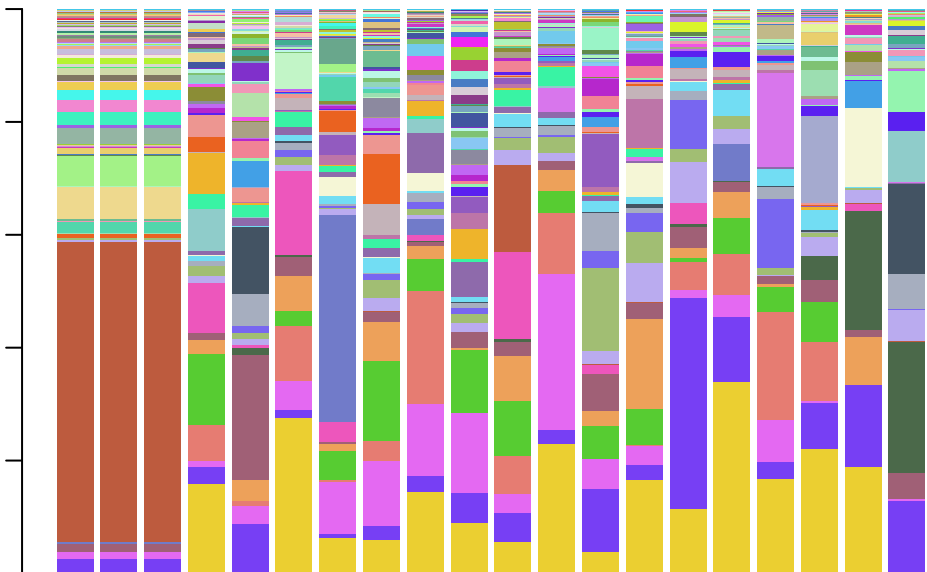
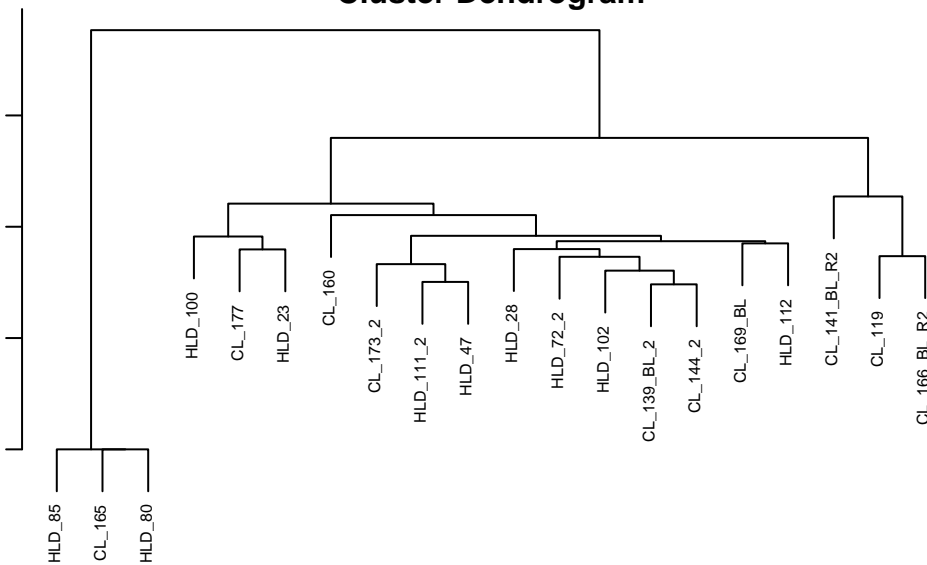
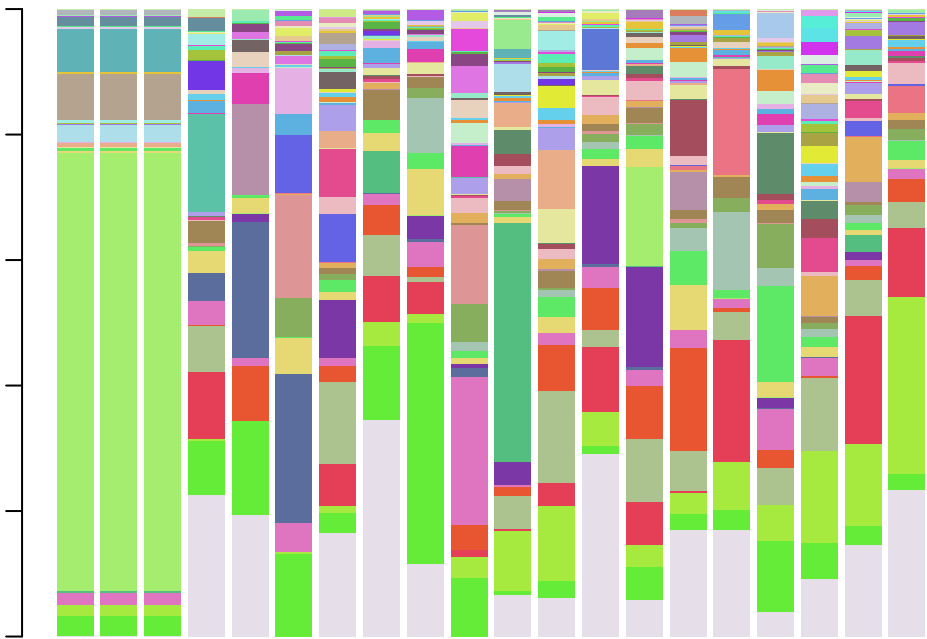
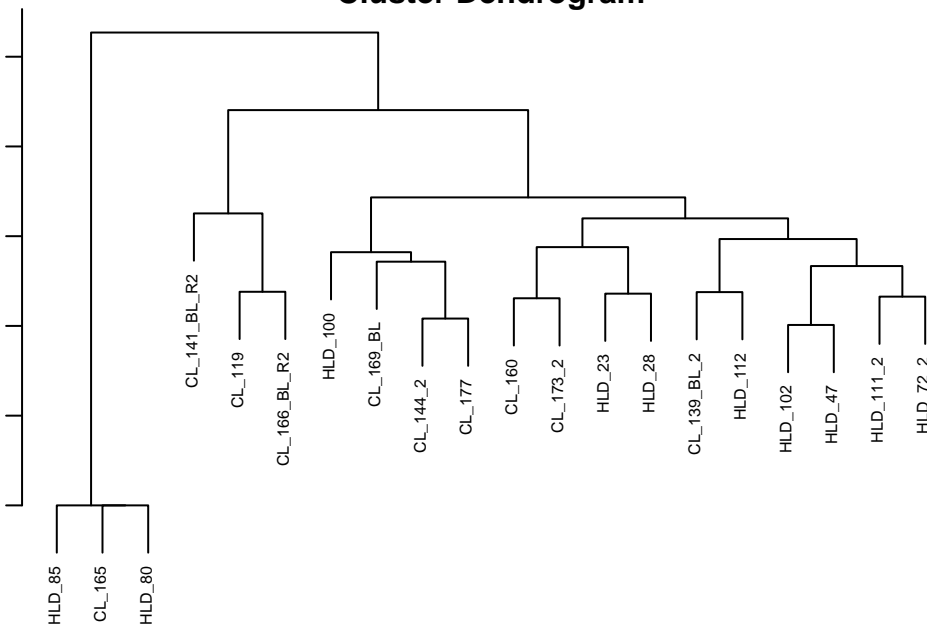


Cluster Dendrogram



- :lostridium_nexile
- achnospiraceae_bacterium_3_1_57F/
- :lostridium_hatthewayi
- acteroides_finegoldii
- arabacteroides_johnsonii
- :lostridium_hiranonis
- oldemania_sp_AP2
- treptococcus_australis
- actobacillus_casei_paracasei
- legamonas_unclassified
- arabacteroides_goldsteinii
- ggerthella_sp_HGA1
- :lostridium_asparagiforme
- uricbacter_unclassified
- seudomonas_unclassified
- uminococcus_flavefaciens
- ubacterium_dolichum
- achnospiraceae_bacterium_9_1_43B/
- treptococcus_anginosus
- :uminococcaceae_bacterium_D16
- ifidobacterium_animalis
- :lostridium_scindens
- naerotruncus_colihominis
- :lostridiales_bacterium_1_7_47FAA
- oseburia_unclassified
- actobacillus_curvatus
- acteroides_intestinalis
- naerotruncus_unclassified
- naerostipes_sp_3_2_56FAA
- eillonella_parvula
- lethanobrevibacter_smithii
- lebsiella_oxytoca
- naerostipes_unclassified
- :lostridium_perfringens
- ilophila_wadsworthia
- :eissella_halotolerans
- lautia_producta
- rysiplotrichaceae_bacterium_2_2_4_
- acteroides_clarus
- ifidobacterium_dentium
- treptococcus_sanguinis
- othia_mucilaginoso
- seudomonas_sp_UK4
- actobacillus_fermentum
- acteroides_cellulosilyticus
- actobacillus_plantarum
- accharomyces_cerevisiae
- achnospiraceae_bacterium_5_1_57F/
- rysiplotrichaceae_bacterium_3_1_5_
- oldemania_unclassified
- actobacillus_sakei
- utyricoccus_pullicaeorum
- :lostridium_bartlettii
- cinetobacter_unclassified
- treptococcus_mitis_oralis_pneumoniae
- eillonella_atypica
- acteroides_salyersiae
- ctinomyces_odontolyticus
- lebsiella_variicola
- treptococcus_infantis
- rysiplotrichaceae_bacterium_21_3_
- ctinomyces_turicensis
- :lostridiaceae_bacterium_JC118
- rysiplotrichaceae_bacterium_5_2_5_
- :itrobacter_unclassified
- legamonas_hypermegale
- treptococcus_sobrinus
- nterobacter_cloacae

Cluster Dendrogram



s__Faecalibacterium_prausnitzii
 s__Dorea_longicatena
 s__Ruminococcus_torques
 s__Ruminococcus_gnavus
 s__Akkermansia_muciniphila
 s__Coprococcus_sp_ART55_1
 s__Clostridium_sp_L2_50
 s__Eubacterium_hallii
 s__Ruminococcus_obeum
 s__Roseburia_intestinalis
 s__Coprococcus_comes
 s__Escherichia_coli
 s__Dorea_formicigenerans
 s__Bifidobacterium_adolescentis
 s__Bacteroides_uniformis
 s__Coprococcus_eutactus
 s__Dialister_invisus
 s__Collinsella_aerofaciens
 s__Alistipes_putredinis
 s__Roseburia_hominis
 s__Eubacterium_siraeum
 s__Roseburia_inulinivorans
 s__Prevotella_copri
 s__Bacteroides_vulgatus
 s__Lactobacillus_mucosae
 s__Lachnospiraceae_bacterium_1_1_
 s__Eubacterium_ramulus
 s__Clostridium_bolteae
 s__Streptococcus_thermophilus
 s__Coprococcus_catus
 s__Streptococcus_salivarius
 s__Eggerthella_unclassified
 s__Parabacteroides_merdae
 s__Bifidobacterium_longum
 s__Bacteroides_stercoris
 s__Clostridium_clostridioforme
 s__Raoultella_ornithinolytica
 s__Bacteroides_coprocola
 s__Lactobacillus_ruminis
 s__Lachnospiraceae_bacterium_3_1_
 s__Ruminococcus_callidus
 s__Lachnospiraceae_bacterium_1_4_
 s__Lachnospiraceae_bacterium_5_1_
 s__Bacteroides_caccae
 s__Streptococcus_parasanguinis
 s__Oscillibacter_unclassified
 s__Barnesiella_intestinihominis
 s__Bacteroides_plebeius
 s__Alistipes_sp_HGB5
 s__Adlercreutzia_equilifaciens
 s__Megasphaera_unclassified
 s__Dialister_succinatiphilus
 s__Bacteroides_thetaiotaomicron
 s__Clostridium_leptum
 s__Collinsella_tanakaiei
 s__Escherichia_unclassified
 s__Parabacteroides_distasonis
 s__Bacteroides_dorei
 s__Anaerostipes_hadrus
 s__Bacteroides_ovatus
 s__Eubacterium_biforme
 s__Collinsella_intestinalis
 s__Eubacterium_eligens
 s__Bifidobacterium_bifidum
 s__Alistipes_nderdonkii
 s__Alistipes_shahii
 s__Lachnospiraceae_bacterium_8_1_
 s__Ruminococcus_lactaris
 s__Bacteroides_faeris