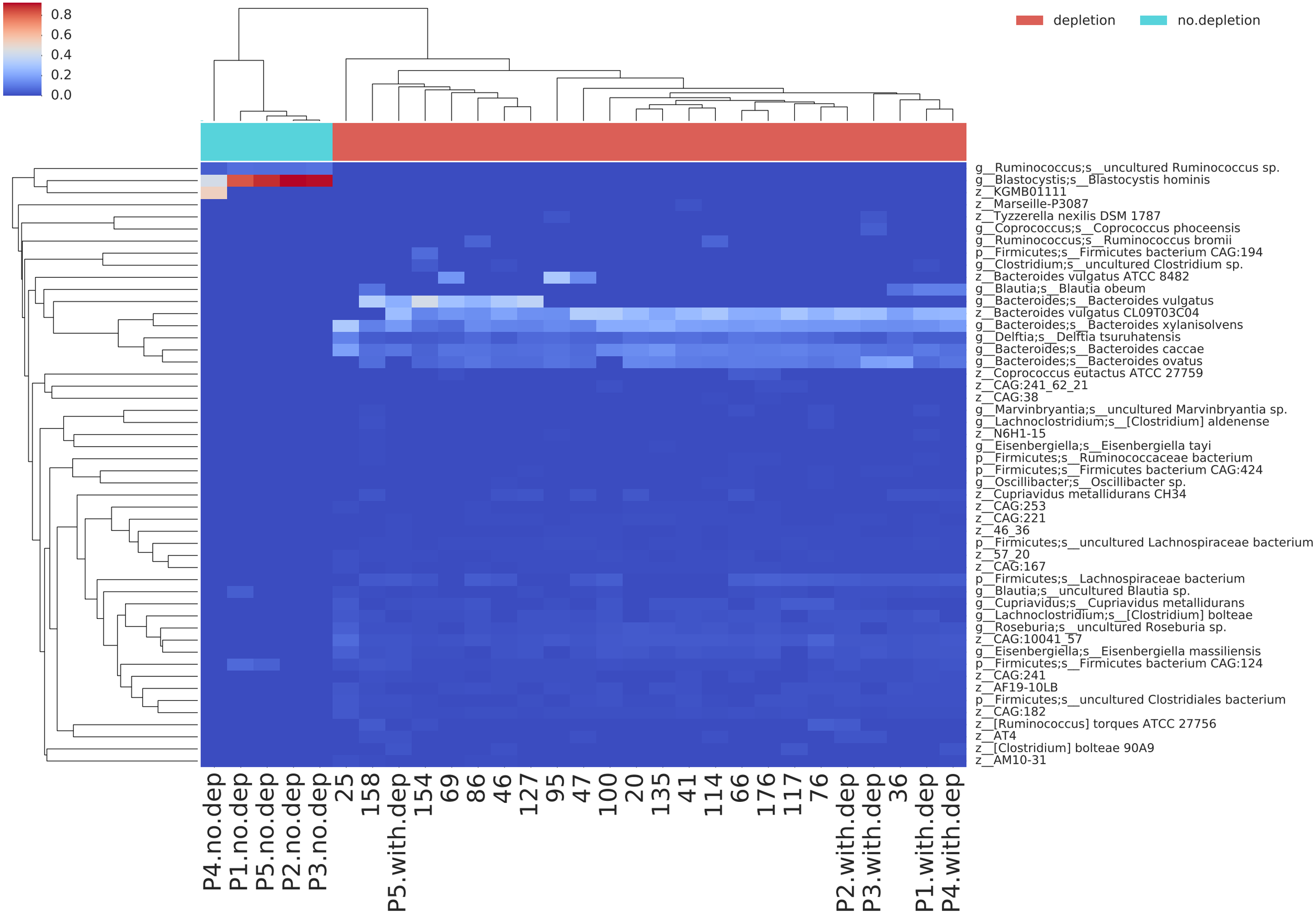
0.8
0.6
0.4
0.2
0.0
depletion
no.depletion
g__Ruminococcus;s__uncultured Ruminococcus sp.
g__Blastocystis;s__Blastocystis hominis
z__KGMB01111
z__Marseille-P3087
z__Tyzzerella nexilis DSM 1787
g__Coprococcus;s__Coprococcus phoceensis
g__Ruminococcus;s__Ruminococcus bromii
p__Firmicutes;s__Firmicutes 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
g__Marvinbryantia;s__uncultured Marvinbryantia sp.
g__Lachnoclostridium;s__[Clostridium] aldenense
z__N6H1-15
g__Eisenbergiella;s__Eisenbergiella tayi
p__Firmicutes;s__Ruminococcaceae bacterium
p__Firmicutes;s__Firmicutes bacterium CAG:424
g__Oscillibacter;s__Oscillibacter 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
g__Blautia;s__uncultured Blautia sp.
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
p__Firmicutes;s__Firmicutes bacterium CAG:124
z__CAG:241
z__AF19-10LB
p__Firmicutes;s__uncultured Clostridiales bacterium
z__CAG:182
z__[Ruminococcus] torques ATCC 27756
z__AT4
z__[Clostridium] bolteae 90A9
z__AM10-31
Taxon
P4.no.dep
P1.no.dep
P5.no.dep
P2.no.dep
P3.no.dep
25
158
P5.with.dep
154
69
86
46
127
95
47
100
20
135
41
114
66
176
117
76
P2.with.dep
P3.with.dep
36
P1.with.dep
P4.with.dep