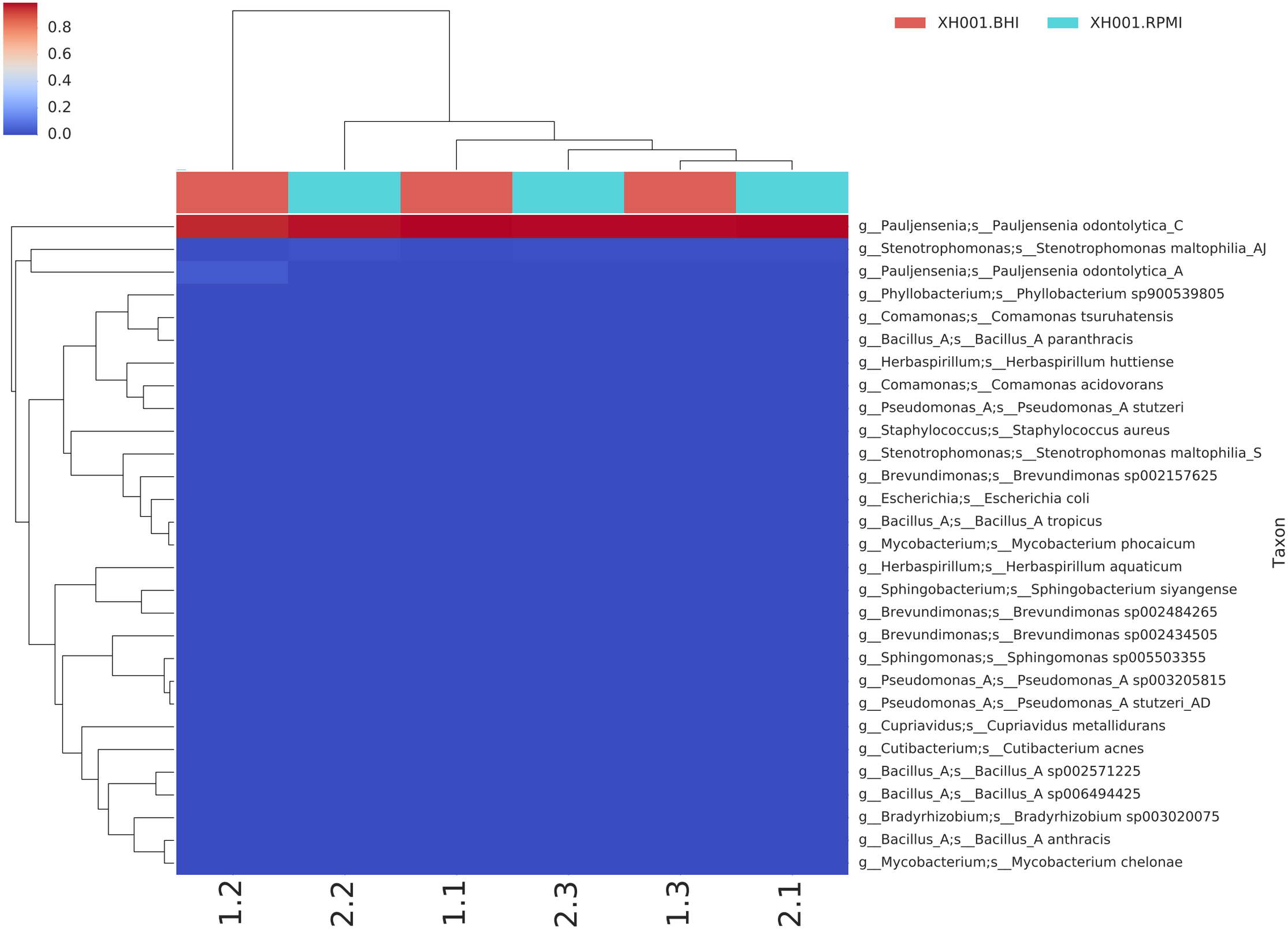

0.8
0.6
0.4
0.2
0.0
XH001.BHI
XH001.RPMI
g__Pauljensenia;s__Pauljensenia odontolytica_C
g__Stenotrophomonas;s__Stenotrophomonas maltophilia_AJ
g__Pauljensenia;s__Pauljensenia odontolytica_A
g__Phyllobacterium;s__Phyllobacterium sp900539805
g__Comamonas;s__Comamonas tsuruhatensis
g__Bacillus_A;s__Bacillus_A paranthracis
g__Herbaspirillum;s__Herbaspirillum huttiense
g__Comamonas;s__Comamonas acidovorans
g__Pseudomonas_A;s__Pseudomonas_A stutzeri
g__Staphylococcus;s__Staphylococcus aureus
g__Stenotrophomonas;s__Stenotrophomonas maltophilia_S
g__Brevundimonas;s__Brevundimonas sp002157625
g__Escherichia;s__Escherichia coli
g__Bacillus_A;s__Bacillus_A tropicus
g__Mycobacterium;s__Mycobacterium phocaicum
g__Herbaspirillum;s__Herbaspirillum aquaticum
g__Sphingobacterium;s__Sphingobacterium siyangense
g__Brevundimonas;s__Brevundimonas sp002484265
g__Brevundimonas;s__Brevundimonas sp002434505
g__Sphingomonas;s__Sphingomonas sp005503355
g__Pseudomonas_A;s__Pseudomonas_A sp003205815
g__Pseudomonas_A;s__Pseudomonas_A stutzeri_AD
g__Cupriavidus;s__Cupriavidus metallidurans
g__Cutibacterium;s__Cutibacterium acnes
g__Bacillus_A;s__Bacillus_A sp002571225
g__Bacillus_A;s__Bacillus_A sp006494425
g__Bradyrhizobium;s__Bradyrhizobium sp003020075
g__Bacillus_A;s__Bacillus_A anthracis
g__Mycobacterium;s__Mycobacterium chelonae
Taxon
1.2
2.2
1.1
2.3
1.3
2.1