

Supplementary Table 4. Bacterial taxa identified by mNGS using an RPM ratio metric

*re-classified to a higher taxonomic rank because reads aligned equally well to multiple different organisms that shared the same species, genus or family

#positive result according to an RPM (reads per million) ratio ≥ 10 , where the RPM ratio = RPM(sample) / RPM(NTC)

&reads correspond to the *Streptococcus agalactiae* PC

Abbreviations: PC, positive control; CSF, cerebrospinal fluid; mNGS, metagenomic next-generation sequencing; RPM, reads per million

| species | genus | family | Normalized Reads Per million | |
|--|-----------------------|-------------------------|------------------------------|-------------------|
| | | | DNA PC | DNA Patient CSF |
| * | Brucella | Brucellaceae | 0.0 | 15.7 [#] |
| Brucella melitensis | Brucella | Brucellaceae | 0.0 | 1.6 |
| Streptococcus agalactiae | Streptococcus | Streptococcaceae | 3,264.1 [#] | 0.0 |
| Klebsiella pneumoniae | Klebsiella | Enterobacteriaceae | 52.9 [#] | 0.0 |
| * | * | Enterobacteriaceae | 0.1 | 0.0 |
| Escherichia coli | Escherichia | Enterobacteriaceae | 0.0 | 0.0 |
| Propionibacterium acnes | Propionibacterium | Propionibacteriaceae | 0.0 | 0.0 |
| * | Streptococcus | Streptococcaceae | 8.7 | 0.0 |
| * | Klebsiella | Enterobacteriaceae | 41.6 [#] | 0.0 |
| * | * | * | 0.0 | 0.0 |
| Streptococcus suis | Streptococcus | Streptococcaceae | 16.2 ^{#&} | 0.0 |
| Streptococcus salivarius | Streptococcus | Streptococcaceae | 1.0 | 0.0 |
| Streptococcus pyogenes | Streptococcus | Streptococcaceae | 30.7 ^{#&} | 0.0 |
| Serratia marcescens | Serratia | Enterobacteriaceae | 2.3 | 0.0 |
| Lactococcus lactis | Lactococcus | Streptococcaceae | 0.4 | 0.0 |
| * | Escherichia | Enterobacteriaceae | 0.0 | 0.0 |
| Pseudomonas sp. TKP | Pseudomonas | Pseudomonadaceae | 0.1 | 0.0 |
| * | Pseudomonas | Pseudomonadaceae | 0.2 | 0.0 |
| Streptococcus macedonicus | Streptococcus | Streptococcaceae | 18.2 ^{#&} | 0.0 |
| Enterobacter cloacae | Enterobacter | Enterobacteriaceae | 0.4 | 0.0 |
| Klebsiella oxytoca | Klebsiella | Enterobacteriaceae | 0.7 | 0.0 |
| Thermoanaerobacterium thermosaccharophilum | Thermoanaerobacterium | Thermoanaerobacteriales | 0.2 | 0.0 |
| Streptococcus sp. VT 162 | Streptococcus | Streptococcaceae | 0.6 | 0.0 |
| Enterococcus faecium | Enterococcus | Enterococcaceae | 0.1 | 0.1 |
| Pseudomonas protegens | Pseudomonas | Pseudomonadaceae | 3.0 | 0.1 |
| Micrococcus luteus | Micrococcus | Micrococcaceae | 0.0 | 0.0 |

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|---------------------------------|-------------------|----------------------|-----|-----|
| Streptococcus infantarius | Streptococcus | Streptococcaceae | 0.5 | 0.0 |
| Staphylococcus epidermidis | Staphylococcus | Staphylococcaceae | 0.0 | 0.0 |
| Enterobacter asburiae | Enterobacter | Enterobacteriaceae | 0.2 | 0.0 |
| Cupriavidus metallidurans | Cupriavidus | Burkholderiaceae | 0.0 | 0.0 |
| Pseudomonas putida | Pseudomonas | Pseudomonadaceae | 0.1 | 0.0 |
| Enterococcus casseliflavus | Enterococcus | Enterococcaceae | 0.1 | 0.1 |
| Streptococcus mitis | Streptococcus | Streptococcaceae | 0.6 | 0.0 |
| Streptococcus equi | Streptococcus | Streptococcaceae | 5.2 | 0.0 |
| Salmonella enterica | Salmonella | Enterobacteriaceae | 4.9 | 0.0 |
| Streptococcus oralis | Streptococcus | Streptococcaceae | 0.5 | 0.0 |
| Klebsiella variicola | Klebsiella | Enterobacteriaceae | 4.3 | 0.0 |
| Burkholderia lata | Burkholderia | Burkholderiaceae | 0.3 | 0.0 |
| Pseudomonas stutzeri | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Streptococcus pneumoniae | Streptococcus | Streptococcaceae | 0.4 | 0.0 |
| Streptococcus dysgalactiae | Streptococcus | Streptococcaceae | 3.7 | 0.0 |
| * | Burkholderia | Burkholderiaceae | 0.1 | 0.0 |
| Pseudomonas fluorescens | Pseudomonas | Pseudomonadaceae | 0.4 | 0.0 |
| Acinetobacter guillouiae | Acinetobacter | Moraxellaceae | 0.1 | 0.0 |
| Veillonella parvula | Veillonella | Veillonellaceae | 0.1 | 0.0 |
| Xanthomonas campestris | Xanthomonas | Xanthomonadaceae | 0.1 | 0.1 |
| Exiguobacterium sp. AT1b | Exiguobacterium | | 0.1 | 0.0 |
| Pseudomonas pseudoalcaligenes | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Streptococcus pasteurianus | Streptococcus | Streptococcaceae | 2.8 | 0.0 |
| Rothia dentocariosa | Rothia | Micrococcaceae | 0.1 | 0.0 |
| * | * | Streptococcaceae | 2.6 | 0.0 |
| Delftia acidovorans | Delftia | Comamonadaceae | 0.5 | 0.0 |
| * | Propionibacterium | Propionibacteriaceae | 0.0 | 0.0 |
| Acinetobacter baumannii | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| Streptococcus parasanguinis | Streptococcus | Streptococcaceae | 0.0 | 0.0 |
| Pseudomonas sp. WCS374 | Pseudomonas | Pseudomonadaceae | 0.8 | 0.0 |
| Enterobacter aerogenes | Enterobacter | Enterobacteriaceae | 2.3 | 0.0 |
| Stenotrophomonas maltophilia | Stenotrophomonas | Xanthomonadaceae | 0.2 | 0.0 |
| Alicyclobacillus acidocaldarius | Alicyclobacillus | Alicyclobacillaceae | 0.2 | 0.0 |
| Haemophilus influenzae | Haemophilus | Pasteurellaceae | 0.1 | 0.0 |
| Rothia mucilaginosa | Rothia | Micrococcaceae | 0.0 | 0.0 |
| Staphylococcus xylosus | Staphylococcus | Staphylococcaceae | 0.2 | 0.0 |
| Acidovorax sp. JS42 | Acidovorax | Comamonadaceae | 0.1 | 0.0 |
| * | Rahnella | Enterobacteriaceae | 1.8 | 0.0 |
| Streptococcus pseudopneumoniae | Streptococcus | Streptococcaceae | 0.3 | 0.0 |
| Streptococcus thermophilus | Streptococcus | Streptococcaceae | 0.1 | 0.0 |
| Pseudomonas aeruginosa | Pseudomonas | Pseudomonadaceae | 0.1 | 0.0 |

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|--|------------------------------|------------------------|-----|-----|
| <i>Corynebacterium kroppenstedtii</i> | <i>Corynebacterium</i> | Corynebacteriaceae | 1.6 | 0.0 |
| <i>Staphylococcus haemolyticus</i> | <i>Staphylococcus</i> | Staphylococcaceae | 0.0 | 0.0 |
| <i>Serratia</i> sp. SCBI | <i>Serratia</i> | Enterobacteriaceae | 1.5 | 0.0 |
| * | <i>Enterococcus</i> | Enterococcaceae | 0.1 | 0.0 |
| <i>Staphylococcus warneri</i> | <i>Staphylococcus</i> | Staphylococcaceae | 0.0 | 0.0 |
| <i>Burkholderia cepacia</i> | <i>Burkholderia</i> | Burkholderiaceae | 0.1 | 0.1 |
| * | <i>Corynebacterium</i> | Corynebacteriaceae | 0.0 | 0.0 |
| <i>Comamonas testosteroni</i> | <i>Comamonas</i> | Comamonadaceae | 0.0 | 0.0 |
| <i>Bifidobacterium thermophilum</i> | <i>Bifidobacterium</i> | Bifidobacteriaceae | 1.3 | 0.0 |
| * | <i>Lactobacillus</i> | Lactobacillaceae | 0.2 | 0.0 |
| <i>Thermoanaerobacterium xylanolyticum</i> | <i>Thermoanaerobacterium</i> | Thermoanaerobacterales | 0.1 | 0.0 |
| <i>Ralstonia pickettii</i> | <i>Ralstonia</i> | Burkholderiaceae | 0.1 | 0.0 |
| <i>Meiothermus ruber</i> | <i>Meiothermus</i> | Thermaceae | 0.0 | 0.0 |
| <i>Acidovorax ebreus</i> | <i>Acidovorax</i> | Comamonadaceae | 0.7 | 0.0 |
| <i>Micrococcus</i> sp. V7 | <i>Micrococcus</i> | Micrococcaceae | 0.2 | 0.0 |
| <i>Leuconostoc mesenteroides</i> | <i>Leuconostoc</i> | Leuconostocaceae | 0.0 | 0.0 |
| <i>Bacillus halodurans</i> | <i>Bacillus</i> | Bacillaceae | 0.7 | 0.0 |
| <i>Corynebacterium variabile</i> | <i>Corynebacterium</i> | Corynebacteriaceae | 0.3 | 0.0 |
| * | <i>Acidovorax</i> | Comamonadaceae | 0.1 | 0.0 |
| <i>Bifidobacterium bifidum</i> | <i>Bifidobacterium</i> | Bifidobacteriaceae | 0.0 | 0.0 |
| <i>Rhizobium</i> sp. IRBG74 | <i>Rhizobium</i> | Rhizobiaceae | 0.0 | 0.0 |
| * | <i>Acinetobacter</i> | Moraxellaceae | 0.0 | 0.0 |
| <i>Acinetobacter calcoaceticus</i> | <i>Acinetobacter</i> | Moraxellaceae | 1.0 | 0.0 |
| <i>Chroococcidiopsis thermalis</i> | <i>Chroococcidiopsis</i> | | 0.2 | 0.0 |
| <i>Streptococcus sanguinis</i> | <i>Streptococcus</i> | Streptococcaceae | 0.1 | 0.0 |
| <i>Acidovorax</i> sp. KKS102 | <i>Acidovorax</i> | Comamonadaceae | 0.1 | 0.0 |
| <i>Raoultella ornithinolytica</i> | <i>Raoultella</i> | Enterobacteriaceae | 0.1 | 0.0 |
| <i>Ochrobactrum anthropi</i> | <i>Ochrobactrum</i> | Brucellaceae | 0.0 | 0.0 |
| <i>Lactobacillus johnsonii</i> | <i>Lactobacillus</i> | Lactobacillaceae | 0.6 | 0.1 |
| <i>Methylobacterium populi</i> | <i>Methylobacterium</i> | Methylobacteriaceae | 0.2 | 0.0 |
| <i>Rhodococcus equi</i> | <i>Rhodococcus</i> | Nocardiaceae | 0.1 | 0.0 |
| <i>Lactobacillus helveticus</i> | <i>Lactobacillus</i> | Lactobacillaceae | 0.9 | 0.0 |
| * | <i>Serratia</i> | Enterobacteriaceae | 0.2 | 0.0 |
| <i>Burkholderia cenocepacia</i> | <i>Burkholderia</i> | Burkholderiaceae | 0.3 | 0.0 |
| * | <i>Staphylococcus</i> | Staphylococcaceae | 0.0 | 0.0 |
| <i>Enterobacter</i> sp. R4-368 | <i>Enterobacter</i> | Enterobacteriaceae | 0.8 | 0.0 |
| <i>Propionibacterium propionicum</i> | <i>Propionibacterium</i> | Propionibacteriaceae | 0.1 | 0.0 |
| <i>Streptococcus gordonii</i> | <i>Streptococcus</i> | Streptococcaceae | 0.1 | 0.0 |
| <i>Corynebacterium singulare</i> | <i>Corynebacterium</i> | Corynebacteriaceae | 0.1 | 0.0 |
| <i>Burkholderia ambifaria</i> | <i>Burkholderia</i> | Burkholderiaceae | 0.7 | 0.1 |
| * | <i>Micrococcus</i> | Micrococcaceae | 0.0 | 0.0 |

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|---------------------------------------|-----------------------|------------------------|-----|-----|
| Fervidobacterium nodosum | Fervidobacterium | Thermotogaceae | 0.0 | 0.0 |
| Aeromonas media | Aeromonas | Aeromonadaceae | 0.0 | 0.0 |
| Cronobacter sakazakii | Cronobacter | Enterobacteriaceae | 0.7 | 0.0 |
| Myroides profundi | Myroides | Flavobacteriaceae | 0.7 | 0.0 |
| Methylobacterium oryzae | Methylobacterium | Methylobacteriaceae | 0.7 | 0.0 |
| * | Xanthomonas | Xanthomonadaceae | 0.3 | 0.0 |
| Thermoanaerobacterium saccharolyticum | Thermoanaerobacterium | Thermoanaerobacterales | 0.1 | 0.0 |
| Pseudomonas mendocina | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Corynebacterium ureicelerivorans | Corynebacterium | Corynebacteriaceae | 0.0 | 0.0 |
| Lactobacillus crispatus | Lactobacillus | Lactobacillaceae | 0.0 | 0.0 |
| Alicyclophilus denitrificans | Alicyclophilus | Comamonadaceae | 0.0 | 0.0 |
| Gardnerella vaginalis | Gardnerella | Bifidobacteriaceae | 0.0 | 0.0 |
| * | Gemella | | 0.6 | 0.0 |
| * | Ralstonia | Burkholderiaceae | 0.6 | 0.0 |
| Eggerthella lenta | Eggerthella | Coriobacteriaceae | 0.6 | 0.0 |
| * | * | Rhizobiaceae | 0.5 | 0.0 |
| Prevotella denticola | Prevotella | Prevotellaceae | 0.2 | 0.0 |
| Prevotella intermedia | Prevotella | Prevotellaceae | 0.2 | 0.0 |
| Psychrobacter sp. PRwf-1 | Psychrobacter | Moraxellaceae | 0.1 | 0.0 |
| Azospira oryzae | Azospira | Rhodocyclaceae | 0.1 | 0.0 |
| Acinetobacter haemolyticus | Acinetobacter | Moraxellaceae | 0.1 | 0.0 |
| * | Delftia | Comamonadaceae | 0.1 | 0.0 |
| Burkholderia contaminans | Burkholderia | Burkholderiaceae | 0.4 | 0.2 |
| Arthrobacter arilaitensis | Arthrobacter | Micrococcaceae | 0.1 | 0.1 |
| Dermacoccus nishinomiyensis | Dermacoccus | Dermacoccaceae | 0.1 | 0.1 |
| Pantoea ananatis | Pantoea | Enterobacteriaceae | 0.1 | 0.0 |
| Staphylococcus saprophyticus | Staphylococcus | Staphylococcaceae | 0.0 | 0.0 |
| Staphylococcus pasteurii | Staphylococcus | Staphylococcaceae | 0.0 | 0.0 |
| Rahnella aquatilis | Rahnella | Enterobacteriaceae | 0.5 | 0.0 |
| Rahnella sp. Y9602 | Rahnella | Enterobacteriaceae | 0.5 | 0.0 |
| Campylobacter concisus | Campylobacter | Campylobacteraceae | 0.5 | 0.0 |
| Geobacillus sp. WCH70 | Geobacillus | Bacillaceae | 0.5 | 0.0 |
| * | Frankia | Frankiaceae | 0.5 | 0.0 |
| Lactobacillus casei | Lactobacillus | Lactobacillaceae | 0.5 | 0.0 |
| Thiomonas intermedia | Thiomonas | | 0.5 | 0.0 |
| Streptococcus gallolyticus | Streptococcus | Streptococcaceae | 0.5 | 0.0 |
| Thioalkalivibrio sulfidiphilus | Thioalkalivibrio | Ectothiorhodospiraceae | 0.3 | 0.0 |
| * | Bradyrhizobium | Bradyrhizobiaceae | 0.2 | 0.0 |
| Bifidobacterium longum | Bifidobacterium | Bifidobacteriaceae | 0.1 | 0.0 |
| Corynebacterium falsenii | Corynebacterium | Corynebacteriaceae | 0.1 | 0.0 |
| Delftia sp. Cs1-4 | Delftia | Comamonadaceae | 0.1 | 0.0 |

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|-----------------------------------|-------------------|----------------------|-----|-----|
| Acinetobacter sp. M131 | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| Prevotella melaninogenica | Prevotella | Prevotellaceae | 0.0 | 0.0 |
| * | * | Comamonadaceae | 0.0 | 0.0 |
| Leuconostoc carnosum | Leuconostoc | Leuconostocaceae | 0.2 | 0.0 |
| Pectobacterium carotovorum | Pectobacterium | Enterobacteriaceae | 0.0 | 0.0 |
| * | Myroides | Flavobacteriaceae | 0.4 | 0.0 |
| * | Erwinia | Enterobacteriaceae | 0.4 | 0.0 |
| Gordonia sp. KTR9 | Gordonia | Gordoniaceae | 0.4 | 0.0 |
| Paenibacillus sp. FSL R7-0273 | Paenibacillus | Paenibacillaceae | 0.4 | 0.0 |
| Paracoccus sp. N81106 | Paracoccus | Rhodobacteraceae | 0.4 | 0.0 |
| Sphingobium fuliginis | Sphingobium | Sphingomonadaceae | 0.4 | 0.0 |
| * | * | | 0.4 | 0.0 |
| * | Geobacillus | Bacillaceae | 0.2 | 0.0 |
| Pseudomonas sp. VLB120 | Pseudomonas | Pseudomonadaceae | 0.2 | 0.0 |
| Pelagibacterium halotolerans | Pelagibacterium | Hyphomicrobiaceae | 0.1 | 0.0 |
| Streptococcus intermedius | Streptococcus | Streptococcaceae | 0.1 | 0.0 |
| Propionibacterium freudenreichii | Propionibacterium | Propionibacteriaceae | 0.1 | 0.0 |
| * | Enterobacter | Enterobacteriaceae | 0.0 | 0.0 |
| Nakamurella multipartita | Nakamurella | Nakamurellaceae | 0.1 | 0.1 |
| Haemophilus parasuis | Haemophilus | Pasteurellaceae | 0.3 | 0.0 |
| Fusobacterium nucleatum | Fusobacterium | Fusobacteriaceae | 0.3 | 0.0 |
| Citrobacter freundii | Citrobacter | Enterobacteriaceae | 0.3 | 0.0 |
| Ruminococcus sp. SR1/5 | Ruminococcus | Ruminococcaceae | 0.2 | 0.0 |
| Pseudoxanthomonas spadix | Pseudoxanthomonas | Xanthomonadaceae | 0.2 | 0.0 |
| Lactococcus garvieae | Lactococcus | Streptococcaceae | 0.1 | 0.0 |
| Neisseria elongata | Neisseria | Neisseriaceae | 0.1 | 0.0 |
| Acidovorax citrulli | Acidovorax | Comamonadaceae | 0.1 | 0.0 |
| Novosphingobium pentaromativorans | Novosphingobium | Sphingomonadaceae | 0.1 | 0.0 |
| Citrobacter koseri | Citrobacter | Enterobacteriaceae | 0.1 | 0.0 |
| Methylobacterium aquaticum | Methylobacterium | Methylobacteriaceae | 0.1 | 0.0 |
| Pseudomonas denitrificans | Pseudomonas | Pseudomonadaceae | 0.1 | 0.0 |
| Rhodococcus erythropolis | Rhodococcus | Nocardiaceae | 0.0 | 0.0 |
| Lactobacillus reuteri | Lactobacillus | Lactobacillaceae | 0.0 | 0.0 |
| Bacteroides fragilis | Bacteroides | Bacteroidaceae | 0.0 | 0.0 |
| Lactobacillus plantarum | Lactobacillus | Lactobacillaceae | 0.0 | 0.0 |
| * | Bacillus | Bacillaceae | 0.0 | 0.0 |
| Pseudomonas rhizosphaerae | Pseudomonas | Pseudomonadaceae | 0.2 | 0.1 |
| Achromobacter xylosoxidans | Achromobacter | Alcaligenaceae | 0.0 | 0.0 |
| Lactobacillus amylovorus | Lactobacillus | Lactobacillaceae | 0.2 | 0.0 |
| Propionibacterium acidipropionici | Propionibacterium | Propionibacteriaceae | 0.2 | 0.0 |
| Leuconostoc gelidum | Leuconostoc | Leuconostocaceae | 0.2 | 0.0 |

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|-------------------------------------|-------------------|------------------------|-----|-----|
| Weissella thailandensis | Weissella | Leuconostocaceae | 0.2 | 0.0 |
| Pandoraea sp. RB-44 | Pandoraea | Burkholderiaceae | 0.2 | 0.0 |
| Escherichia vulneris | Escherichia | Enterobacteriaceae | 0.2 | 0.0 |
| Yersinia intermedia | Yersinia | Enterobacteriaceae | 0.2 | 0.0 |
| Flavobacteriaceae bacterium 3519-10 | | Flavobacteriaceae | 0.2 | 0.0 |
| * | Rhodococcus | Nocardiaceae | 0.2 | 0.0 |
| Streptococcus sp. (N1) | Streptococcus | Streptococcaceae | 0.2 | 0.0 |
| * | Methylobacterium | Methylobacteriaceae | 0.1 | 0.0 |
| Sphingomonas sp. MM-1 | Sphingomonas | Sphingomonadaceae | 0.1 | 0.0 |
| Rhizobium etli | Rhizobium | Rhizobiaceae | 0.1 | 0.0 |
| * | Agrobacterium | Rhizobiaceae | 0.0 | 0.0 |
| Thermus scotoductus | Thermus | Thermaceae | 0.0 | 0.0 |
| Methylobacterium extorquens | Methylobacterium | Methylobacteriaceae | 0.0 | 0.0 |
| Streptococcus sp. I-P16 | Streptococcus | Streptococcaceae | 0.0 | 0.0 |
| Pantoea sp. PSNIH1 | Pantoea | Enterobacteriaceae | 0.0 | 0.0 |
| Methylobacterium radiotolerans | Methylobacterium | Methylobacteriaceae | 0.0 | 0.0 |
| [Ruminococcus] torques | Blautia | Lachnospiraceae | 0.1 | 0.4 |
| Lactobacillus sakei | Lactobacillus | Lactobacillaceae | 0.1 | 0.1 |
| * | * | Rhodocyclaceae | 0.1 | 0.1 |
| Bacillus licheniformis | Bacillus | Bacillaceae | 0.0 | 0.1 |
| Corynebacterium accolens | Corynebacterium | Corynebacteriaceae | 0.1 | 0.0 |
| Corynebacterium sp. ATCC 6931 | Corynebacterium | Corynebacteriaceae | 0.1 | 0.0 |
| Serratia symbiotica | Serratia | Enterobacteriaceae | 0.1 | 0.0 |
| Lactobacillus delbrueckii | Lactobacillus | Lactobacillaceae | 0.1 | 0.0 |
| Bacillus sp. YP1 | Bacillus | Bacillaceae | 0.1 | 0.0 |
| Klebsiella sp. PG122E | Klebsiella | Enterobacteriaceae | 0.1 | 0.0 |
| * | Rathayibacter | Microbacteriaceae | 0.1 | 0.0 |
| Pseudoalteromonas sp. P30 | Pseudoalteromonas | Pseudoalteromonadaceae | 0.1 | 0.0 |
| Staphylococcus sp. CDC25 | Staphylococcus | Staphylococcaceae | 0.1 | 0.0 |
| Corynebacterium resistens | Corynebacterium | Corynebacteriaceae | 0.1 | 0.0 |
| Shigella dysenteriae | Shigella | Enterobacteriaceae | 0.1 | 0.0 |
| * | * | Xanthomonadaceae | 0.1 | 0.0 |
| Agrobacterium fabrum | Agrobacterium | Rhizobiaceae | 0.0 | 0.0 |
| Gordonia polyisoprenivorans | Gordonia | Gordoniaceae | 0.0 | 0.0 |
| Pseudomonas balearica | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| * | * | Pseudomonadaceae | 0.0 | 0.0 |
| Ruminococcus bromii | Ruminococcus | Ruminococcaceae | 0.0 | 0.0 |
| Brachybacterium faecium | Brachybacterium | Dermabacteraceae | 0.0 | 0.0 |
| Acinetobacter johnsonii | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| Micrococcus sp. A1 | Micrococcus | Micrococcaceae | 0.0 | 0.0 |
| Filifactor alocis | Filifactor | Peptostreptococcaceae | 0.0 | 0.0 |

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|--|--------------------------|---------------------|-----|-----|
| <i>Pantoea vagans</i> | <i>Pantoea</i> | Enterobacteriaceae | 0.0 | 0.0 |
| <i>Haemophilus parainfluenzae</i> | <i>Haemophilus</i> | Pasteurellaceae | 0.0 | 0.0 |
| <i>Pantoea rwandensis</i> | <i>Pantoea</i> | Enterobacteriaceae | 0.0 | 0.3 |
| <i>Corynebacterium vitaeruminis</i> | <i>Corynebacterium</i> | Corynebacteriaceae | 0.0 | 0.2 |
| <i>Pseudomonas poae</i> | <i>Pseudomonas</i> | Pseudomonadaceae | 0.0 | 0.1 |
| * | * | Brucellaceae | 0.0 | 0.1 |
| <i>Lactobacillus fermentum</i> | <i>Lactobacillus</i> | Lactobacillaceae | 0.0 | 0.1 |
| <i>Anabaena variabilis</i> | <i>Anabaena</i> | Nostocaceae | 0.0 | 0.1 |
| <i>Sphingobacterium</i> sp. ML3W | <i>Sphingobacterium</i> | Sphingobacteriaceae | 0.0 | 0.1 |
| sugarcane isolate 74-1 | | | 0.0 | 0.1 |
| * | * | Geodermatophilaceae | 0.0 | 0.1 |
| <i>Megasphaera elsdenii</i> | <i>Megasphaera</i> | Veillonellaceae | 0.0 | 0.1 |
| <i>Pseudoxanthomonas suwonensis</i> | <i>Pseudoxanthomonas</i> | Xanthomonadaceae | 0.0 | 0.1 |
| <i>Corynebacterium glutamicum</i> | <i>Corynebacterium</i> | Corynebacteriaceae | 0.0 | 0.0 |
| <i>Sphingomonas taxi</i> | <i>Sphingomonas</i> | Sphingomonadaceae | 0.0 | 0.0 |
| <i>Pseudomonas graminis</i> | <i>Pseudomonas</i> | Pseudomonadaceae | 0.0 | 0.0 |
| <i>Bradyrhizobium</i> sp. BTAi1 | <i>Bradyrhizobium</i> | Bradyrhizobiaceae | 0.0 | 0.0 |
| <i>Enterococcus hirae</i> | <i>Enterococcus</i> | Enterococcaceae | 0.0 | 0.0 |
| <i>Corynebacterium</i> sp. L2-79-05 | <i>Corynebacterium</i> | Corynebacteriaceae | 0.0 | 0.0 |
| <i>Arthrobacter phenanthrenivorans</i> | <i>Arthrobacter</i> | Micrococcaceae | 0.0 | 0.0 |
| <i>Corynebacterium maris</i> | <i>Corynebacterium</i> | Corynebacteriaceae | 0.0 | 0.0 |
| <i>Gordonia bronchialis</i> | <i>Gordonia</i> | Gordoniaceae | 0.0 | 0.0 |
| <i>Kytococcus sedentarius</i> | <i>Kytococcus</i> | Dermacoccaceae | 0.0 | 0.0 |
| <i>Kosakonia cowanii</i> | <i>Kosakonia</i> | Enterobacteriaceae | 0.0 | 0.0 |
| <i>Xenorhabdus bovienii</i> | <i>Xenorhabdus</i> | Enterobacteriaceae | 0.0 | 0.0 |
| <i>Paracoccus haeundaensis</i> | <i>Paracoccus</i> | Rhodobacteraceae | 0.0 | 0.0 |
| <i>Methylobacterium</i> sp. 238 | <i>Methylobacterium</i> | Methylobacteriaceae | 0.0 | 0.0 |
| <i>Acinetobacter</i> sp. BW3 | <i>Acinetobacter</i> | Moraxellaceae | 0.0 | 0.0 |
| <i>Aeromonas sobria</i> | <i>Aeromonas</i> | Aeromonadaceae | 0.0 | 0.0 |
| <i>Bacillus lehensis</i> | <i>Bacillus</i> | Bacillaceae | 0.0 | 0.0 |
| <i>Ralstonia solanacearum</i> | <i>Ralstonia</i> | Burkholderiaceae | 0.0 | 0.0 |
| <i>Citrobacter</i> sp. FPO3 | <i>Citrobacter</i> | Enterobacteriaceae | 0.0 | 0.0 |
| <i>Citrobacter</i> sp. I91-3 | <i>Citrobacter</i> | Enterobacteriaceae | 0.0 | 0.0 |
| <i>Erwinia amylovora</i> | <i>Erwinia</i> | Enterobacteriaceae | 0.0 | 0.0 |
| <i>Klebsiella milletis</i> | <i>Klebsiella</i> | Enterobacteriaceae | 0.0 | 0.0 |
| <i>Salmonella bongori</i> | <i>Salmonella</i> | Enterobacteriaceae | 0.0 | 0.0 |
| <i>Serratia grimesii</i> | <i>Serratia</i> | Enterobacteriaceae | 0.0 | 0.0 |
| <i>Yersinia pestis</i> | <i>Yersinia</i> | Enterobacteriaceae | 0.0 | 0.0 |
| * | | Enterobacteriaceae | 0.0 | 0.0 |
| <i>Lactobacillus brevis</i> | <i>Lactobacillus</i> | Lactobacillaceae | 0.0 | 0.0 |
| <i>Kocuria</i> sp. starX | <i>Kocuria</i> | Micrococcaceae | 0.0 | 0.0 |

| | | | | |
|--|---------------------|-----------------------|-----|-----|
| Acinetobacter sp. 26 | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| Peptoclostridium difficile | Peptoclostridium | Peptostreptococcaceae | 0.0 | 0.0 |
| Sorangium cellulosum | Sorangium | Polyangiaceae | 0.0 | 0.0 |
| Pseudomonas sp. NSi14 | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Synergistetes oral clone 03 5 D05 | | | 0.0 | 0.0 |
| bacterium EBAD26 | | | 0.0 | 0.0 |
| bacterium NLAE-zl-G351 | | | 0.0 | 0.0 |
| rumen bacterium enrichment culture clone Y74 | | | 0.0 | 0.0 |
| unidentified marine bacterioplankton | | | 0.0 | 0.0 |
| Escherichia albertii | Escherichia | Enterobacteriaceae | 0.0 | 0.0 |
| Mobiluncus curtisii | Mobiluncus | Actinomycetaceae | 0.0 | 0.0 |
| * | Caulobacter | Caulobacteraceae | 0.0 | 0.0 |
| Methylothermobacter versatilis | Methylothermobacter | Methylophilaceae | 0.0 | 0.0 |
| Propionibacterium sp. KPL1849 | Propionibacterium | Propionibacteriaceae | 0.0 | 0.0 |
| bacterium EBAD25 | | | 0.0 | 0.0 |
| Bacillus sp. Pc3 | Bacillus | Bacillaceae | 0.0 | 0.0 |
| Acinetobacter sp. EVA14 | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| Agrobacterium sp. | Agrobacterium | Rhizobiaceae | 0.0 | 0.0 |
| Alistipes shahii | Alistipes | Rikenellaceae | 0.0 | 0.0 |
| * | Thermus | Thermaceae | 0.0 | 0.0 |
| * | Methylobacterium | | 0.0 | 0.0 |
| butyrate-producing bacterium SSC/2 | | | 0.0 | 0.0 |
| Escherichia fergusonii | Escherichia | Enterobacteriaceae | 0.0 | 0.0 |
| Enterobacter sp. Ni15 | Enterobacter | Enterobacteriaceae | 0.0 | 0.0 |
| Capnocytophaga ochracea | Capnocytophaga | Flavobacteriaceae | 0.0 | 0.0 |
| Thauera sp. 6NLG | Thauera | Rhodocyclaceae | 0.0 | 0.0 |
| Desulfovibrio alaskensis | Desulfovibrio | Desulfovibrionaceae | 0.0 | 0.0 |
| Variovorax sp. Alb14 | Variovorax | Comamonadaceae | 0.0 | 0.0 |
| * | Shigella | Enterobacteriaceae | 0.0 | 0.0 |
| * | * | Micromonosporaceae | 0.0 | 0.0 |
| * | Micromonospora | Micromonosporaceae | 0.0 | 0.0 |
| Thermobifida fusca | Thermobifida | Nocardiopsaceae | 0.0 | 0.0 |
| Turneriella parva | Turneriella | Leptospiraceae | 0.0 | 0.0 |
| [Clostridium] sticklandii | Peptoclostridium | Peptostreptococcaceae | 0.0 | 0.0 |
| Acinetobacter sp. Ooi24 | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| Ochrobactrum sp. SJY1 | Ochrobactrum | Brucellaceae | 0.0 | 0.0 |
| Carnobacterium sp. WN1359 | Carnobacterium | Carnobacteriaceae | 0.0 | 0.0 |
| Iamia majanohamensis | Iamia | Iamiaceae | 0.0 | 0.0 |
| Saccharomonospora viridis | Saccharomonospora | Pseudonocardiaceae | 0.0 | 0.0 |
| Rhizobium sp. | Rhizobium | Rhizobiaceae | 0.0 | 0.0 |
| Staphylococcus sp. CDC3 | Staphylococcus | Staphylococcaceae | 0.0 | 0.0 |

| | | | | |
|-------------------------------------|-------------------|------------------------|-----|-----|
| Shigella sonnei | Shigella | Enterobacteriaceae | 0.0 | 0.0 |
| Pseudomonas syringae | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Burkholderia vietnamiensis | Burkholderia | Burkholderiaceae | 0.0 | 0.0 |
| Shigella boydii | Shigella | Enterobacteriaceae | 0.0 | 0.0 |
| Bacillus weihenstephanensis | Bacillus | Bacillaceae | 0.0 | 0.0 |
| Erythrobacter litoralis | Erythrobacter | Erythrobacteraceae | 0.0 | 0.0 |
| Pseudoalteromonas haloplanktis | Pseudoalteromonas | Pseudoalteromonadaceae | 0.0 | 0.0 |
| Pseudomonas sp. FG1182 | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| * | Rhizobium | Rhizobiaceae | 0.0 | 0.0 |
| * | Rickettsia | Rickettsiaceae | 0.0 | 0.0 |
| Sphingobium yanoikuyae | Sphingobium | Sphingomonadaceae | 0.0 | 0.0 |
| Stenotrophomonas rhizophila | Stenotrophomonas | Xanthomonadaceae | 0.0 | 0.0 |
| * | Leuconostoc | Leuconostocaceae | 0.0 | 0.0 |
| Aquicola tertiarycarbonis | Aquicola | | 0.0 | 0.0 |
| Nocardiopsis dassonvillei | Nocardiopsis | Nocardiopsaceae | 0.0 | 0.0 |
| Carnobacterium maltaromaticum | Carnobacterium | Carnobacteriaceae | 0.0 | 0.0 |
| * | Haemophilus | Pasteurellaceae | 0.0 | 0.0 |
| Bordetella parapertussis | Bordetella | Alcaligenaceae | 0.0 | 0.0 |
| * | Dietzia | Dietziaceae | 0.0 | 0.0 |
| Shewanella sp. W3-18-1 | Shewanella | Shewanellaceae | 0.0 | 0.0 |
| Sphingomonas sp. NP5 | Sphingomonas | Sphingomonadaceae | 0.0 | 0.0 |
| Staphylococcus gallinarum | Staphylococcus | Staphylococcaceae | 0.0 | 0.0 |
| Micavibrio aeruginosavorus | Micavibrio | | 0.0 | 0.0 |
| Paracoccus denitrificans | Paracoccus | Rhodobacteraceae | 0.0 | 0.0 |
| [Cellvibrio] gilvus | Cellulomonas | Cellulomonadaceae | 0.0 | 0.0 |
| Corynebacterium jeikeium | Corynebacterium | Corynebacteriaceae | 0.0 | 0.0 |
| * | * | Staphylococcaceae | 0.0 | 0.0 |
| Meiothermus silvanus | Meiothermus | Thermaceae | 0.0 | 0.0 |
| Asticcacaulis excentricus | Asticcacaulis | Caulobacteraceae | 0.0 | 0.0 |
| * | Atopobium | Coriobacteriaceae | 0.0 | 0.0 |
| Streptococcus constellatus | Streptococcus | Streptococcaceae | 0.0 | 0.0 |
| Microcystis aeruginosa | Microcystis | | 0.0 | 0.0 |
| agricultural soil bacterium SC-I-13 | | | 0.0 | 0.0 |
| [Ruminococcus] obeum | Blautia | Lachnospiraceae | 0.0 | 0.0 |
| Thermus thermophilus | Thermus | Thermaceae | 0.0 | 0.0 |
| Shigella flexneri | Shigella | Enterobacteriaceae | 0.0 | 0.0 |
| * | Mycobacterium | Mycobacteriaceae | 0.0 | 0.0 |
| Pseudomonas savastanoi | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Staphylococcus capitis | Staphylococcus | Staphylococcaceae | 0.0 | 0.0 |
| * | Cupriavidus | Burkholderiaceae | 0.0 | 0.0 |
| Dyadobacter fermentans | Dyadobacter | Cytophagaceae | 0.0 | 0.0 |

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|-------------------------------|----------------------|--------------------|-----|-----|
| Dietzia sp. CQ4 | Dietzia | Dietziaceae | 0.0 | 0.0 |
| * | * | Methylophilaceae | 0.0 | 0.0 |
| * | Neisseria | Neisseriaceae | 0.0 | 0.0 |
| Corynebacterium aurimucosum | Corynebacterium | Corynebacteriaceae | 0.0 | 0.0 |
| * | | | 0.0 | 0.0 |
| Pseudomonas fulva | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Chromohalobacter salexigens | Chromohalobacter | Halomonadaceae | 0.0 | 0.0 |
| Brevundimonas diminuta | Brevundimonas | Caulobacteraceae | 0.0 | 0.0 |
| Streptococcus lutetiensis | Streptococcus | Streptococcaceae | 0.0 | 0.0 |
| Bordetella petrii | Bordetella | Alcaligenaceae | 0.0 | 0.0 |
| Erythrobacter sp. JP13.1 | Erythrobacter | Erythrobacteraceae | 0.0 | 0.0 |
| Methylobacillus glycogenes | Methylobacillus | Methylophilaceae | 0.0 | 0.0 |
| Candidatus Rhodoluna ladicola | Candidatus Rhodoluna | Microbacteriaceae | 0.0 | 0.0 |
| Arthrobacter sp. JBH1 | Arthrobacter | Micrococcaceae | 0.0 | 0.0 |
| Aggregatibacter aphrophilus | Aggregatibacter | Pasteurellaceae | 0.0 | 0.0 |
| Thauera sp. B4 | Thauera | Rhodocyclaceae | 0.0 | 0.0 |
| Lysobacter dokdonensis | Lysobacter | Xanthomonadaceae | 0.0 | 0.0 |
| Clostridiales genomosp. BVAB3 | | | 0.0 | 0.0 |
| Streptococcus sp. I-G2 | Streptococcus | Streptococcaceae | 0.0 | 0.0 |
| Pseudomonas mandelii | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Bradyrhizobium sp. S23321 | Bradyrhizobium | Bradyrhizobiaceae | 0.0 | 0.0 |
| Phenylobacterium zucineum | Phenylobacterium | Caulobacteraceae | 0.0 | 0.0 |
| Pseudomonas mosselii | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Staphylococcus lugdunensis | Staphylococcus | Staphylococcaceae | 0.0 | 0.0 |
| Proteus mirabilis | Proteus | Enterobacteriaceae | 0.0 | 0.0 |
| * | * | Neisseriaceae | 0.0 | 0.0 |
| Arthrobacter sp. J3-40 | Arthrobacter | Micrococcaceae | 0.0 | 0.0 |
| * | Pantoea | Enterobacteriaceae | 0.0 | 0.0 |
| Corynebacterium efficiens | Corynebacterium | Corynebacteriaceae | 0.0 | 0.0 |
| * | Halomonas | Halomonadaceae | 0.0 | 0.0 |
| Trueperella pyogenes | Trueperella | Actinomycetaceae | 0.0 | 0.0 |
| Streptomyces coelicolor | Streptomyces | Streptomycetaceae | 0.0 | 0.0 |
| Kocuria rhizophila | Kocuria | Micrococcaceae | 0.0 | 0.0 |
| Bacillus cereus | Bacillus | Bacillaceae | 0.0 | 0.0 |
| Tannerella forsythia | Tannerella | Porphyromonadaceae | 0.0 | 0.0 |
| * | Alkalibacterium | Carnobacteriaceae | 0.0 | 0.0 |
| Atopobium parvulum | Atopobium | Coriobacteriaceae | 0.0 | 0.0 |
| Serinicoccus profundi | Serinicoccus | Intrasporangiaceae | 0.0 | 0.0 |
| * | Leptotrichia | Leptotrichiaceae | 0.0 | 0.0 |
| * | * | Planococcaceae | 0.0 | 0.0 |
| Planomicrobium okeanokoites | Planomicrobium | Planococcaceae | 0.0 | 0.0 |

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|---|--------------------|----------------------|-----|-----|
| Jannaschia sp. CCS1 | Jannaschia | Rhodobacteraceae | 0.0 | 0.0 |
| Paracoccus aestuarii | Paracoccus | Rhodobacteraceae | 0.0 | 0.0 |
| Rhodobacter blasticus | Rhodobacter | Rhodobacteraceae | 0.0 | 0.0 |
| Agrobacterium tumefaciens | Agrobacterium | Rhizobiaceae | 0.0 | 0.0 |
| Shewanella sp. ANA-3 | Shewanella | Shewanellaceae | 0.0 | 0.0 |
| Pseudomonas cichorii | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Halomonas sp. A3H3 | Halomonas | Halomonadaceae | 0.0 | 0.0 |
| Serratia liquefaciens | Serratia | Enterobacteriaceae | 0.0 | 0.0 |
| Sphingopyxis alaskensis | Sphingopyxis | Sphingomonadaceae | 0.0 | 0.0 |
| * | Brevundimonas | Caulobacteraceae | 0.0 | 0.0 |
| Deinococcus deserti | Deinococcus | Deinococcaceae | 0.0 | 0.0 |
| Desulfovibrio vulgaris | Desulfovibrio | Desulfovibrionaceae | 0.0 | 0.0 |
| Propionibacterium sp. NTS31307302 | Propionibacterium | Propionibacteriaceae | 0.0 | 0.0 |
| Paracoccus alcaliphilus | Paracoccus | Rhodobacteraceae | 0.0 | 0.0 |
| Vibrio parahaemolyticus | Vibrio | Vibrionaceae | 0.0 | 0.0 |
| Candidatus Saccharibacteria oral taxon TM7x | | | 0.0 | 0.0 |
| Geitlerinema sp. PCC 7407 | Geitlerinema | | 0.0 | 0.0 |
| * | Actinomyces | Actinomycetaceae | 0.0 | 0.0 |
| Brevundimonas vesicularis | Brevundimonas | Caulobacteraceae | 0.0 | 0.0 |
| Acinetobacter sp. YS0810 | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| * | Prevotella | Prevotellaceae | 0.0 | 0.0 |
| Methyloceanibacter caenitepidi | Methyloceanibacter | | 0.0 | 0.0 |
| Leuconostoc citreum | Leuconostoc | Leuconostocaceae | 0.0 | 0.0 |
| * | Bacteroides | Bacteroidaceae | 0.0 | 0.0 |
| Pseudomonas alcaligenes | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Methylibium petroleiphilum | Methylibium | | 0.0 | 0.0 |
| Moraxella catarrhalis | Moraxella | Moraxellaceae | 0.0 | 0.0 |
| Sphingopyxis sp. Kp5.2 | Sphingopyxis | Sphingomonadaceae | 0.0 | 0.0 |
| Pandoraea apista | Pandoraea | Burkholderiaceae | 0.0 | 0.0 |
| * | Cellulomonas | Cellulomonadaceae | 0.0 | 0.0 |
| Olsenella uli | Olsenella | Coriobacteriaceae | 0.0 | 0.0 |
| Acinetobacter oleivorans | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| Sphingomonas sp. 133 | Sphingomonas | Sphingomonadaceae | 0.0 | 0.0 |
| Meiothermus taiwanensis | Meiothermus | Thermaceae | 0.0 | 0.0 |
| Deinococcus geothermalis | Deinococcus | Deinococcaceae | 0.0 | 0.0 |
| Lactobacillus sanfranciscensis | Lactobacillus | Lactobacillaceae | 0.0 | 0.0 |
| Acinetobacter sp. LUH5605 | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| Staphylococcus simulans | Staphylococcus | Staphylococcaceae | 0.0 | 0.0 |
| Arsenophonus nasoniae | Arsenophonus | Enterobacteriaceae | 0.0 | 0.0 |
| Buchnera aphidicola | Buchnera | Enterobacteriaceae | 0.0 | 0.0 |
| Weissella koreensis | Weissella | Leuconostocaceae | 0.0 | 0.0 |

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|-------------------------------|------------------|---------------------|-----|-----|
| Psychrobacter sp. G | Psychrobacter | Moraxellaceae | 0.0 | 0.0 |
| Amycolobicoccus subflavus | Amycolobicoccus | Mycobacteriaceae | 0.0 | 0.0 |
| Staphylococcus hyicus | Staphylococcus | Staphylococcaceae | 0.0 | 0.0 |
| Morganella morganii | Morganella | Enterobacteriaceae | 0.0 | 0.0 |
| Kyrpidia tusciae | Kyrpidia | Alicyclobacillaceae | 0.0 | 0.0 |
| Ramlibacter tataouinensis | Ramlibacter | Comamonadaceae | 0.0 | 0.0 |
| Weeksella virosa | Weeksella | Flavobacteriaceae | 0.0 | 0.0 |
| Acinetobacter junii | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| Acinetobacter sp. 26B2 | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| Mycobacterium abscessus | Mycobacterium | Mycobacteriaceae | 0.0 | 0.0 |
| Neisseria gonorrhoeae | Neisseria | Neisseriaceae | 0.0 | 0.0 |
| Sphingomonas wittichii | Sphingomonas | Sphingomonadaceae | 0.0 | 0.0 |
| Bacteroides dorei | Bacteroides | Bacteroidaceae | 0.0 | 0.0 |
| Corynebacterium halotolerans | Corynebacterium | Corynebacteriaceae | 0.0 | 0.0 |
| Pelomonas aquatica | Pelomonas | Comamonadaceae | 0.0 | 0.0 |
| Janibacter sp. TYM3221 | Janibacter | Intrasporangiaceae | 0.0 | 0.0 |
| * | Arthrobacter | Micrococcaceae | 0.0 | 0.0 |
| Mycobacterium gordonae | Mycobacterium | Mycobacteriaceae | 0.0 | 0.0 |
| Pimelobacter simplex | Pimelobacter | Nocardoidaceae | 0.0 | 0.0 |
| Pseudomonas sp. OM2164 | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Streptomyces albus | Streptomyces | Streptomycetaceae | 0.0 | 0.0 |
| Halomonas halocynthiae | Halomonas | Halomonadaceae | 0.0 | 0.0 |
| Nitrosomonas sp. AL212 | Nitrosomonas | Nitrosomonadaceae | 0.0 | 0.0 |
| Sphingobacterium mizutaii | Sphingobacterium | Sphingobacteriaceae | 0.0 | 0.0 |
| Vibrio cholerae | Vibrio | Vibrionaceae | 0.0 | 0.0 |
| Rickettsia felis | Rickettsia | Rickettsiaceae | 0.0 | 0.0 |
| * | * | Moraxellaceae | 0.0 | 0.0 |
| Bacteroidetes bacterium WD317 | | | 0.0 | 0.0 |
| Corynebacterium casei | Corynebacterium | Corynebacteriaceae | 0.0 | 0.0 |
| Corynebacterium marinum | Corynebacterium | Corynebacteriaceae | 0.0 | 0.0 |
| * | * | Flavobacteriaceae | 0.0 | 0.0 |
| Caulobacter segnis | Caulobacter | Caulobacteraceae | 0.0 | 0.0 |
| Lactobacillus gasseri | Lactobacillus | Lactobacillaceae | 0.0 | 0.0 |
| * | Meiothermus | Thermaceae | 0.0 | 0.0 |
| Rhizobium sp. NT-26 | Rhizobium | Rhizobiaceae | 0.0 | 0.0 |
| Bacillus coagulans | Bacillus | Bacillaceae | 0.0 | 0.0 |
| * | Sphingomonas | Sphingomonadaceae | 0.0 | 0.0 |
| * | Brevibacterium | Brevibacteriaceae | 0.0 | 0.0 |
| Nitrosomonas europaea | Nitrosomonas | Nitrosomonadaceae | 0.0 | 0.0 |
| Pseudomonas alkylphenolia | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Terrabacter sp. DBF63 | Terrabacter | Intrasporangiaceae | 0.0 | 0.0 |

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|--------------------------------------|------------------|---------------------|-----|-----|
| beta proteobacterium CB | | | 0.0 | 0.0 |
| Moraxella ovis | Moraxella | Moraxellaceae | 0.0 | 0.0 |
| Shewanella baltica | Shewanella | Shewanellaceae | 0.0 | 0.0 |
| Mycobacterium gilvum | Mycobacterium | Mycobacteriaceae | 0.0 | 0.0 |
| * | Exiguobacterium | | 0.0 | 0.0 |
| * | Ochrobactrum | Brucellaceae | 0.0 | 0.0 |
| Geodermatophilus obscurus | Geodermatophilus | Geodermatophilaceae | 0.0 | 0.0 |
| * | Devosia | Hyphomicrobiaceae | 0.0 | 0.0 |
| Moraxella osloensis | Moraxella | Moraxellaceae | 0.0 | 0.0 |
| Exiguobacterium sp. 11-28 | Exiguobacterium | | 0.0 | 0.0 |
| Nocardioides sp. JS614 | Nocardioides | Nocardioidaceae | 0.0 | 0.0 |
| Nocardioides sp. USM2 | Nocardioides | Nocardioidaceae | 0.0 | 0.0 |
| Burkholderia gladioli | Burkholderia | Burkholderiaceae | 0.0 | 0.0 |
| Renibacterium salmoninarum | Renibacterium | Micrococcaceae | 0.0 | 0.0 |
| Pseudomonas syringae group genom | Pseudomonas | Pseudomonadaceae | 0.0 | 0.0 |
| Bifidobacterium pseudolongum | Bifidobacterium | Bifidobacteriaceae | 0.0 | 0.0 |
| toluene-degrading bacterium UCR 021t | | | 0.0 | 0.0 |
| Corynebacterium imitans | Corynebacterium | Corynebacteriaceae | 0.0 | 0.0 |
| Corynebacterium callunae | Corynebacterium | Corynebacteriaceae | 0.0 | 0.0 |
| Bosea sp. WAO | Bosea | Bradyrhizobiaceae | 0.0 | 0.0 |
| Xanthobacter autotrophicus | Xanthobacter | Xanthobacteraceae | 0.0 | 0.0 |
| Corynebacterium diphtheriae | Corynebacterium | Corynebacteriaceae | 0.0 | 0.0 |
| Bacteroides thetaiotaomicron | Bacteroides | Bacteroidaceae | 0.0 | 0.0 |
| Caulobacter vibrioides | Caulobacter | Caulobacteraceae | 0.0 | 0.0 |
| * | * | Pasteurellaceae | 0.0 | 0.0 |
| Fingoldia magna | Fingoldia | Peptoniphilaceae | 0.0 | 0.0 |
| Anaerococcus prevotii | Anaerococcus | Peptoniphilaceae | 0.0 | 0.0 |
| Azorhizobium caulinodans | Azorhizobium | Xanthobacteraceae | 0.0 | 0.0 |
| Erysipelothrix rhusiopathiae | Erysipelothrix | Erysipelotrichaceae | 0.0 | 0.0 |
| Porphyromonas asaccharolytica | Porphyromonas | Porphyromonadaceae | 0.0 | 0.0 |
| * | Sphingopyxis | Sphingomonadaceae | 0.0 | 0.0 |
| Eubacterium rectale | Eubacterium | Eubacteriaceae | 0.0 | 0.0 |
| Acinetobacter venetianus | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| Variovorax paradoxus | Variovorax | Comamonadaceae | 0.0 | 0.0 |
| Acinetobacter sp. ED45-25 | Acinetobacter | Moraxellaceae | 0.0 | 0.0 |
| Bradyrhizobium diazoefficiens | Bradyrhizobium | Bradyrhizobiaceae | 0.0 | 0.0 |
| Bradyrhizobium japonicum | Bradyrhizobium | Bradyrhizobiaceae | 0.0 | 0.0 |
| Megamonas hypermegale | Megamonas | Veillonellaceae | 0.0 | 0.0 |
| * | Methylobacillus | Methylophilaceae | 0.0 | 0.0 |
| Nocardiopsis alba | Nocardiopsis | Nocardiopsaceae | 0.0 | 0.0 |
| Modestobacter marinus | Modestobacter | Geodermatophilaceae | 0.0 | 0.0 |

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|-------------------------------------|-------------------------|---------------------|-----|-----|
| <i>Corynebacterium doosanense</i> | <i>Corynebacterium</i> | Corynebacteriaceae | 0.0 | 0.0 |
| <i>Blastococcus saxobidens</i> | <i>Blastococcus</i> | Geodermatophilaceae | 0.0 | 0.0 |
| <i>Anoxybacillus flavithermus</i> | <i>Anoxybacillus</i> | Bacillaceae | 0.0 | 0.0 |
| <i>Aeromonas caviae</i> | <i>Aeromonas</i> | Aeromonadaceae | 0.0 | 0.0 |
| <i>Bacillus subtilis</i> | <i>Bacillus</i> | Bacillaceae | 0.0 | 0.0 |
| <i>Elizabethkingia anophelis</i> | <i>Elizabethkingia</i> | Flavobacteriaceae | 0.0 | 0.0 |
| <i>Staphylococcus hominis</i> | <i>Staphylococcus</i> | Staphylococcaceae | 0.0 | 0.0 |
| <i>Ruminococcus bicirculans</i> | <i>Ruminococcus</i> | Ruminococcaceae | 0.0 | 0.0 |
| <i>Paracoccus marcusii</i> | <i>Paracoccus</i> | Rhodobacteraceae | 0.0 | 0.0 |
| * | <i>Psychrobacter</i> | Moraxellaceae | 0.0 | 0.0 |
| <i>Sphingomonas sanxanigenens</i> | <i>Sphingomonas</i> | Sphingomonadaceae | 0.0 | 0.0 |
| * | * | Leuconostocaceae | 0.0 | 0.0 |
| * | * | Burkholderiaceae | 0.0 | 0.0 |
| <i>Bacteroides vulgatus</i> | <i>Bacteroides</i> | Bacteroidaceae | 0.0 | 0.0 |
| <i>Rhodopseudomonas palustris</i> | <i>Rhodopseudomonas</i> | Bradyrhizobiaceae | 0.0 | 0.0 |
| <i>Pantoea agglomerans</i> | <i>Pantoea</i> | Enterobacteriaceae | 0.0 | 0.0 |
| * | * | Sphingomonadaceae | 0.0 | 0.0 |
| <i>Mycobacterium kansasii</i> | <i>Mycobacterium</i> | Mycobacteriaceae | 0.0 | 0.0 |
| * | <i>Streptomyces</i> | Streptomycetaceae | 0.0 | 0.0 |
| <i>Enterococcus faecalis</i> | <i>Enterococcus</i> | Enterococcaceae | 0.0 | 0.0 |
| <i>Acinetobacter</i> sp. NFM2 | <i>Acinetobacter</i> | Moraxellaceae | 0.0 | 0.0 |
| <i>Shewanella putrefaciens</i> | <i>Shewanella</i> | Shewanellaceae | 0.0 | 0.0 |
| <i>Bifidobacterium adolescentis</i> | <i>Bifidobacterium</i> | Bifidobacteriaceae | 0.0 | 0.0 |
| * | <i>Bifidobacterium</i> | Bifidobacteriaceae | 0.0 | 0.0 |
| <i>Porphyromonas gingivalis</i> | <i>Porphyromonas</i> | Porphyromonadaceae | 0.0 | 0.0 |
| <i>Neisseria meningitidis</i> | <i>Neisseria</i> | Neisseriaceae | 0.0 | 0.0 |
| <i>Rhodococcus pyridinivorans</i> | <i>Rhodococcus</i> | Nocardiaceae | 0.0 | 0.0 |
| <i>Aeromonas salmonicida</i> | <i>Aeromonas</i> | Aeromonadaceae | 0.0 | 0.0 |
| <i>Planococcus</i> sp. PAMC 21323 | <i>Planococcus</i> | Planococcaceae | 0.0 | 0.0 |
| <i>Pseudomonas simiae</i> | <i>Pseudomonas</i> | Pseudomonadaceae | 0.0 | 0.0 |
| <i>Faecalibacterium prausnitzii</i> | <i>Faecalibacterium</i> | Ruminococcaceae | 0.0 | 0.0 |
| <i>Acinetobacter lwoffii</i> | <i>Acinetobacter</i> | Moraxellaceae | 0.0 | 0.0 |
| <i>Exiguobacterium</i> sp. N139 | <i>Exiguobacterium</i> | | 0.0 | 0.0 |
| <i>Streptococcus anginosus</i> | <i>Streptococcus</i> | Streptococcaceae | 0.0 | 0.0 |
| <i>Thauera</i> sp. MZ1T | <i>Thauera</i> | Rhodocyclaceae | 0.0 | 0.0 |
| * | <i>Shewanella</i> | Shewanellaceae | 0.0 | 0.0 |
| * | <i>Aeromonas</i> | Aeromonadaceae | 0.0 | 0.0 |
| <i>Staphylococcus aureus</i> | <i>Staphylococcus</i> | Staphylococcaceae | 0.0 | 0.0 |
| <i>Aeromonas hydrophila</i> | <i>Aeromonas</i> | Aeromonadaceae | 0.0 | 0.0 |
| <i>Aeromonas veronii</i> | <i>Aeromonas</i> | Aeromonadaceae | 0.0 | 0.0 |
| <i>Bifidobacterium breve</i> | <i>Bifidobacterium</i> | Bifidobacteriaceae | 0.0 | 0.0 |

| | | | | |
|---------------------|----------|-------------|-----|-----|
| Bacillus megaterium | Bacillus | Bacillaceae | 0.0 | 0.0 |
|---------------------|----------|-------------|-----|-----|