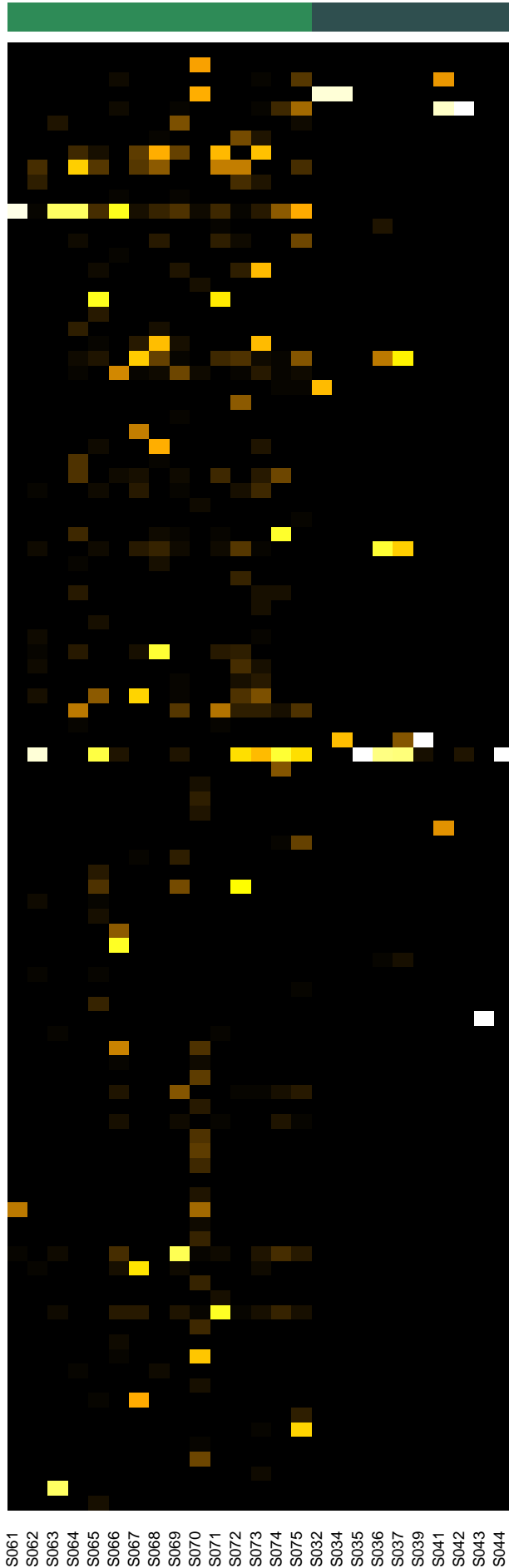




Description2  
 RC\_VP\_PP  
 RC\_NP\_PP



- Rothia mucilaginosa
- Micrococcus cohnii
- Peptostreptococcaceae\_[XII][G-5] [Eubacterium]\_saphenum
- Campylobacter gracilis
- Peptostreptococcaceae\_[XII][G-9] [Eubacterium]\_brachy
- Streptococcus sanguinis
- Olsenella ulii
- Bacteroidaceae\_[G-1] bacterium\_HMT\_272
- Stomatobaculum sp.\_HMT\_373
- Oribacterium sp.\_HMT\_078
- Desulfovibrio desulfuricans
- Enterococcus faecalis
- Peptostreptococcaceae\_[XII][G-6] [Eubacterium]\_nodatum
- Peptostreptococcaceae\_[XII][G-6] [Eubacterium]\_minutum
- Porphyromonas gingivalis
- Atopobium sp.\_HMT\_199
- Sphingomonas hankookensis
- Peptoniphilaceae\_[G-1] bacterium\_HMT\_113
- Atopobium sp.\_HMT\_416
- Fretibacterium sp.\_HMT\_360
- Erysipelotrichaceae\_[G-1] bacterium\_HMT\_904
- Mogibacterium timidum
- Schaalia odontolytica
- Peptostreptococcaceae\_[XII][G-4] bacterium\_HMT\_369
- Prevotella sp.\_HMT\_376
- Enterobacter mori
- Atopobium sp.\_HMT\_810
- Phocaeicola abscessus
- Anaeroglobus geminatus
- Desulfobulbus sp.\_HMT\_041
- Dialister invisus
- Capnocytophaga sputigena
- Granulicatella adiacens
- Fretibacterium fastidiosum
- Pseudoramibacter alactolyticus
- Anaerolineae\_[G-1] bacterium\_HMT\_439
- Erysipelotrichaceae\_[G-1] bacterium\_HMT\_905
- Tannerella forsythia
- Porphyromonas endodontalis
- Bulleidia extracta
- Peptostreptococcaceae\_[XII][G-1] [Eubacterium]\_infirmum
- Peptostreptococcaceae\_[XII][G-1] bacterium\_HMT\_383
- Solobacterium moorei
- Dialister pneumosintes
- Peptostreptococcus stomatis
- Filifactor aloccis
- Peptostreptococcaceae\_[XII][G-2] bacterium\_HMT\_091
- Paracoccus speluncae
- Parvimonas micra
- Methylobacterium radiotolerans
- Brevundimonas bacteroides
- Sphingomonas aestuarii
- Alishewanella alkalitolerans
- Peptostreptococcaceae\_[XII][G-4] bacterium\_HMT\_103
- Streptococcus gordonii
- Oribacterium sp.\_HMT\_102
- Butyrivibrio sp.\_HMT\_090\_nov\_95.724%
- Oribacterium sp.\_HMT\_078\_nov\_97.537%
- Stomatobaculum sp.\_HMT\_373\_nov\_94.581%
- Peptococcus simiae\_nov\_92.326%
- Caedimonas varicaedens\_nov\_93.069%
- Pradoshia eiseniae\_nov\_93.764%
- Parvimonas micra\_nov\_93.971%
- Peptostreptococcus stomatis\_nov\_95.037%
- Schnuerera ultunensis\_nov\_92.840%
- Peptoniphilaceae\_[G-1] bacterium\_HMT\_113\_nov\_96.782%
- Aridibacter famidurans\_nov\_96.552%
- Enterococcus faecalis\_nov\_97.669%
- Paracoccus carotinifaciens\_hibiscisoli\_marcusii\_nototheniae
- Neobacillus cucumis\_drentensis\_novalis\_soli\_vireti
- Kocuria atrinae\_carniphila\_gwangalliensis
- Streptococcus cristatus\_downii\_gwangjuense\_infantis\_infantis\_cla
- Acinetobacter\_Prolinoborus fasciculus\_lwoffii
- Streptococcus salivarius\_vestibularis
- Chryseobacterium binzhouense\_echinoideorum
- Micrococcus antarcticus\_endophyticus\_luteus\_yunnanensis
- Blastomonas\_Sphingomonas natatoria\_ursincola
- Sphingomonas aquatilis\_melonis
- Janibacter\_Knoellia\_Pedococcus\_Phycococcus anophelis\_badiiscori
- Paracoccus aestuarii\_beibuensis\_hibisci\_marinus\_pueri
- Sphingomonas paucimobilis\_pseudosanguinis\_sanguinis\_yabuuchi
- Brevundimonas bullata\_halotolerans
- Aerococcus urinaequi\_viridans
- Parvimonas\_Peptostreptococcus Candidatus massiliensis\_sp.\_HMT
- Paracoccus aminovorans\_caeni\_chinensis\_huijuniae\_subflavus
- Staphylococcus capitis\_caprae
- Staphylococcus capitis\_caprae\_epidermidis
- Sphingomonas hankookensis\_panni
- Enterococcus casseliflavus\_gallinarum
- Pseudomonas alcaliphila\_chengduensis\_oleovorans\_toyotomiensis
- Veillonellaceae\_[G-1] bacterium\_HMT\_132\_bacterium\_HMT\_150
- Brachybacterium horti\_nesterenkovi\_rharnosum
- Olsenella phocaeensis\_sp.\_HMT\_809
- Streptococcus cristatus\_cristatus\_clade\_578
- Streptococcus anginosus\_constellatus
- Enterobacter asburiae\_cancerogenus\_cloacae\_hormaechei
- Brevundimonas albigilva\_nasdae\_vesicularis
- Fusobacterium nucleatum\_nucleatum\_subsp.\_vincentii
- multigenus multispecies\_sppn3\_5\_nov\_96.737%
- multigenus multispecies\_sppn6\_3\_nov\_96.535%

Species

Samples