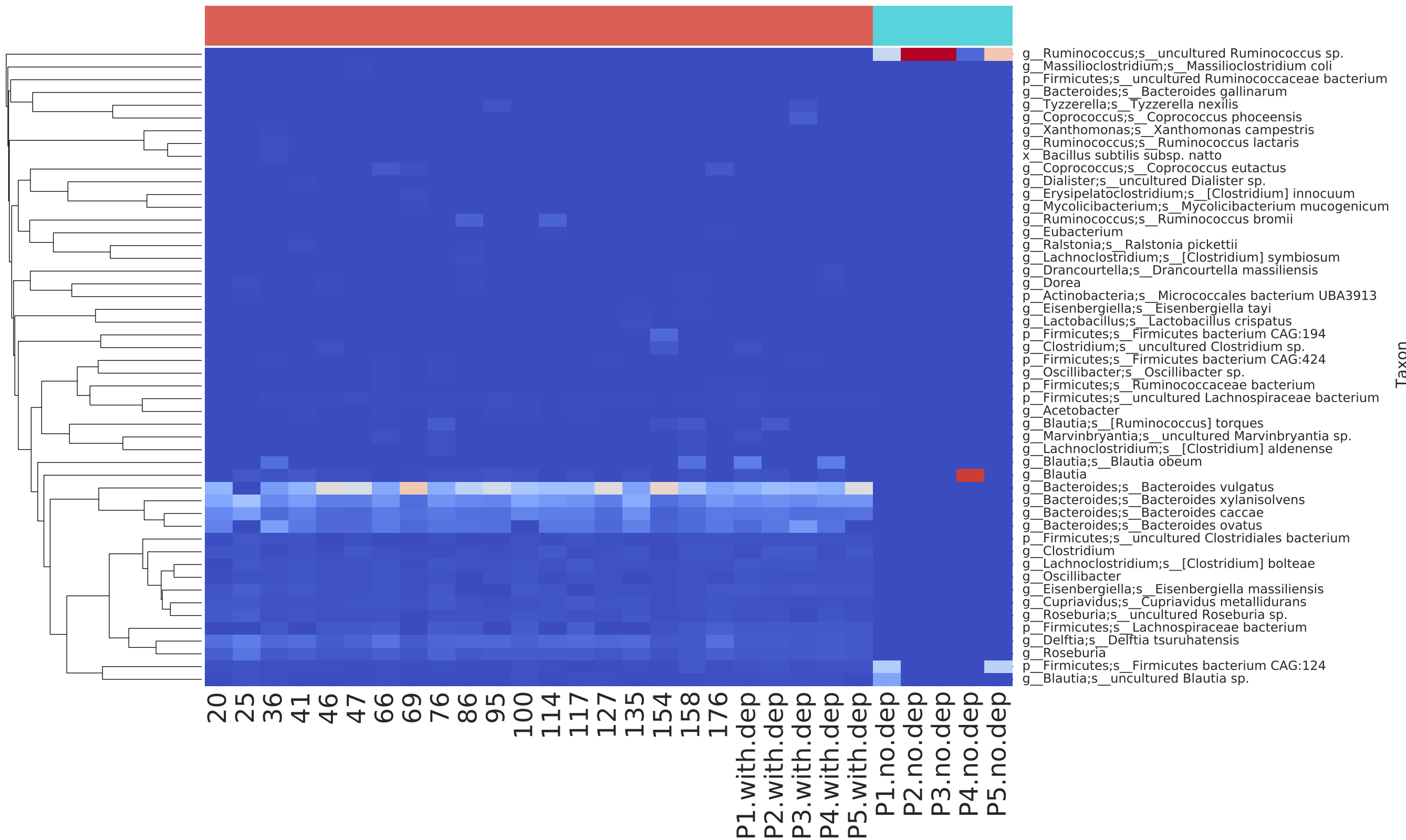


depletion no.depletion



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

g_Ruminococcus;s_uncultured Ruminococcus sp.
g_Massilioclostridium;s_Massilioclostridium coli
p_Firmicutes;s_uncultured Ruminococcaceae bacterium
g_Bacteroides;s_Bacteroides gallinarum
g_Tyzzereella;s_Tyzzereella nexilis
g_Coprococcus;s_Coprococcus phoceensis
g_Xanthomonas;s_Xanthomonas campestris
g_Ruminococcus;s_Ruminococcus lactaris
x_Bacillus subtilis subsp. natto
g_Coprococcus;s_Coprococcus eutactus
g_Dialister;s_uncultured Dialister sp.
g_Erysipelatoclostridium;s_[Clostridium] innocuum
g_Mycolicibacterium;s_Mycolicibacterium mucogenicum
g_Ruminococcus;s_Ruminococcus bromii
g_Eubacterium
g_Ralstonia;s_Ralstonia pickettii
g_Lachnoclostridium;s_[Clostridium] symbiosum
g_Drancourtella;s_Drancourtella massiliensis
g_Dorea
p_Actinobacteria;s_Micrococcales bacterium UBA3913
g_Eisenbergiella;s_Eisenbergiella tayi
g_Lactobacillus;s_Lactobacillus crispatus
p_Firmicutes;s_Firmicutes bacterium CAG:194
g_Clostridium;s_uncultured Clostridium sp.
p_Firmicutes;s_Firmicutes bacterium CAG:424
g_Oscillibacter;s_Oscillibacter sp.
p_Firmicutes;s_Ruminococcaceae bacterium
p_Firmicutes;s_uncultured Lachnospiraceae bacterium
g_Acetobacter
g_Blautia;s_[Ruminococcus] torques
g_Marvinbryantia;s_uncultured Marvinbryantia sp.
g_Lachnoclostridium;s_[Clostridium] aldenense
g_Blautia;s_Blautia obeum
g_Blautia
g_Bacteroides;s_Bacteroides vulgatus
g_Bacteroides;s_Bacteroides xylanisolvens
g_Bacteroides;s_Bacteroides caccae
g_Bacteroides;s_Bacteroides ovatus
p_Firmicutes;s_uncultured Clostridiales bacterium
g_Clostridium
g_Lachnoclostridium;s_[Clostridium] bolteae
g_Oscillibacter
g_Eisenbergiella;s_Eisenbergiella massiliensis
g_Cupriavidus;s_Cupriavidus metallidurans
g_Roseburia;s_uncultured Roseburia sp.
p_Firmicutes;s_Lachnospiraceae bacterium
g_Delftia;s_Delftia tsuruhatensis
g_Roseburia
p_Firmicutes;s_Firmicutes bacterium CAG:124
g_Blautia;s_uncultured Blautia sp.

Taxon