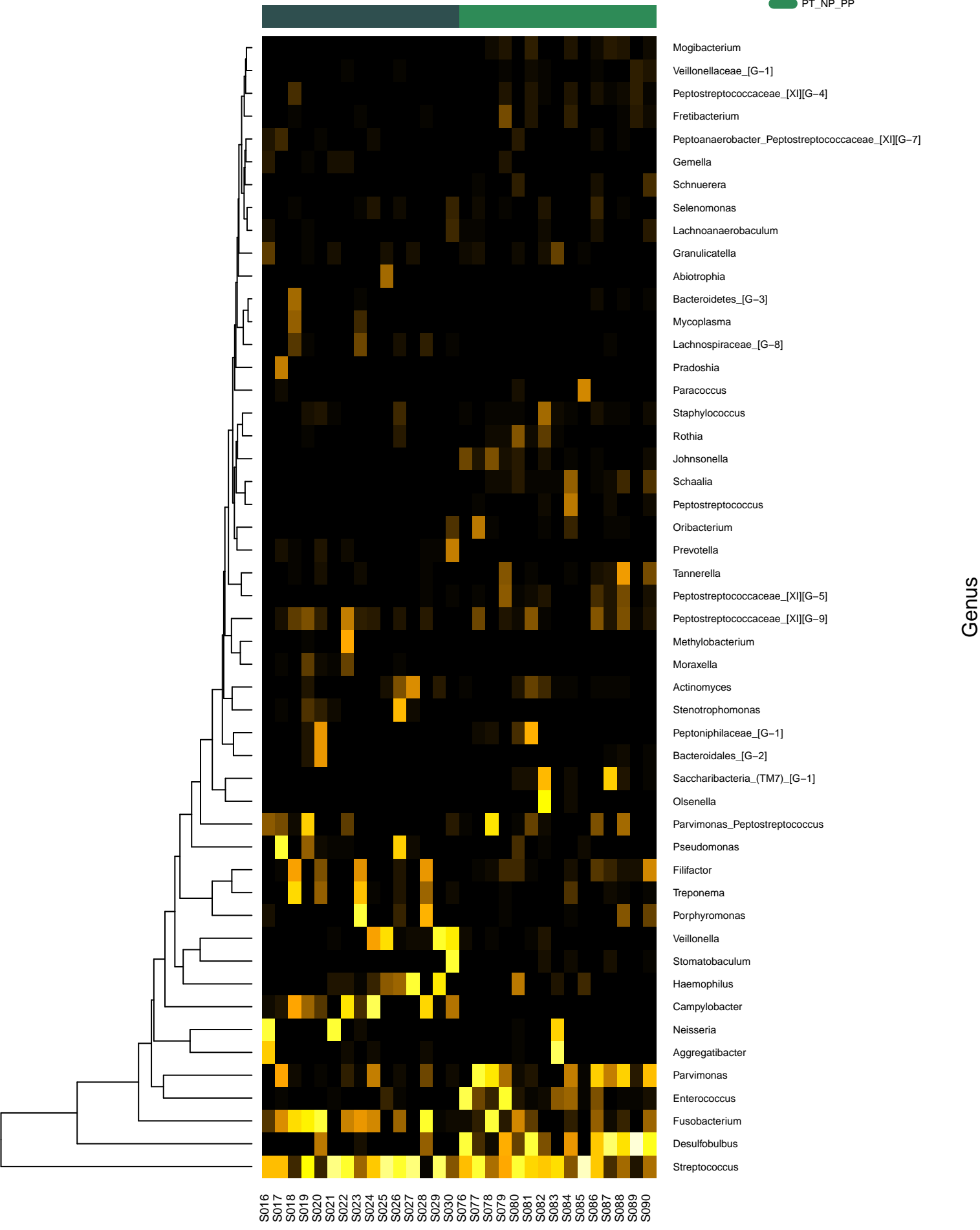




Description2
 PT_NP_HPT
 PT_NP_PP



- Mogibacterium
- Veillonellaceae_[G-1]
- Peptostreptococcaceae_[XI][G-4]
- Fretibacterium
- Peptoanaerobacter_Peptostreptococcaceae_[XI][G-7]
- Gemella
- Schnuerera
- Selenomonas
- Lachnoanaerobaculum
- Granulicatella
- Abiotrophia
- Bacteroidetes_[G-3]
- Mycoplasma
- Lachnospiraceae_[G-8]
- Pradoshia
- Paracoccus
- Staphylococcus
- Rothia
- Johnsonella
- Schaalia
- Peptostreptococcus
- Oribacterium
- Prevotella
- Tannerella
- Peptostreptococcaceae_[XI][G-5]
- Peptostreptococcaceae_[XI][G-9]
- Methylobacterium
- Moraxella
- Actinomyces
- Stenotrophomonas
- Peptoniphilaceae_[G-1]
- Bacteroidales_[G-2]
- Saccharibacteria_(TM7)_[G-1]
- Olsenella
- Parvimonas_Peptostreptococcus
- Pseudomonas
- Filifactor
- Treponema
- Porphyromonas
- Veillonella
- Stomatobaculum
- Haemophilus
- Campylobacter
- Neisseria
- Aggregatibacter
- Parvimonas
- Enterococcus
- Fusobacterium
- Desulfobulbus
- Streptococcus

S016 S017 S018 S019 S020 S021 S022 S023 S024 S025 S026 S027 S028 S029 S030 S076 S077 S078 S079 S080 S081 S082 S083 S084 S085 S086 S087 S088 S089 S090