

1.0 -

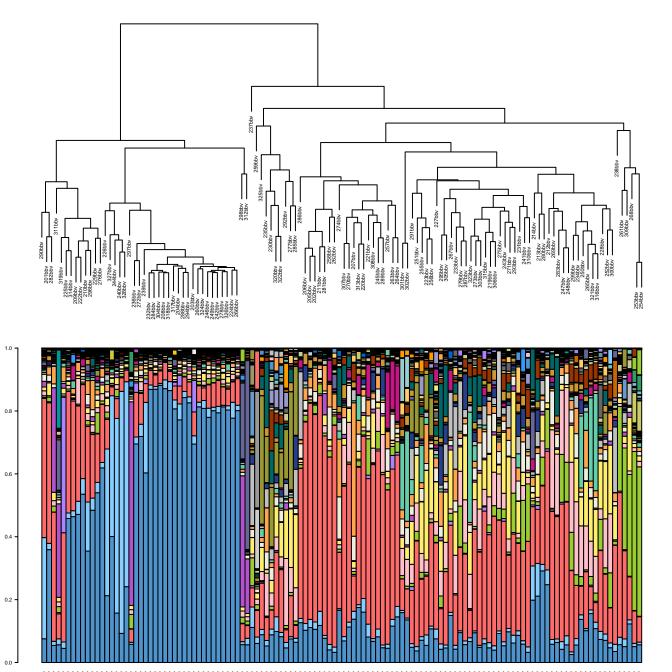
0.8

0.6

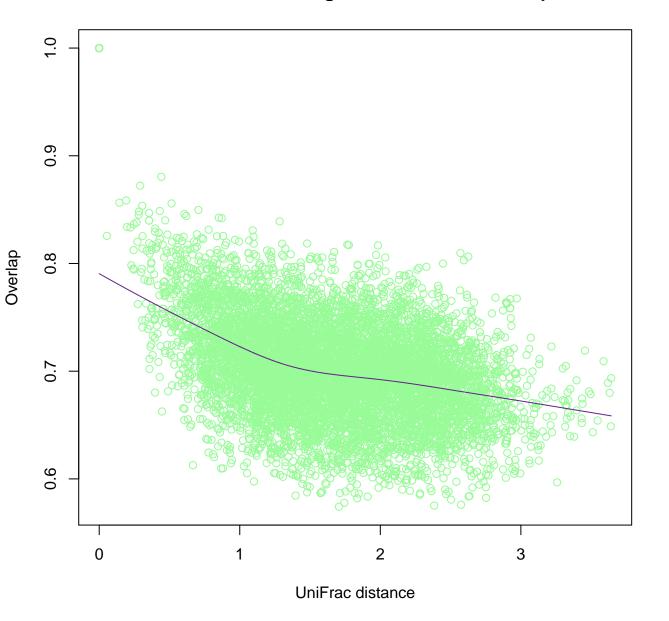
0.4

0.2 -

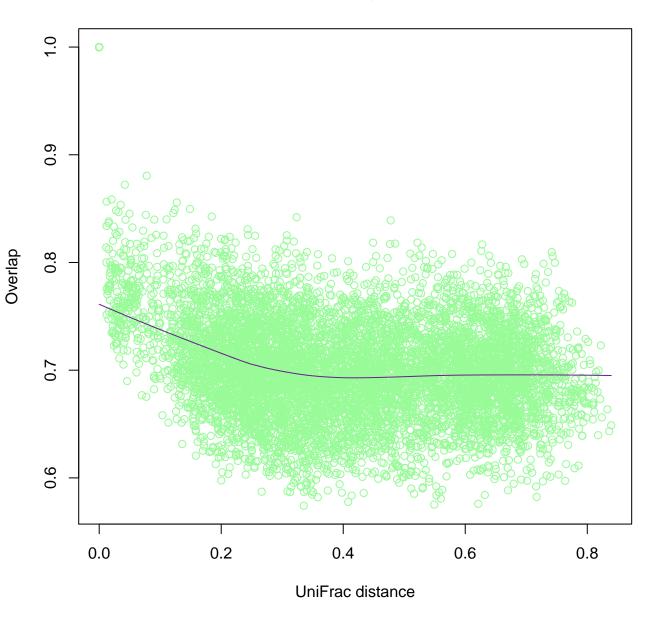
0.0



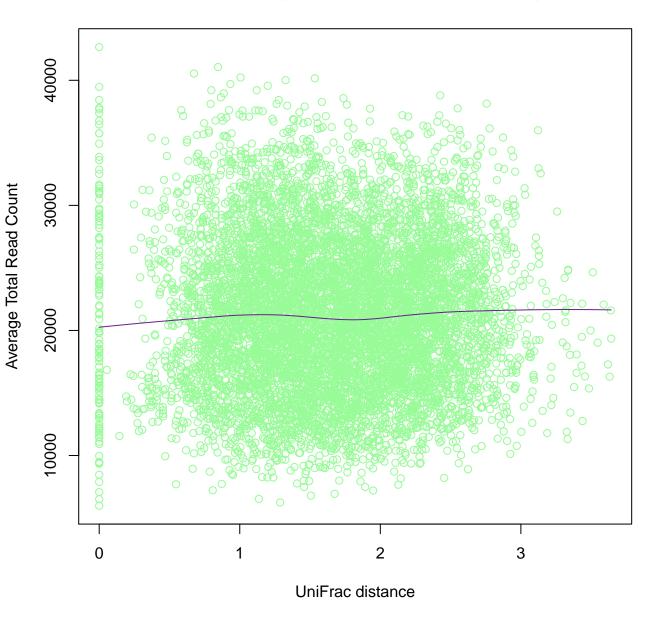
CLR transform weighted UniFrac vs. overlap



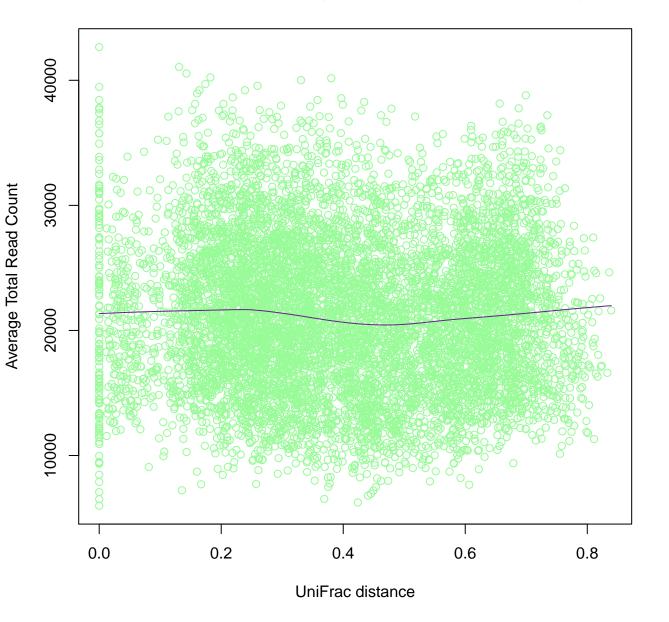
Proportional abundance weighted UniFrac vs. overlap



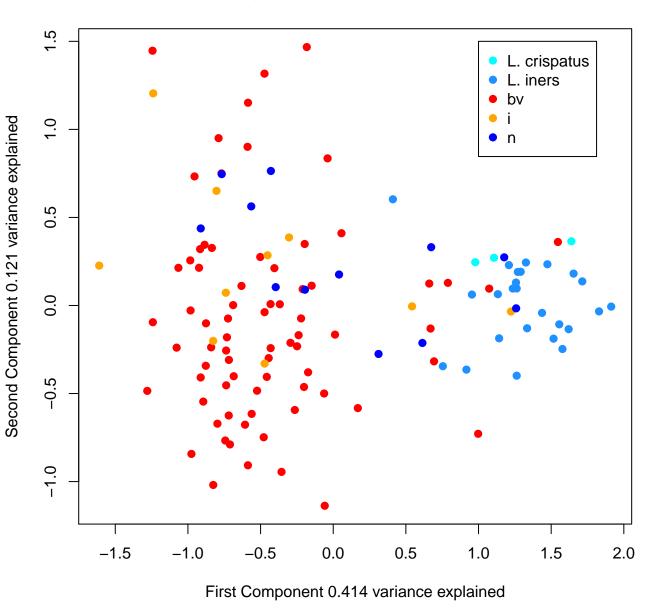
CLR transform weighted UniFrac vs. sequencing depth



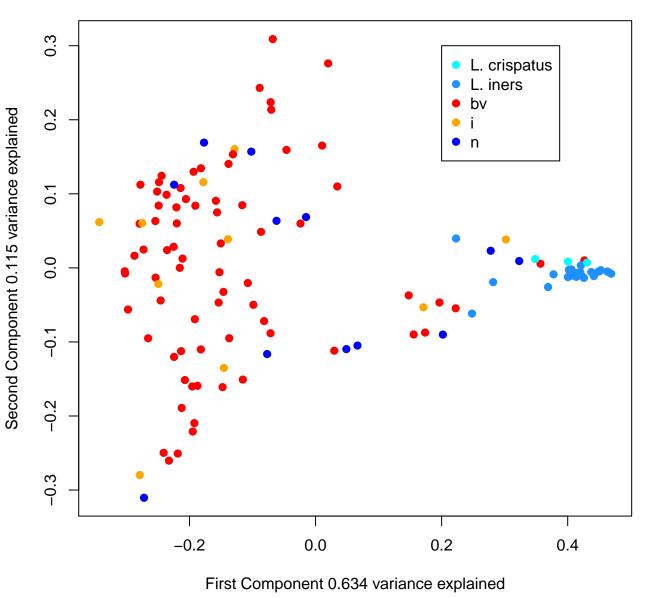
Proportional abundance weighted UniFrac vs. sequencing depth



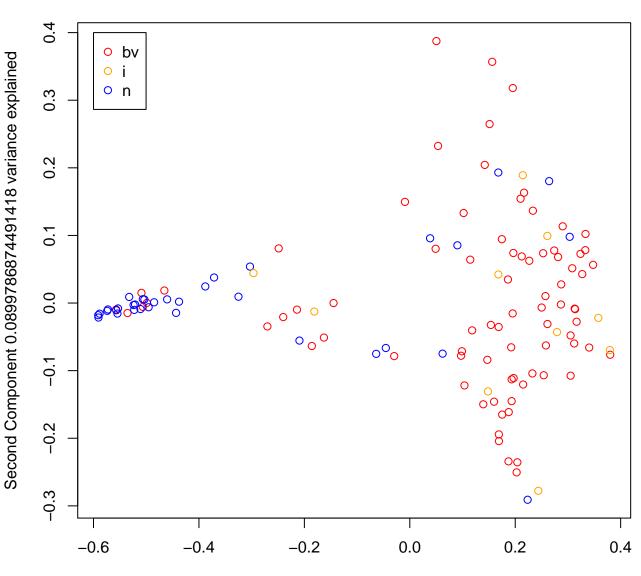
CLR transformed weighted UniFrac principle coordinates analysis



Proportional abundance weighted UniFrac principle coordinates analysis

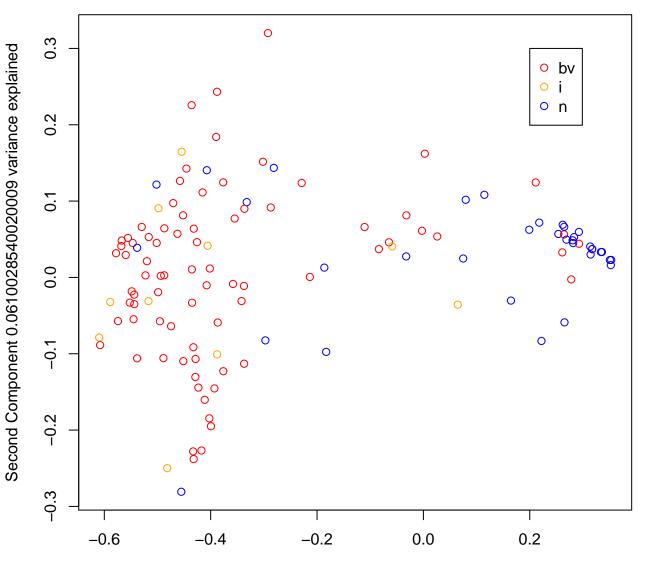


pcoa from qiime unifrac distances



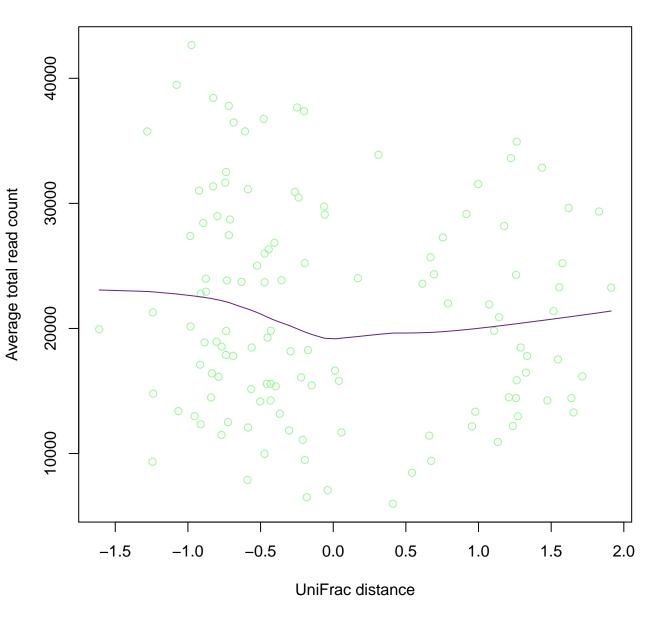
First Component 0.683924586703643 variance explained

qiime pcoa



First Component 0.593588246926922 variance explained

Proportional abundance weighted UniFrac vs. PCoA first component



CLR transform weighted UniFrac vs. PCoA first component

