococcus comes	40 0.1%
Ruthenibacterium lactatiformans	39 0.1%
Blautia massiliensis	38 0.1%
unclassified Prevotellaceae	38 0.1%
Enterorhabdus caecimuris	37 0.09%
Erysipelothrix inopinata	37 0.09%
Desulfovibrio fairfieldensis	37 0.09%
Mitsuokella jalaludinii	36 0.09%
Eubacterium hallii	35 0.09%
Ruminococcus bromii	34 0.09%

Christensenella minuta	32 0.08%
Eubacterium ruminantium	32 0.08%
Eubacterium eligens	30 0.08%
Selenomonas ruminantium	30 0.08%
Prevotella ruminicola	29 0.07%
Lactobacillus helveticus	29 0.07%
Prevotella genomosp.	29 0.07%
Brassicibacter thermophilus	29 0.07%
Butyrivibrio fibrisolvens	28 0.07%
Adlercreutzia equolifaciens	28 0.07%
Roseburia inulinivorans	27 0.07%
Collinsella intestinalis	27 0.07%
cyanobacterium enrichment	26 0.07%
Clostridium cellobioparum	26 0.07%
Ruminococcus bicirculans	26 0.07%
unclassified Rikenella	26 0.07%
Pseudoflavonifractor capillosus	26 0.07%
Candidatus Soleaferrea	25 0.06%
Lutispora thermophila	24 0.06%
Geosporobacter ferrireducens	24 0.06%
Ruminococcus flavefaciens	24 0.06%
Clostridium quinii	24 0.06%
unclassified Clostridium	24 0.06%
Papillibacter cinnamivorans	23 0.06%
unclassified Lachnospiraceae	23 0.06%
Oscillibacter valericigenes	22 0.06%
Barnesiella viscericola	21 0.05%
Asaccharospora irregularis	21 0.05%
Dorea formicigenerans	21 0.05%
Prevotella paludivivens	20 0.05%
Dorea longicatena	20 0.05%
Lactobacillus frumenti	20 0.05%
Lactobacillus acidophilus	19 0.05%
Saccharofermentans acetigenes	19 0.05%
Prevotella conceptionensis	19 0.05%
Slackia exigua	19 0.05%
Blautia faecis	19 0.05%
unclassified Deltaproteobacteria	19 0.05%
Anaerobacterium chartisolvens	19 0.05%
Murimonas intestini	18 0.05%
Hungatella hathewayi	18 0.05%
Bifidobacterium pseudolongum	17 0.04%
Flavonifractor plautii	17 0.04%
Vallitalea pronyensis	17 0.04%
Clostridium amylolyticum	17 0.04%
unclassified Ruminococcaceae	16 0.04%
Oribacterium sinus	16 0.04%
Campylobacter lanienae	15 0.04%
unclassified Porphyromonadaceae	15 0.04%
Natranaerovirga pectinivora	15 0.04%

<i>Clostridium aldenense</i>	15 0.04%
<i>Clostridium bovipellis</i>	14 0.04%
unclassified <i>Olsenella</i>	14 0.04%
<i>Parasutterella secunda</i>	14 0.04%
unclassified <i>Anaerovibrio</i>	14 0.04%
<i>Clostridium methylpentosum</i>	13 0.03%
<i>Ruminococcus torques</i>	13 0.03%
<i>Roseburia intestinalis</i>	13 0.03%
<i>Collinsella massiliensis</i>	12 0.03%
unclassified Clostridia	12 0.03%
<i>Succiniclasticum ruminis</i>	12 0.03%
<i>Peptococcus simiae</i>	12 0.03%
<i>Clostridium chartatabidum</i>	12 0.03%
<i>Roseburia hominis</i>	12 0.03%
<i>Prevotella denticola</i>	12 0.03%
<i>Collinsella stercoris</i>	12 0.03%
<i>Anaeromassilibacillus senegalensis</i>	12 0.03%
<i>Desulfovibrio desulfuricans</i>	12 0.03%
<i>Acetivibrio cellulolyticus</i>	12 0.03%
<i>Clostridium polysaccharolyticum</i>	11 0.03%
<i>Gorbachella massiliensis</i>	11 0.03%
<i>Anaerotaenia torta</i>	11 0.03%
<i>Intestinimonas gabonensis</i>	11 0.03%
unclassified <i>Paludibacter</i>	11 0.03%
unclassified Clostridiales	11 0.03%
actinobacterium enrichment	11 0.03%
unclassified Chlamydia	11 0.03%
Candidatus <i>Heliomonas</i>	10 0.03%
<i>Methylocystis rosea</i>	10 0.03%
<i>Eubacterium siraeum</i>	10 0.03%
unclassified Mollicutes	10 0.03%
<i>Caloramator fervidus</i>	10 0.03%
<i>Eisenbergiella tayi</i>	10 0.03%
<i>Megasphaera hominis</i>	10 0.03%
<i>Clostridium leptum</i>	10 0.03%
<i>Acidaminococcus fermentans</i>	10 0.03%
<i>Clostridium cadaveris</i>	10 0.03%
<i>Clostridium oroticum</i>	10 0.03%
<i>Eubacterium oxidoreducens</i>	10 0.03%
<i>Clostridium disporicum</i>	9 0.02%
<i>Subdoligranulum variabile</i>	9 0.02%
<i>Selenomonas sputigena</i>	9 0.02%
Bifidobacteriaceae genomosp.	9 0.02%
Bacteroidales genomosp.	9 0.02%
unclassified Bacteroidia	9 0.02%
<i>Anaerostipes hadrus</i>	9 0.02%
<i>Blautia stercoris</i>	9 0.02%
<i>Clostridium cellulolyticum</i>	9 0.02%
<i>Clostridium xylanolyticum</i>	9 0.02%
<i>Sphaerochaeta coccoides</i>	8 0.02%

Falcatimonas natans	8 0.02%
Clostridium chauvoei	8 0.02%
Blautia luti	8 0.02%
Slackia piriformis	8 0.02%
Clostridium sartagoforme	8 0.02%
Blautia glucersea	8 0.02%
unclassified Turicibacter	8 0.02%
Anaerovorax odorimutans	8 0.02%
Lachnospira pectinoschiza	7 0.02%
Helicobacter rodentium	7 0.02%
Garciella nitratreducens	7 0.02%
Eubacteriaceae oral	7 0.02%
Clostridium intestinale	7 0.02%
Holdemania filiformis	7 0.02%
Prevotella dentasini	7 0.02%
Herbinix luporum	7 0.02%
Desulfotomaculum halophilum	7 0.02%
Mucispirillum schaedleri	7 0.02%
Prevotella baroniae	7 0.02%
Ruminococcus gnavus	7 0.02%
Faecalicoccus acidiformans	7 0.02%
Acetivibrio ethanolgignens	7 0.02%
Ruminococcus callidus	7 0.02%
Clostridium aurantibutyricum	6 0.02%
Eubacterium rangiferina	6 0.02%
Clostridium saccharolyticum	6 0.02%
Zhizhongheella caldifontis	6 0.02%
Bacteroides heparinolyticus	6 0.02%
Pseudomonas fluorescens	6 0.02%
Clostridium hiranonis	6 0.02%
Porphyromonas catoniae	6 0.02%
Lactobacillus mucosae	6 0.02%
Propionispira arcuata	6 0.02%
unclassified Wautersiella	6 0.02%
Lactobacillus gallinarum	6 0.02%
Ethanoligenens harbinense	6 0.02%
Clostridium hveragerdense	6 0.02%
Treponema porcinum	6 0.02%
Clostridium clariflavum	6 0.02%
Mitsuokella multacida	5 0.01%
Abyssivirga alkaniphila	5 0.01%
unclassified Spirochaetia	5 0.01%
Eubacterium plexicaudatum	5 0.01%
Eubacterium infirmum	5 0.01%
Terrisporobacter petrolearius	5 0.01%
unclassified Eubacterium	5 0.01%
Clostridium tertium	5 0.01%
Prevotella loescheii	5 0.01%
Coprococcus eutactus	5 0.01%
Alloprevotella tanneriae	5 0.01%

Ruminobacter amylophilus	5 0.01%
Elbe River	5 0.01%
unclassified Bulleidia	5 0.01%
Clostridium clostridioforme	5 0.01%
Propionispira paucivorans	5 0.01%
Bacteroides vulgatus	5 0.01%
Clostridium papyrosolvens	5 0.01%
unclassified Enterococcus	5 0.01%
Holdemania massiliensis	4 0.01%
Prevotella buccalis	4 0.01%
Parabacteroides chinchillae	4 0.01%
Asteroleplasma anaerobium	4 0.01%
Bacteroides zoogloformans	4 0.01%
Clostridium hungatei	4 0.01%
Candidatus Dorea	4 0.01%
Parabacteroides merdae	4 0.01%
Clostridium viride	4 0.01%
Corynebacterium provencense	4 0.01%
Clostridium butyricum	4 0.01%
unclassified Oscillospira	4 0.01%
Ruminococcus champanellensis	4 0.01%
unclassified Actinobacteria	4 0.01%
Clostridium lavalense	4 0.01%
Clostridium aminobutyricum	4 0.01%
Faecalitalea cylindroides	4 0.01%
Mogibacterium diversum	4 0.01%
Prevotella fusca	4 0.01%
Acidaminococcus intestini	4 0.01%
Caminiella sporogenes	4 0.01%
Clostridium isatidis	4 0.01%
Catabacter hongkongensis	4 0.01%
Escherichia coli	4 0.01%
Lactobacillus amylolyticus	4 0.01%
Paeniclostridium sordellii	4 0.01%
unclassified Roseburia	4 0.01%
Dialister propionificiens	4 0.01%
Butyrivibrio crossotus	4 0.01%
Caproiciproducens galactitolivorans	4 0.01%
Ruminococcus albus	3 0.01%
Eggerthella lenta	3 0.01%
Treponema zioleckii	3 0.01%
Bacteroides ovatus	3 0.01%
Clostridium longisporum	3 0.01%
Prevotella timonensis	3 0.01%
unclassified Oscillibacter	3 0.01%
unclassified Candidatus Solibacter	3 0.01%
Allisonella histaminiformans	3 0.01%
Bacteroides nordii	3 0.01%
unclassified Faecalibacterium	3 0.01%
unclassified Coriobacteriaceae	3 0.01%

Agathobacter ruminis	3 0.01%
Prevotella shahii	3 0.01%
Clostridium fusiformis	3 0.01%
Sutterella parvirubra	3 0.01%
Lactobacillus vaginalis	3 0.01%
Lactonifactor longoviformis	3 0.01%
Hespellia porcina	3 0.01%
unclassified Dialister	3 0.01%
Eubacterium contortum	3 0.01%
Sphaerochaeta pleomorpha	3 0.01%
Bacteroides caecicola	3 0.01%
Clostridium populeti	3 0.01%
Clostridium fimetarium	3 0.01%
Robinsoniella peoriensis	3 0.01%
Desulfotomaculum tongense	3 0.01%
Lactobacillus psittaci	3 0.01%
Clostridium colicanis	3 0.01%
Bifidobacterium boum	3 0.01%
Streptococcus gallolyticus	3 0.01%
Bacteroides uniformis	3 0.01%
Bacteroides helcogenes	3 0.01%
unclassified Candidatus Glomeribacter	3 0.01%
Oceanirhabdus sediminicola	3 0.01%
Clostridium moniliforme	3 0.01%
Lactobacillus johnsonii	3 0.01%
alpha proteobacterium	3 0.01%
Natronincola histidinovorans	3 0.01%
Clostridium celerecrescens	2 0.01%
Porphyromonas cangingivalis	2 0.01%
Paludibacter jiangxiensis	2 0.01%
Lentzea violacea	2 0.01%
metal-contaminated soil	2 0.01%
unclassified Bacteroides	2 0.01%
Pseudomonas lurida	2 0.01%
Bifidobacterium kashiwanohense	2 0.01%
Clostridioides difficile	2 0.01%
Coriobacterium glomerans	2 0.01%
Bifidobacterium choerinum	2 0.01%
Bacteroides clarus	2 0.01%
Clostridium glycyrrhizinilyticum	2 0.01%
unclassified Collinsella	2 0.01%
Actinomyces denticolens	2 0.01%
Prevotella jejuni	2 0.01%
Prevotella corporis	2 0.01%
Clostridium stercorarium	2 0.01%
Eubacterium tenue	2 0.01%
Bifidobacterium breve	2 0.01%
Blautia schinkii	2 0.01%
Lactobacillus hamsteri	2 0.01%
Lactobacillus fermentum	2 0.01%

Moorella humiferrea	2 0.01%
Bacteroides cellulosilyticus	2 0.01%
Anaerospobacter mobilis	2 0.01%
unclassified Clostridiaceae	2 0.01%
Thermotalea metallivorans	2 0.01%
Collinsella tanakaei	2 0.01%
Bacteroides stercoris	2 0.01%
Terrisporobacter mayombeii	2 0.01%
Cellulosibacter alkalithermophilus	2 0.01%
Geosporobacter subterraneus	2 0.01%
Magaeibacillus indolicus	2 0.01%
unclassified Solobacterium	2 0.01%
Clostridium aminophilum	2 0.01%
Prevotella bryantii	2 0.01%
Anaerostipes butyraticus	2 0.01%
Laceyella putida	2 0.01%
Hallella seregens	2 0.01%
gamma proteobacterium	2 0.01%
unclassified Acetivibrio	2 0.01%
Bacteroides barnesiae	2 0.01%
Lachnoanaerobaculum umeaense	2 0.01%
Campylobacter jejuni	2 0.01%
Bifidobacterium animalis	2 0.01%
Eisenbergiella massiliensis	2 0.01%
Prevotella bivia	2 0.01%
Treponema brennaborensis	2 0.01%
Bacteroides fragilis	2 0.01%
Clostridium straminisolvens	2 0.01%
unclassified Acidobacterium	2 0.01%
unclassified Treponema	2 0.01%
Clostridium asparagiforme	2 0.01%
unclassified Veillonellaceae	2 0.01%
unclassified Sutterella	2 0.01%
Marvinbryantia formatexigens	2 0.01%
Clostridium tarantellae	2 0.01%
Defluviitalea raffinosedens	2 0.01%
Anaerocolumna cellulosilytica	2 0.01%
unclassified Lactobacillus	2 0.01%
Blautia hansenii	1 0%
Prevotella albensis	1 0%
Methylocystis echinoides	1 0%
Lachnospiraceae oral	1 0%
unclassified Subdoligranulum	1 0%
Nakamurella panacisegetis	1 0%
Roseburia cecicola	1 0%
Leifsonia kafniensis	1 0%
Anaerocolumna aminovalerica	1 0%
Slackia faecicanis	1 0%
Clostridium taeniosporum	1 0%
Gordonibacter urolithinifaciens	1 0%

Megasphaera genomosp.	1 0%
Gottschalkia acidurici	1 0%
Atopobium vaginae	1 0%
Bacteroides faecichinchillae	1 0%
Eubacterium pyruvativorans	1 0%
Pleomorphochaeta multiformis	1 0%
Tyzzera nexilis	1 0%
Nostocoida limicola	1 0%
Catenisphaera adipataaccumulans	1 0%
Parabacteroides johnsonii	1 0%
Blautia producta	1 0%
Sphingopyxis panaciterrae	1 0%
Streptococcus danieliae	1 0%
Bacteroides caecigallinarum	1 0%
Mycobacterium tuberculosis	1 0%
proteobacterium enrichment	1 0%
Treponema berlinense	1 0%
Ercella succinigenes	1 0%
Peptoclostridium litorale	1 0%
Oligotropha carboxidovorans	1 0%
unclassified Spirochaeta	1 0%
Hespellia stercorisuis	1 0%
Thermanaerovibrio velox	1 0%
Streptomyces scabrisporus	1 0%
Bacteroides dorei	1 0%
Prevotella oulorum	1 0%
Butyrivibrio cf.	1 0%
Prevotella amnii	1 0%
Slackia equolifaciens	1 0%
Byssovorax cruenta	1 0%
Lactobacillus rodentium	1 0%
Tindallia magadiensis	1 0%
Parasporobacterium paucivorans	1 0%
methanogenic archaeon	1 0%
Megasphaera indica	1 0%
Prevotella scopos	1 0%
unclassified Victivallaceae	1 0%
Bacteroides timonensis	1 0%
unclassified Catonella	1 0%
unclassified Beggiatoa	1 0%
Bacteroides salanitronis	1 0%
Paludibaculum fermentans	1 0%
Pseudobutyrvibrio ruminis	1 0%
Clostridium indolis	1 0%
Bacteroides salyersiae	1 0%
Corynebacterium variabile	1 0%
Bacillus pumilus	1 0%
Clostridium polyendosporum	1 0%
unclassified Staphylococcus	1 0%
Clostridium amygdalinum	1 0%

unclassified Candidatus Rhabdochlamydia	1 0%
Bacteroides coprophilus	1 0%
unclassified Rhodoferax	1 0%
Pedosphaera parvula	1 0%
Desulfotomaculum nigrificans	1 0%
Prevotella multiformis	1 0%
Desulfovibrio multispirans	1 0%
Lactobacillus gastricus	1 0%
Bacteroides luti	1 0%
Clostridium botulinum	1 0%
Streptococcus macedonicus	1 0%
Bifidobacterium longum	1 0%
Stomatobaculum longum	1 0%
Nitrospirillum amazonense	1 0%
Clostridium sufflavum	1 0%
unclassified Alloprevotella	1 0%
Mobilitalea sibirica	1 0%
Marinilabilia salmonicolor	1 0%
Anaeroplasma bactoclasticum	1 0%
Clostridium elmenteitii	1 0%
Clostridium tyrobutyricum	1 0%
Anaerofilum pentosovorans	1 0%
Ruminococcus gauvreauii	1 0%
beta proteobacterium	1 0%
Novosphingobium sediminicola	1 0%
Caloramator australicus	1 0%
CFB group	1 0%
Stenotrophobacter terrae	1 0%
unclassified Achromobacter	1 0%
Lactobacillus camelliae	1 0%
Aeriscardovia aeriphila	1 0%
Eubacterium cellulosolvens	1 0%
Casaltella massiliensis	1 0%
Janthinobacterium agaricidamnosum	1 0%
Rhodoplanes roseus	1 0%
Schlegelella aquatica	1 0%
Anaerotruncus colihominis	1 0%
Desulfomonas oviles	1 0%
Streptococcus lutetiensis	1 0%
unclassified Bifidobacterium	1 0%
Hathewayia histolytica	1 0%
Actinocorallia cavernae	1 0%
Clostridium frigoriphilum	1 0%
unclassified Gordonibacter	1 0%
Bulleidia extracta	1 0%
Clostridium symbiosum	1 0%
unclassified Alphaproteobacteria	1 0%
Clostridium bornimense	1 0%
Lactobacillus secaliphilus	1 0%
Lachnospira multipara	1 0%

<i>Alistipes senegalensis</i>	1 0%
<i>Prevotella micans</i>	1 0%
unclassified <i>Kopriimonas</i>	1 0%
<i>Bifidobacterium saguini</i>	1 0%
<i>Eubacterium limosum</i>	1 0%
<i>Ruminococcus lactaris</i>	1 0%
<i>Methanosphaera cuniculi</i>	1 0%
<i>Breznakia pachnodae</i>	1 0%
<i>Acetatifactor muris</i>	1 0%
<i>Clostridium lactatifermentans</i>	1 0%
<i>Tannerella forsythia</i>	1 0%
<i>Bacteroides acidifaciens</i>	1 0%
<i>Tetracoccus cechii</i>	1 0%
<i>Propionispira arboris</i>	1 0%
<i>Eubacterium saphenum</i>	1 0%
<i>Conexibacter woesei</i>	1 0%
<i>Vitis hybrid</i>	1 0%
unclassified <i>Desulfovibrio</i>	1 0%
<i>Eggerthella sinensis</i>	1 0%
<i>Catonella morbi</i>	1 0%
<i>Pseudomonas putida</i>	1 0%
<i>Actinomyces naeslundii</i>	1 0%
<i>Clostridium sulfidigenes</i>	1 0%
<i>Bariatricus massiliensis</i>	1 0%
unclassified <i>Sphingobacterium</i>	1 0%
<i>Methyloceanibacter caenitepidi</i>	1 0%
<i>Rarimicrobium hominis</i>	1 0%
<i>Lachnoanaerobaculum</i> cf.	1 0%
<i>Parabacteroides faecis</i>	1 0%
<i>Eubacterium xylanophilum</i>	1 0%
<i>Paraclostridium benzoelyticum</i>	1 0%
unclassified <i>Erysipelotrichia</i>	1 0%
unclassified <i>Sporobacter</i>	1 0%
<i>Centipeda periodontii</i>	1 0%
<i>Methylosinus sporium</i>	1 0%
<i>Pediococcus inopinatus</i>	1 0%
<i>Lactobacillus gigeriorum</i>	1 0%
<i>Clostridium alkalicellulosi</i>	1 0%
<i>Rhodanobacter spathiphylli</i>	1 0%
<i>Helicobacter canadensis</i>	1 0%
<i>Clostridium septicum</i>	1 0%
<i>Parabacteroides goldsteinii</i>	1 0%