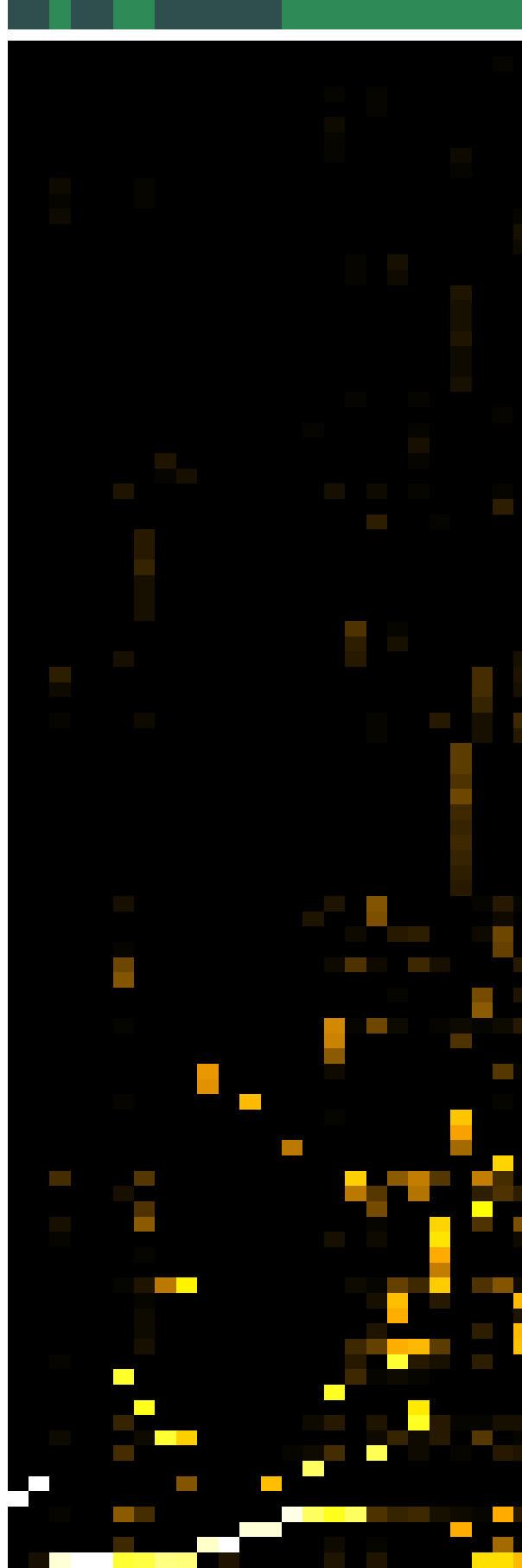


Description2
 RC_VP_PP
 RC_NP_PP



- Species
- Spingomonas aquatilis_melonis
 - Granulicatella adiacens
 - Rothia mucilaginos
 - Desulfovibrio desulfuricans
 - Enterobacter mori
 - Enterococcus casseliflavus_gallinarum
 - Porphyromonas gingivalis
 - Neobacillus cucumis_drentensis_novalis_soli_vireti
 - Enterobacter asburiae_cancerogenus_cloacae_hormaechei
 - Stomatobaculum sp._HMT_373_nov_94.581%
 - Peptostreptococcus stomatis_nov_95.037%
 - Peptostreptococcaceae_[X1][G-1] [Eubacterium]_infirmum
 - Porphyromonas endodontalis
 - Fusobacterium nucleatum_nucleatum_subsp._vincentii
 - Anaerolineae_[G-1] bacterium_HMT_439
 - Veillonellaceae_[G-1] bacterium_HMT_132_bacterium_HMT_150
 - Janibacter_Knoellia_Pedococcus_Phycococcus anophelis_badiiscori
 - Spingomonas hankookensis
 - Brevundimonas bacteroides
 - Alishewanella alkalitolerans
 - Spingomonas paucimobilis_pseudosanguinis_sanguinis_yabuuchi
 - Capnocytophaga sputigena
 - Brachybacterium horti_nesterenkovi_rhamnosum
 - Peptostreptococcaceae_[X1][G-2] bacterium_HMT_091
 - Schnuerera ultunensis_nov_92.840%
 - Enterococcus faecalis_nov_97.669%
 - Staphylococcus capitis_caprae
 - Peptostreptococcaceae_[X1][G-6] [Eubacterium]_nodatum
 - Parvimonas micra_nov_93.971%
 - Streptococcus salivarius_vestibularis
 - Streptococcus cristatus_cristatus_clade_578
 - Oribacterium sp._HMT_102
 - Atopobium sp._HMT_416
 - Butyrivibrio sp._HMT_090_nov_95.724%
 - Peptoniphilaceae_[G-1] bacterium_HMT_113_nov_96.782%
 - multigenus multispecies_sppn6_3_nov_96.535%
 - Peptococcus simiae_nov_92.326%
 - Bulleidia extracta
 - Anaeroglobus geminatus
 - Fretibacterium sp._HMT_360
 - Tannerella forsythia
 - Oribacterium sp._HMT_078
 - Solobacterium moorei
 - Erysipelotrichaceae_[G-1] bacterium_HMT_905
 - Dialister invisus
 - Dialister pneumosintes
 - Kocuria atrinae_carniphila_gwangalliensis
 - Micrococcus antarcticus_endophyticus_luteus_yunnanensis
 - Chryseobacterium binzhouense_echinoideorum
 - Brevundimonas albigilva_nasdae_vesicularis
 - Spingomonas hankookensis_panni
 - Paracoccus aminovorans_caeni_chinensis_huijuniae_subflavus
 - Blastomonas_Spingomonas natatoria_ursincola
 - Brevundimonas bullata_halotolerans
 - Spingomonas aestuarii
 - Acinetobacter_Prolinoborus fasciculus_lwoffii
 - Streptococcus cristatus_downii_gwangjuense_infantis_infantis_cla_.
 - Streptococcus sanguinis
 - Peptostreptococcaceae_[X1][G-6] [Eubacterium]_minutum
 - Streptococcus gordonii
 - Desulfobulbus sp._HMT_041
 - Methylobacterium radiotolerans
 - Olsenella uli
 - Prevotella sp._HMT_376
 - Schaalia odontolytica
 - Paracoccus carotinifaciens_hibiscisoli_marcusii_nototheniae
 - Caedimonas varicaedens_nov_93.069%
 - Peptostreptococcaceae_[X1][G-5] [Eubacterium]_saphenum
 - Peptostreptococcaceae_[X1][G-4] bacterium_HMT_103
 - Peptostreptococcaceae_[X1][G-4] bacterium_HMT_369
 - Pseudomonas alcaliphila_chengduensis_oleovorans_toyotomiensis
 - Micrococcus cohnii
 - Paracoccus aestuarii_beibuensis_hibisci_marinus_pueri
 - Streptococcus anginosus_constellatus
 - Stomatobaculum sp._HMT_373
 - Fillifactor alocis
 - Oribacterium sp._HMT_078_nov_97.537%
 - Peptostreptococcus stomatis
 - Parvimonas_Peptostreptococcus Candidatus massiliensis_sp._HMT
 - Olsenella phocaeensis_sp._HMT_809
 - Atopobium sp._HMT_810
 - Mogibacterium timidum
 - Erysipelotrichaceae_[G-1] bacterium_HMT_904
 - Phocaeicola abscessus
 - Atopobium sp._HMT_199
 - Bacteroidaceae_[G-1] bacterium_HMT_272
 - Peptostreptococcaceae_[X1][G-1] bacterium_HMT_383
 - Fretibacterium fastidiosum
 - Pradoshia eiseniae_nov_93.764%
 - Peptoniphilaceae_[G-1] bacterium_HMT_113
 - Staphylococcus capitis_caprae_epidermidis
 - Pseudoramibacter alactolyticus
 - Aerococcus urinaequi_viridans
 - multigenus multispecies_sppn3_5_nov_96.737%
 - Paracoccus speluncae
 - Aridibacter famidurans_nov_96.552%
 - Enterococcus faecalis
 - Aridibacter famidurans_nov_96.552%
 - Campylobacter gracilis
 - Peptostreptococcaceae_[X1][G-9] [Eubacterium]_brachy
 - Parvimonas micra

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Samples