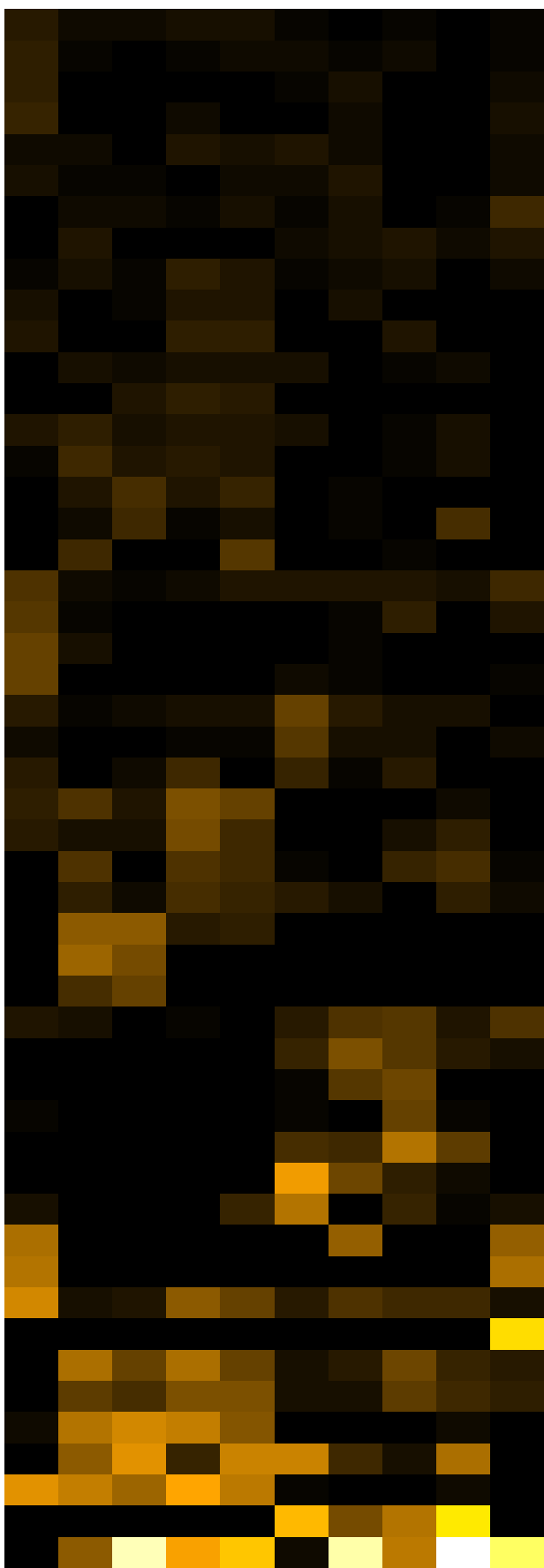
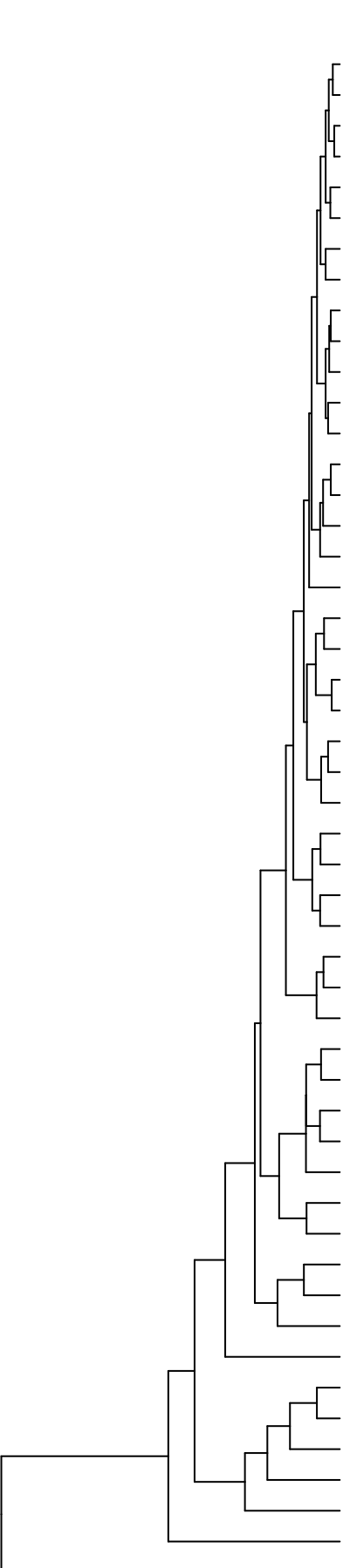




F31987.S10
F31987.S09
F31987.S08
F31987.S07
F31987.S06
F31987.S15
F31987.S14
F31987.S13
F31987.S12
F31987.S11

Lachnospiraceae [G-14] bacterium_MOT-184_nov_92.793%
 Lachnospiraceae [G-11] bacterium_MOT-176_nov_92.713%
 Lachnospiraceae [G-7] bacterium_MOT-172_nov_93.279%
 Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.969%
 Anaerobutyricum hallii_nov_90.249%
 Lachnospiraceae [G-12] bacterium_MOT-179_nov_94.501%
 Rikenella microfusum_nov_88.957%
 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_92.292%
 Saccharibacteria (TM7) [G-3] bacterium_HMT_351_nov_94.200%
 Mediterraneibacter [Ruminococcus] torques_nov_93.933%
 Fusicatenibacter saccharivorans_nov_90.041%
 Duncaniella freteri_nov_89.379%
 Lachnoclostridium [Clostridium] scindens_nov_89.648%
 Duncaniella freteri_nov_90.408%
 Duncaniella freteri_nov_93.293%
 Duncaniella freteri_nov_87.375%
 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_92.902%
 Lachnospiraceae [G-14] bacterium_MOT-185_nov_92.964%
 Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.549%
 Lachnoclostridium [Clostridium] scindens_nov_87.942%
 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_85.804%
 Lachnospiraceae [G-7] bacterium_MOT-172_nov_88.958%
 Lacrimispora xylanolytica_nov_94.363%
 Lachnospiraceae [G-11] bacterium_MOT-176_nov_94.929%
 Marvinbryantia formatexigens_nov_91.649%
 Duncaniella freteri
 Duncaniella freteri_nov_87.755%
 Oribacterium parvum_nov_89.979%
 Muribaculaceae [G-2] bacterium_MOT-104_nov_86.373%
 Duncaniella freteri_nov_91.002%
 Faecalibaculum rodentium
 Lactobacillus johnsonii
 Lachnospiraceae [G-1] bacterium_MOT-166
 Bacteroides uniformis_nov_96.099%
 Kineothrix alysoides_nov_86.667%
 Lachnoclostridium [Clostridium] scindens_nov_89.234%
 Lachnospiraceae [G-14] bacterium_MOT-185_nov_96.781%
 Eisenbergiella massiliensis_nov_87.578%
 Lachnoclostridium [Clostridium] herbivorans_nov_92.917%
 Lachnospiraceae [G-7] bacterium_MOT-172_nov_85.892%
 Lachnospiraceae [G-14] bacterium_MOT-184_nov_92.584%
 Alistipes senegalensis_nov_93.443%
 Turicibacter sanguinis_nov_95.923%
 Muribaculaceae [G-1] bacterium_MOT-129_nov_83.636%
 Muribaculaceae [G-1] bacterium_MOT-129_nov_86.290%
 Muribaculaceae [G-1] bacterium_MOT-129_nov_89.431%
 Bacteroides acidifaciens
 Duncaniella freteri_nov_89.069%
 Prevotella sp._HMT_820_nov_90.204%
 Ligilactobacillus murinus