



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

Family

- F0488.S52
- F0488.S53
- F0488.S54
- F0488.S55
- F0488.S56
- F0488.S57
- F0488.S58
- F0488.S59
- F0488.S60
- F0488.S61
- F0488.S72
- F0488.S73
- F0488.S74
- F0488.S75
- F0488.S76
- F0488.S78
- F0488.S79

Samples