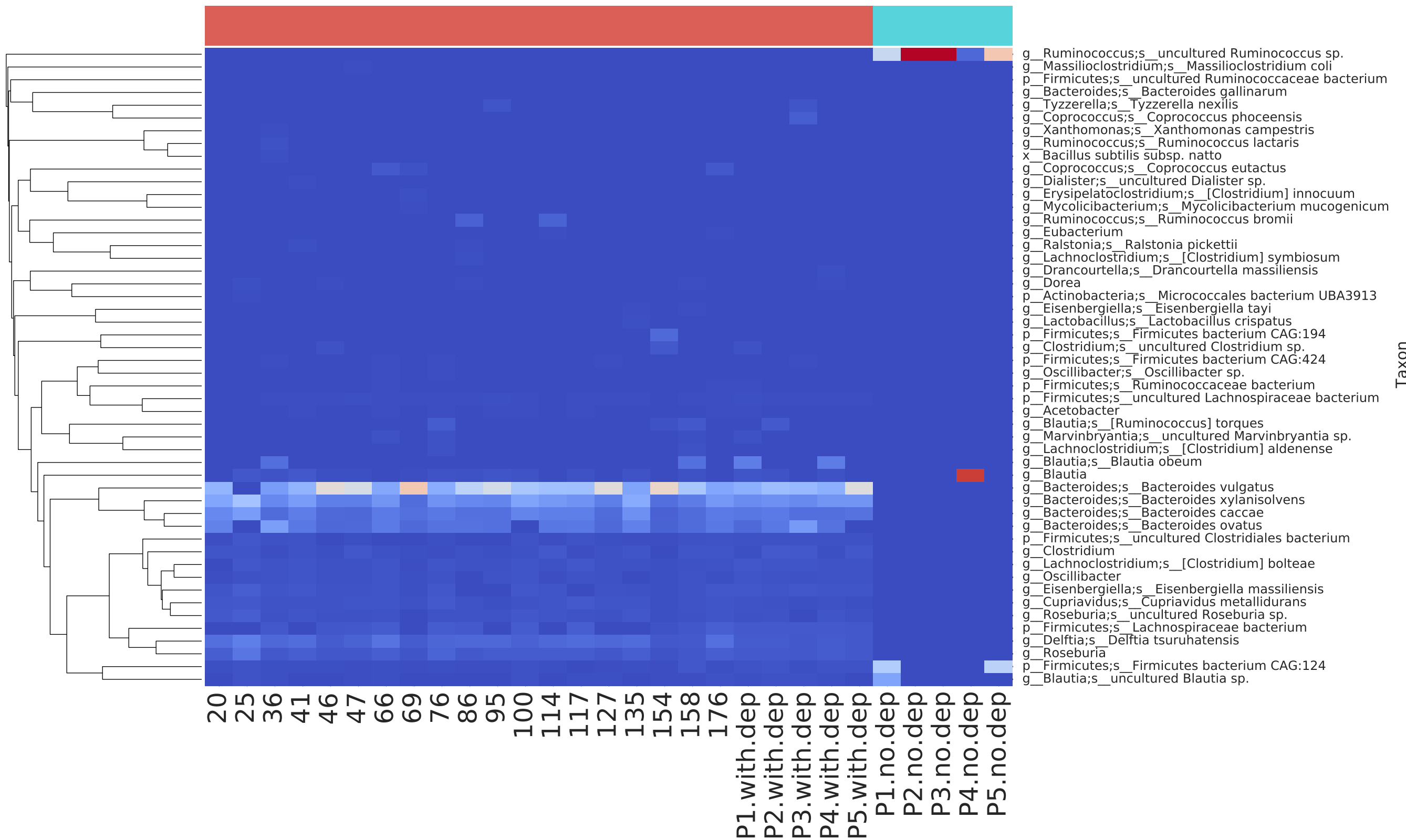


depletion no.depletion



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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