



- Arthrospira platensis\_nov\_88.249%
- Sphingomonas lutea
- Arthrospira platensis\_nov\_89.425%
- Rubellimicrobium mesophilum\_nov\_95.592%
- Sphingomonas zeicaulis
- Sporichthya polymorpha\_nov\_93.305%
- Arthrospira platensis\_nov\_89.041%
- Goodfellowiella coeruleoviolacea\_nov\_94.612%
- Oscillochloris trichoides\_nov\_85.202%
- Pseudonocardia profundimaris
- Kosakonia sacchari
- Lactiplantibacillus\_Lactobacillus pentosus\_plantarum
- Romboutsia timonensis
- Blautia faecicola\_nov\_93.111%
- Gloeobacter multispecies\_sppn35\_2\_nov\_92.361%
- Gordonia soli
- Kocuria himachalensis\_rosea
- Corynebacterium mycetoides\_nov\_97.629%
- Corynebacterium tuberculostearicum
- Actinidia eriantha\_nov\_97.297%
- Atopostipes sp.\_MOT-200
- Rothia terrae
- Anaerosacchariphilus polymeriproducens\_nov\_95.218%
- Staphylococcus hominis
- Micrococcus aloeverae
- Staphylococcus haemolyticus
- Veillonella parvula
- Chryseobacterium culicis\_nov\_97.280%
- Lactiplantibacillus plantarum
- Blautia multispecies\_sppn28\_2\_nov\_95.198%
- Streptococcus mitis\_sp.\_HMT\_064
- Eubacterium coprostanoligenes\_nov\_92.424%
- Shigella flexneri
- Shigella sonnei
- Escherichia coli
- Shigella dysenteriae
- Roseomonas aquatica\_nov\_97.912%
- Dyadobacter sediminis\_nov\_97.240%
- Streptococcus sp.\_HMT\_064
- Schlegella brevitalia\_nov\_97.510%
- Microbacterium aurum
- Corynebacterium amycolatum
- Cutibacterium acnes
- Blautia luti\_nov\_96.429%
- Fusicatenibacter saccharivorans\_nov\_97.689%
- Phocaeicola vulgatus
- Staphylococcus saprophyticus\_xylosus
- Sphingomonas aurantiaca\_faeni
- Mammaliococcus sciuri
- Arthrospira platensis\_nov\_87.838%
- Arthrospira platensis\_nov\_88.713%
- Roseiflexus castenholzii\_nov\_84.842%
- Arthrospira platensis\_nov\_90.092%
- Arthrospira platensis\_nov\_90.138%
- Arthrospira platensis\_nov\_89.450%
- Sphingomonas suaedae\_nov\_96.970%
- Aggregatilinea lenta\_nov\_77.847%
- Arthrospira platensis\_nov\_87.963%
- Microlunatus aurantiacus\_nov\_92.903%
- Arthrospira platensis\_nov\_88.046%
- Craurococcus roseus\_nov\_95.602%
- Microvirga subterranea\_nov\_95.150%
- Arthrospira platensis\_nov\_84.510%
- Arthrospira platensis\_nov\_88.073%
- Roseiflexus castenholzii\_nov\_84.876%
- Arthrospira platensis\_nov\_86.605%
- Rubellimicrobium mesophilum\_nov\_94.919%
- Enterococcus faecalis
- Gordonia sediminis
- Paracoccus marinus
- Rubellimicrobium mesophilum
- Sphingomonas leidyi
- Amaricoccus tamworthensis\_nov\_97.448%
- Paracraurococcus ruber\_nov\_97.448%
- Arboricoccus pini\_nov\_88.276%
- Gordonia soli\_nov\_97.198%
- Salmonella enterica
- Granulicatella adiacens
- Collinsella aerofaciens
- Haemophilus parainfluenzae
- Veillonella dispar\_parvula
- Staphylococcus warneri
- Staphylococcus capitis\_epidermidis
- Citrobacter koseri
- Amaricoccus kaplicensis\_nov\_95.833%
- Staphylococcus argenteus\_aureus\_rotterdamii
- Romboutsia ilealis
- Arthrospira platensis\_nov\_90.805%
- Arthrospira platensis\_nov\_88.736%
- Streptococcus agalactiae
- Bifidobacterium pseudolongum
- Modestobacter marinus
- Ligilactobacillus murinus
- Streptococcus parasanguinis\_parasanguinis\_clade\_721
- Lactobacillus intestinalis
- Paracoccus carotinifaciens\_haeundaensis\_marcusii
- Staphylococcus ureilyticus
- Streptococcus mutans
- Rothia nasimurium\_nov\_97.228%
- Streptococcus sp.\_MOT-045\_nov\_97.764%

Species



F18249.S17  
F18249.S18  
F18249.S19  
F18249.S20  
F18249.S21  
F18249.S22  
F18249.S23  
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F18249.S25  
F18249.S26  
F18249.S27  
F18249.S28  
F18249.S29  
F18249.S30  
F18249.S31  
F18249.S32

Samples