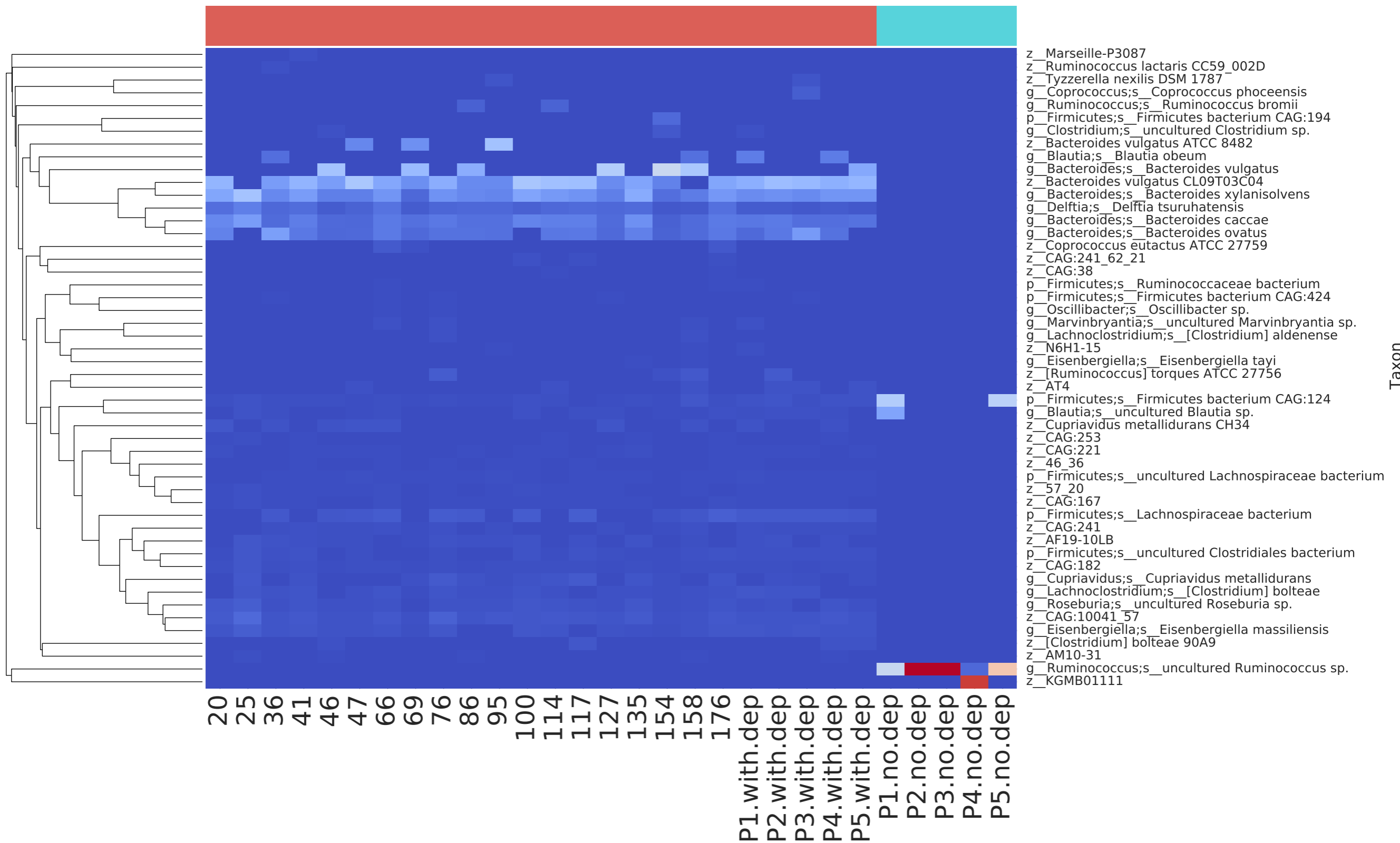


depletion no.depletion



20
25
36
41
46
47
66
69
76
86
95
100
114
117
127
135
154
158
176
P1.with.dep
P2.with.dep
P3.with.dep
P4.with.dep
P5.with.dep
P1.no.dep
P2.no.dep
P3.no.dep
P4.no.dep
P5.no.dep

z_Marseille-P3087
z_Ruminococcus lactaris CC59_002D
z_Tyzzarella nexilis DSM 1787
g_Coprococcus;s_Coprococcus phoceensis
g_Ruminococcus;s_Ruminococcus bromii
p_Firmicutes;s_Firmicutes bacterium CAG:194
g_Clostridium;s_uncultured Clostridium sp.
z_Bacteroides vulgatus ATCC 8482
g_Blautia;s_Blautia obeum
g_Bacteroides;s_Bacteroides vulgatus
z_Bacteroides vulgatus CL09T03C04
g_Bacteroides;s_Bacteroides xylanisolvens
g_Delftia;s_Delftia tsuruhatensis
g_Bacteroides;s_Bacteroides caccae
g_Bacteroides;s_Bacteroides ovatus
z_Coprococcus eutactus ATCC 27759
z_CAG:241_62_21
z_CAG:38
p_Firmicutes;s_Ruminococcaceae bacterium
p_Firmicutes;s_Firmicutes bacterium CAG:424
g_Oscillibacter;s_Oscillibacter sp.
g_Marvinbryantia;s_uncultured Marvinbryantia sp.
g_Lachnoclostridium;s_[Clostridium] aldenense
z_N6H1-15
g_Eisenbergiella;s_Eisenbergiella tayi
z_[Ruminococcus] torques ATCC 27756
z_AT4
p_Firmicutes;s_Firmicutes bacterium CAG:124
g_Blautia;s_uncultured Blautia sp.
z_Cupriavidus metallidurans CH34
z_CAG:253
z_CAG:221
z_46_36
p_Firmicutes;s_uncultured Lachnospiraceae bacterium
z_57_20
z_CAG:167
p_Firmicutes;s_Lachnospiraceae bacterium
z_CAG:241
z_AF19-10LB
p_Firmicutes;s_uncultured Clostridiales bacterium
z_CAG:182
g_Cupriavidus;s_Cupriavidus metallidurans
g_Lachnoclostridium;s_[Clostridium] bolteae
g_Roseburia;s_uncultured Roseburia sp.
z_CAG:10041_57
g_Eisenbergiella;s_Eisenbergiella massiliensis
z_[Clostridium] bolteae 90A9
z_AM10-31
g_Ruminococcus;s_uncultured Ruminococcus sp.
z_KGMB01111

Taxon