s__Veillonella_rogosae s__Veillonella_dispar s_Neisseria_subflava _Leptotrichia_sp_oral_taxon_215 s_Prevotella_nigrescens s_GGB4937_SGB6891 s_s_LacherBangacene_Bangerium_oral_taxon_500 s_Alloprevotella_SGB1410 s_Solo Balcopticwotello spi_oral_taxon_473 s_Leptotriemay hassing SGB 96438 s_Capnocytophaga_sputigena s__GGB4533_SGB94822 s__Granulicatella_SGB8239 s__Sphingomonas_melonis s_Peptostreptococcus_SGB749 s_Capnocytophaga_SGB2480 s__GGB12441_SGB19290 s__GGB1088_SGB1395 s__Prevotella_melaninogenica s_Streptococcus_australis s__Capnocytophaga_leadbetteri s__Leptotrichia_wadei s__Leptotrichia_SGB6069 s_Eubacterium_sulci s_Prevotelle atypica elenomonas_SGB5880 s__Aggregatibacter_segnis s_Campylobacter_concisus s_GGB1 44_SGB1468 Cluster |r-value| Correlation (±) 0.2 positive 1 **2** 0.4 negative 3 0.6

s_Streptococcus_parasanguinis

s__Rothia_mucilaginosa