



- Saccharibacteria_(TM7)_[F-1]
- Actinomycetaceae
- Pasteurellaceae
- Bacteroidales_[F-2]
- Micrococcaceae
- Barnesiellaceae
- Veillonellaceae
- Odoribacteraceae
- Methanobacteriaceae
- Porphyromonadaceae
- Rikenellaceae
- Micrococcaceae
- Lactobacillaceae
- Eubacteriaceae_[XV]
- Peptostreptococcaceae
- Prevotellaceae
- Desulfovibrionaceae
- Gemellaceae
- Lachnospiraceae_[XIVa]
- Erysipelotrichaceae
- Acidaminococcaceae
- Akkermansiaceae
- Bacteroidaceae
- Sutterellaceae
- Ruminococcaceae
- Oscillospiraceae
- Coriobacteriaceae
- Lachnospiraceae
- Bifidobacteriaceae
- Streptococcaceae
- Clostridiales_[F-1]
- Lachnospiraceae_[XIV]
- Eubacteriaceae
- Anaeroplasmataceae
- Succinivibrionaceae
- Muribaculaceae
- Veillonellaceae
- Erysipelotrichaceae
- Neisseriaceae
- Carnobacteriaceae
- Bacteroidaceae
- Bifidobacteriaceae
- Enterobacteriaceae
- Lachnospiraceae_Lachnospiraceae_[XIVa]
- Alcaligenaceae
- Veillonellaceae
- Lachnospiraceae
- Christensenellaceae
- Clostridiaceae
- multifamily

F6488.S52
F6488.S53
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F6488.S58
F6488.S59
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F6488.S78
F6488.S79

Samples

Family