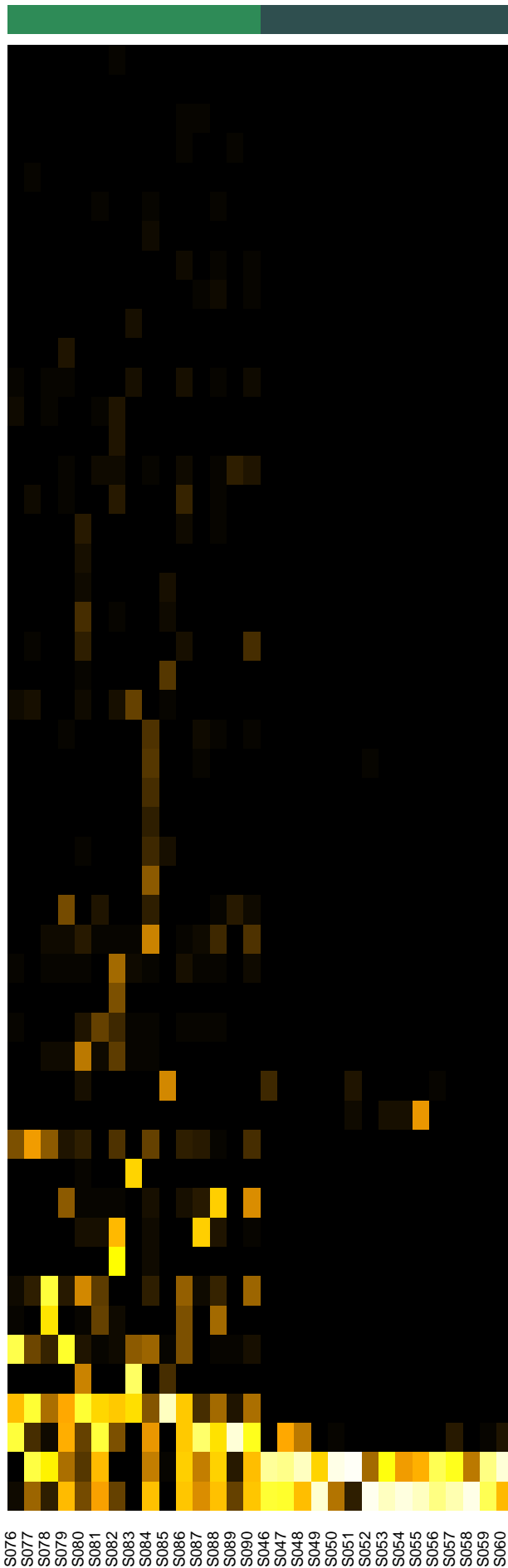




Description2
 PT_VP_PP
 PT_NP_PP



- Saccharibacteria_(TM7)_[F-2]
- Enterobacteriaceae
- Lachnospiraceae
- Clostridiaceae
- multifamily
- Anaerolineaceae
- Prevotellaceae
- Bacteroidetes_[F-1]
- Bacteroidales_[F-2]
- Burkholderiaceae
- Gemellaceae
- Aerococcaceae
- Veillonellaceae
- Bifidobacteriaceae
- Veillonellaceae
- Selenomonadaceae
- Peptostreptococcaceae_Peptostreptococcaceae_[XI]
- Corynebacteriaceae
- Moraxellaceae_Neisseriaceae
- Pseudomonadaceae
- Tissierellaceae
- Caulobacteraceae
- Carnobacteriaceae
- Treponemataceae
- Eubacteriaceae
- Blastocatellaceae
- Xanthobacteraceae
- Sphingomonadaceae
- Acidobacteriaceae
- Synergistaceae
- Actinomycetaceae
- Staphylococcaceae
- Lactobacillaceae
- Actinomycetaceae
- Micrococcaceae
- Rhodobacteraceae
- Campylobacteraceae
- Lachnospiraceae_[XIV]
- Neisseriaceae
- Porphyromonadaceae
- Saccharibacteria_(TM7)_[F-1]
- Atopobiaceae
- Fusobacteriaceae
- Peptoniphilaceae_Peptostreptococcaceae_[XI]
- Enterococcaceae
- Pasteurellaceae
- Streptococcaceae
- Desulfobulbaceae
- Peptoniphilaceae
- Peptostreptococcaceae_[XI]

Family

S076 S077 S078 S079 S080 S081 S082 S083 S084 S085 S086 S087 S088 S089 S090 S046 S047 S048 S049 S050 S051 S052 S053 S054 S055 S056 S057 S058 S059 S060

Samples