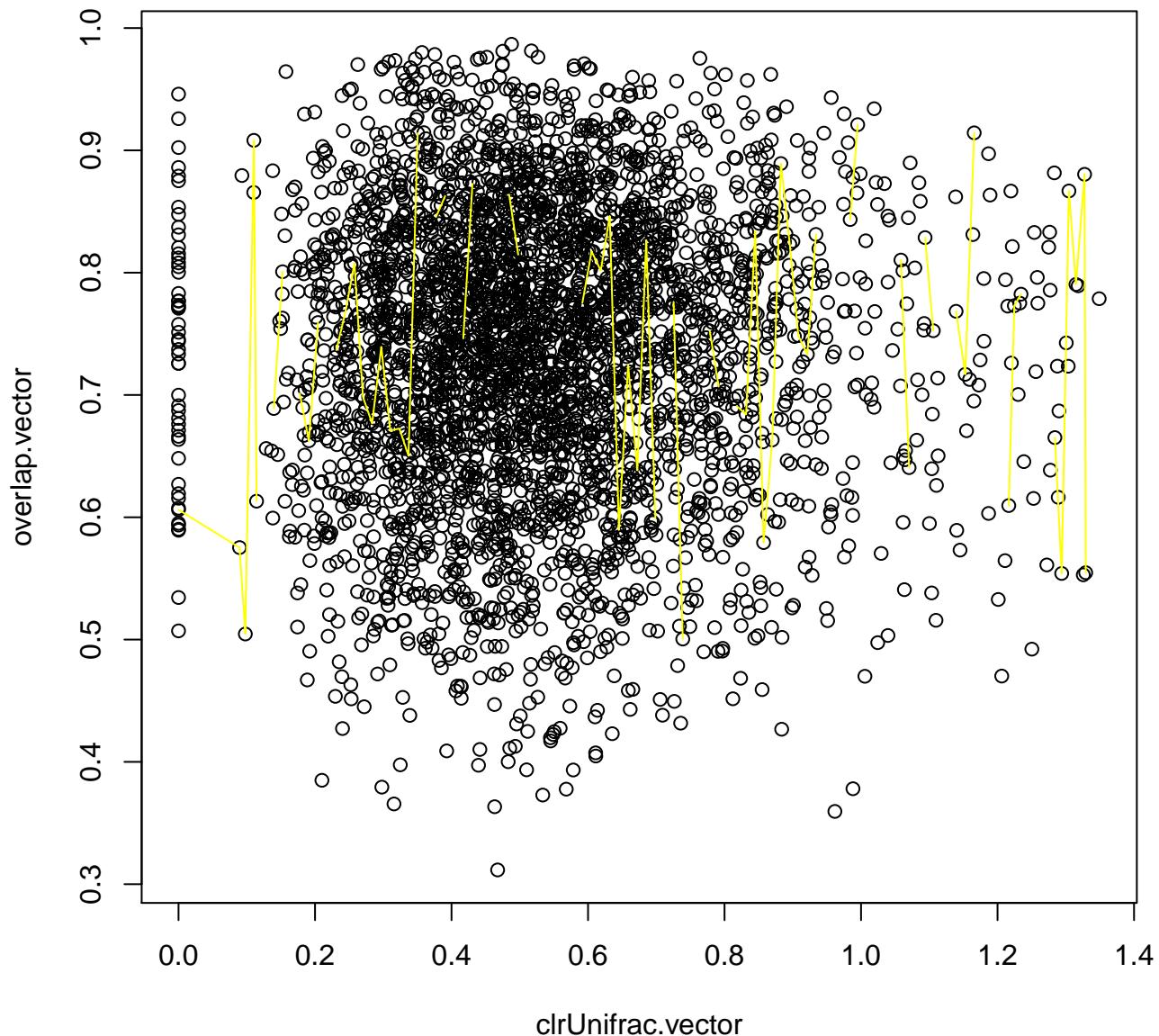
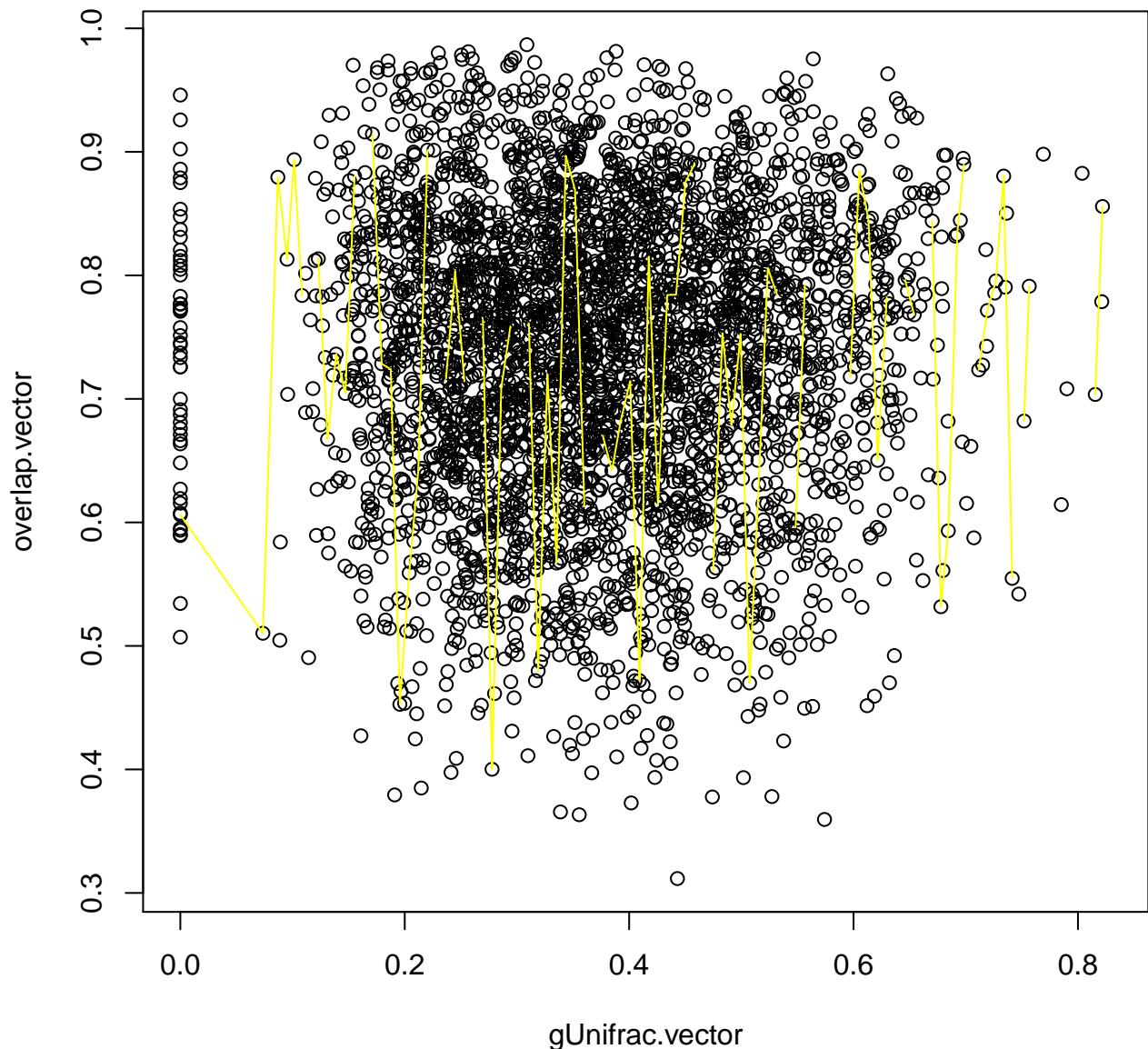


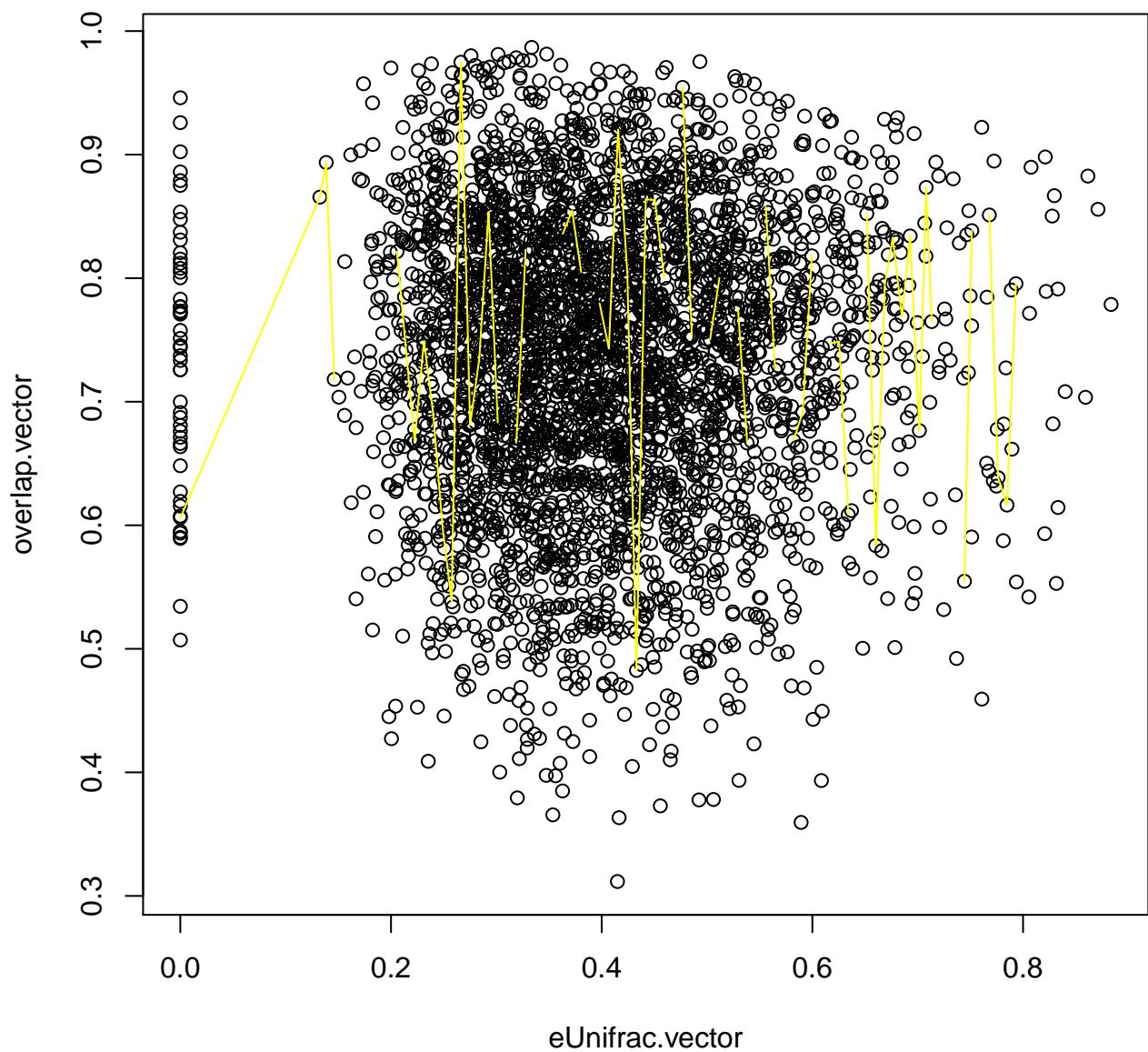
clr combination weights vs overlap



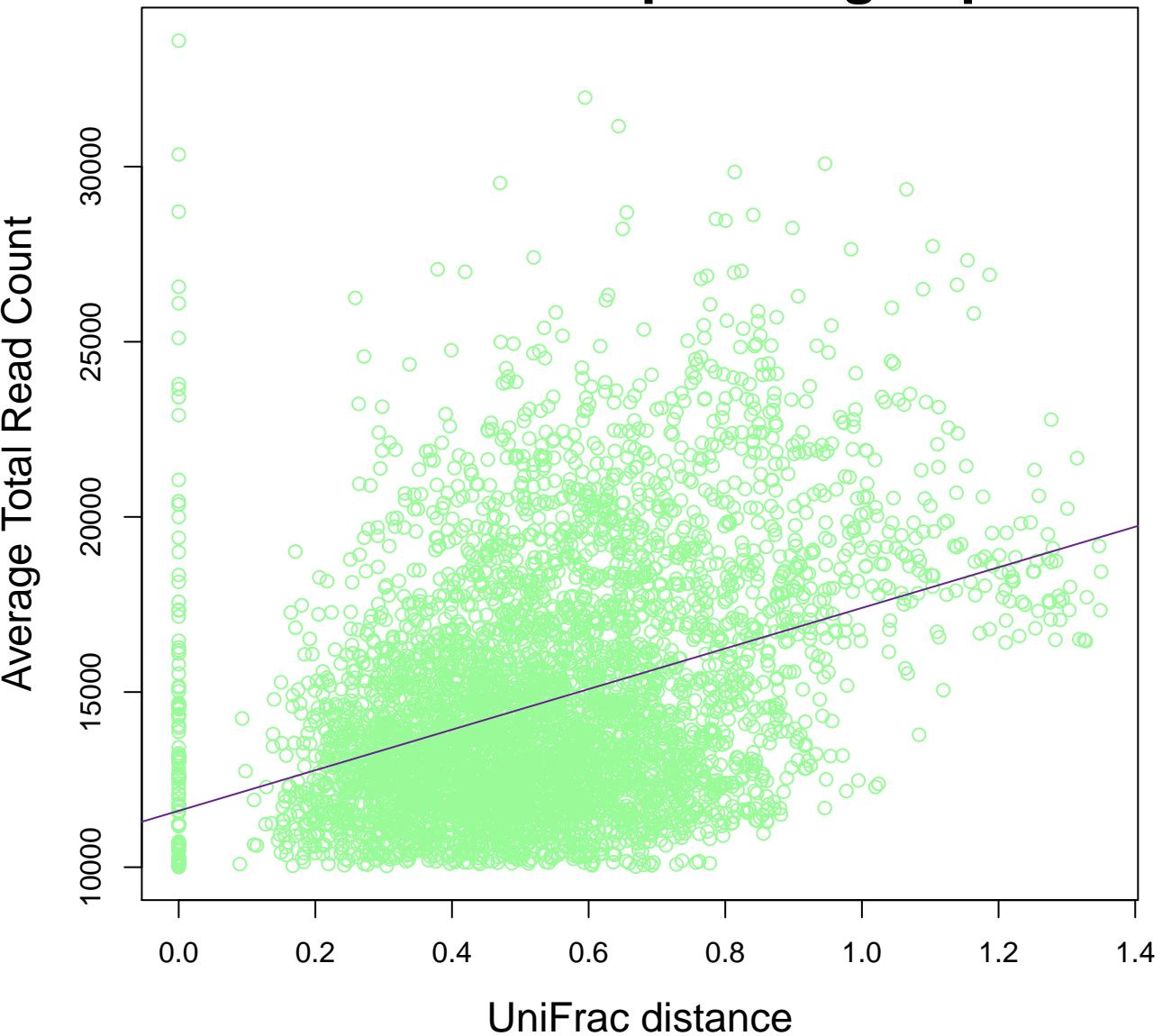
gunifrac vs overlap



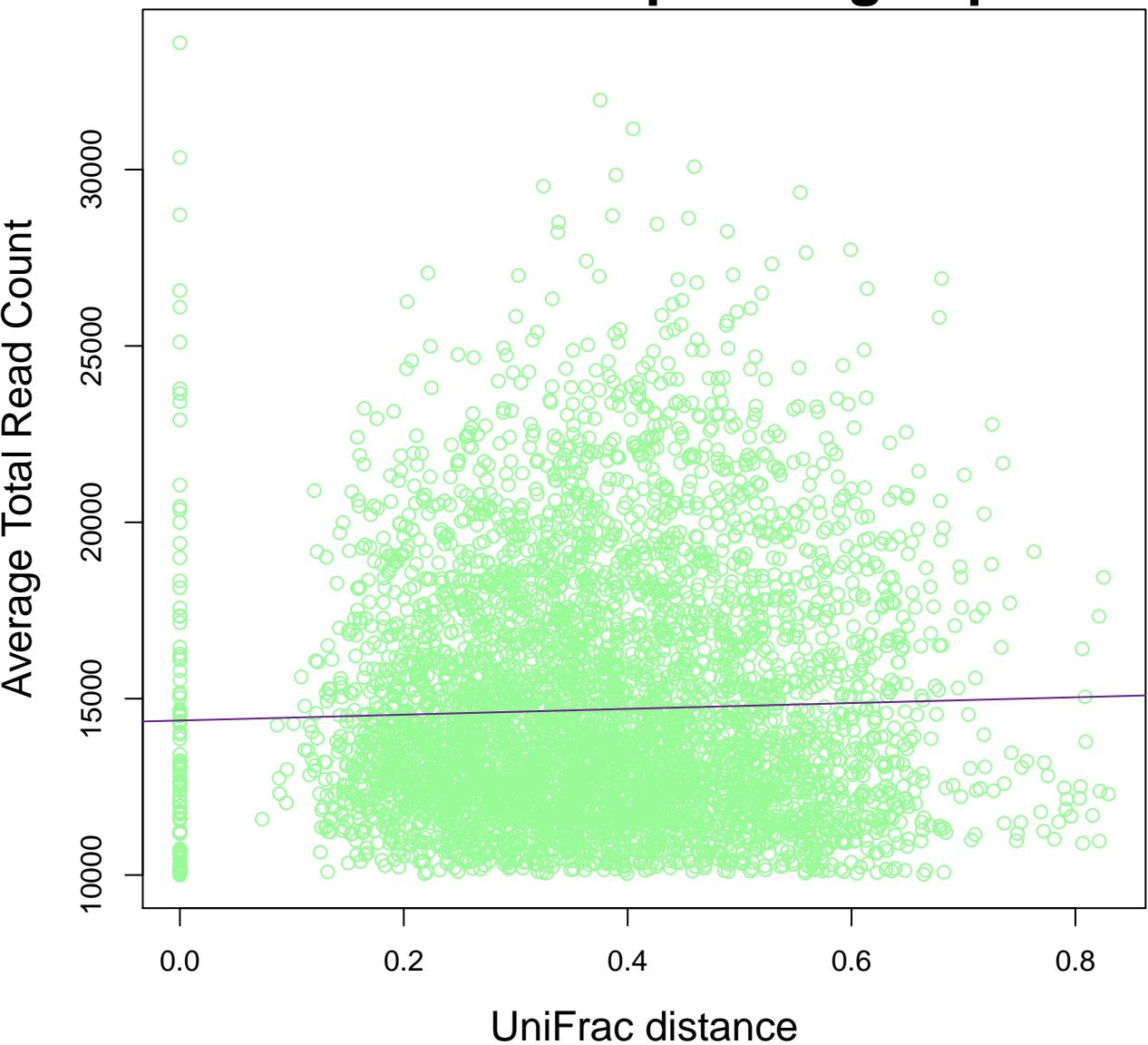
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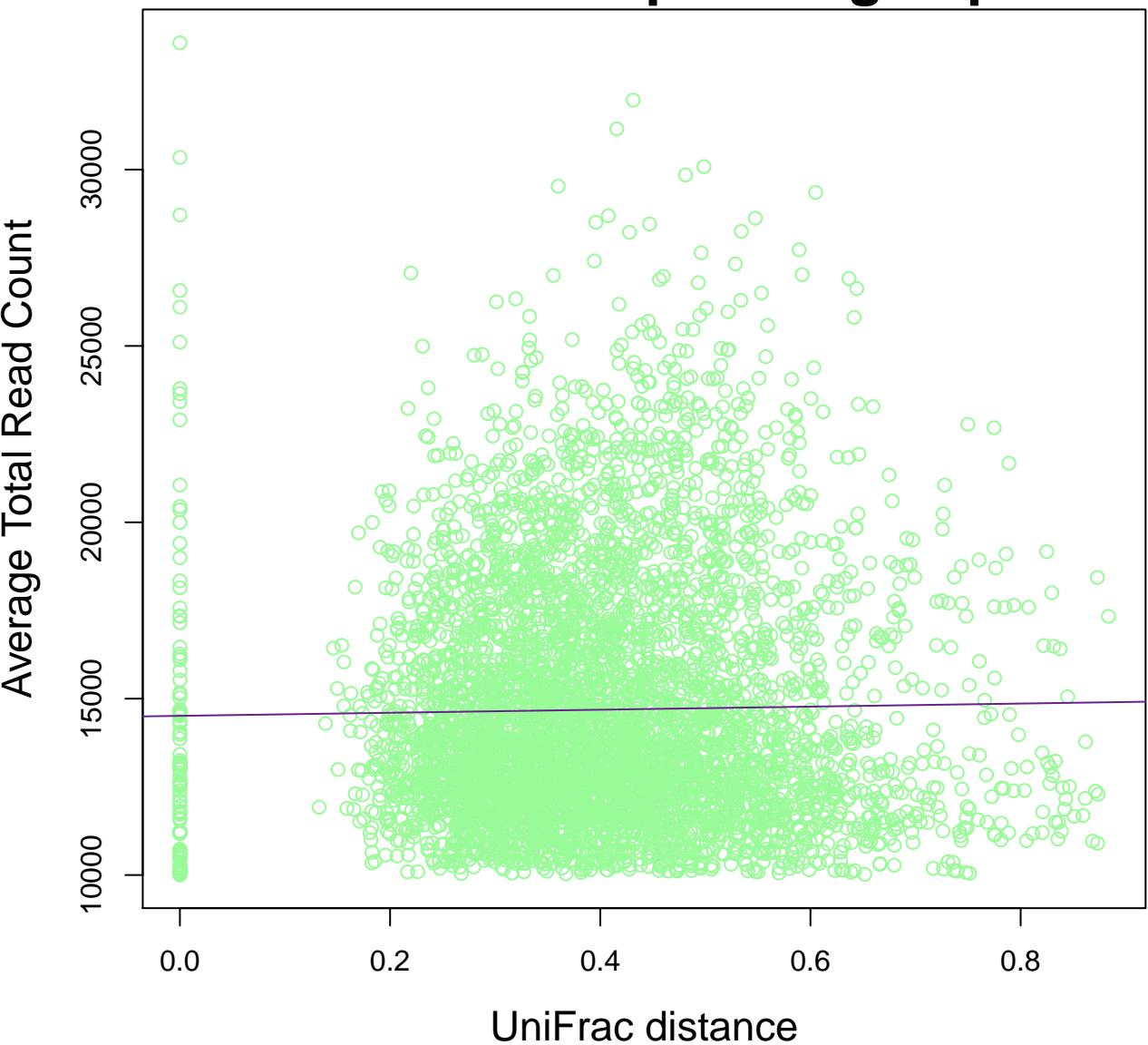
CLR transform weighted UniFrac vs. sequencing depth

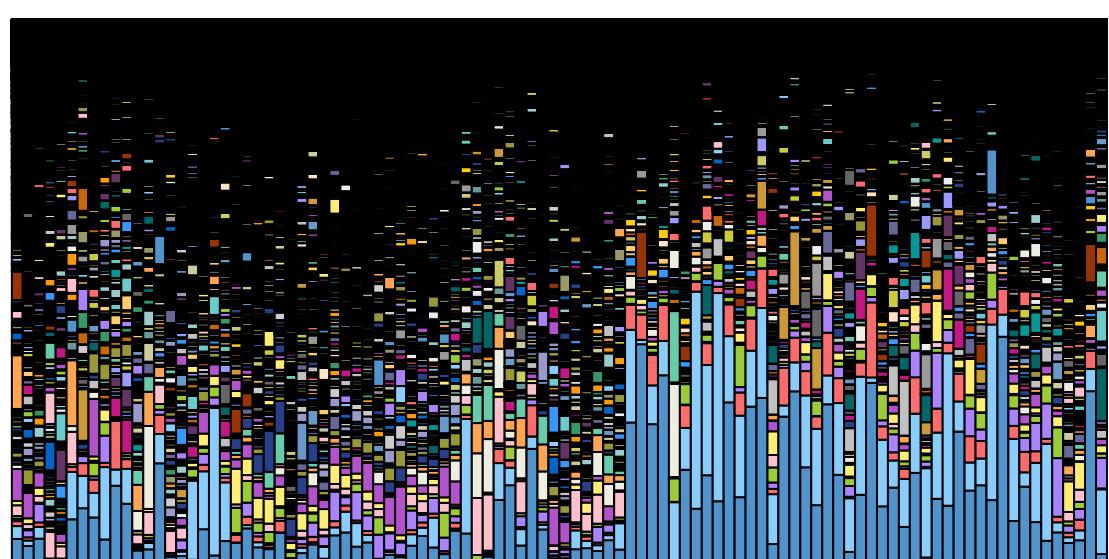
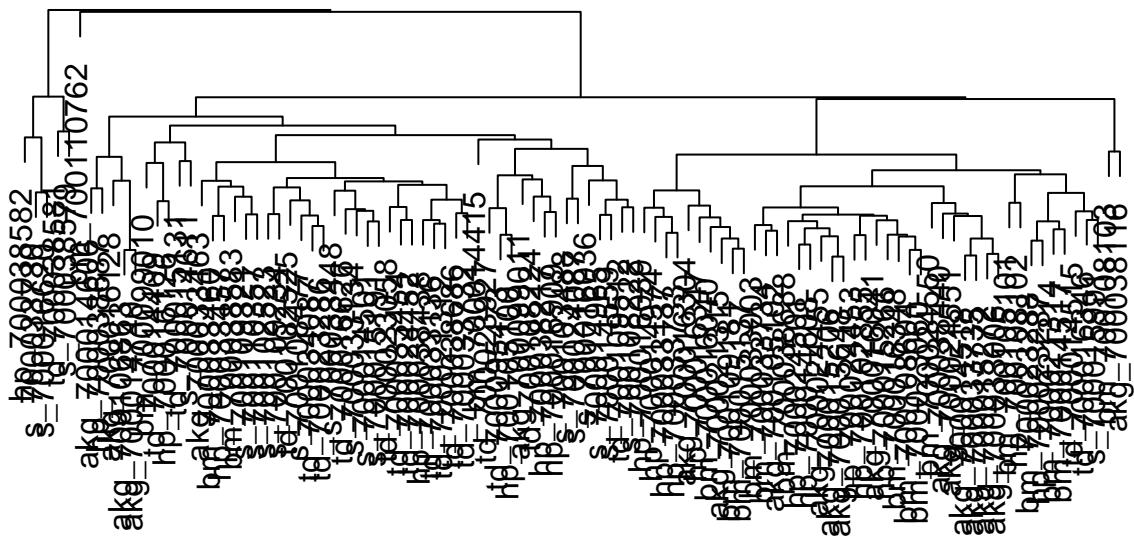


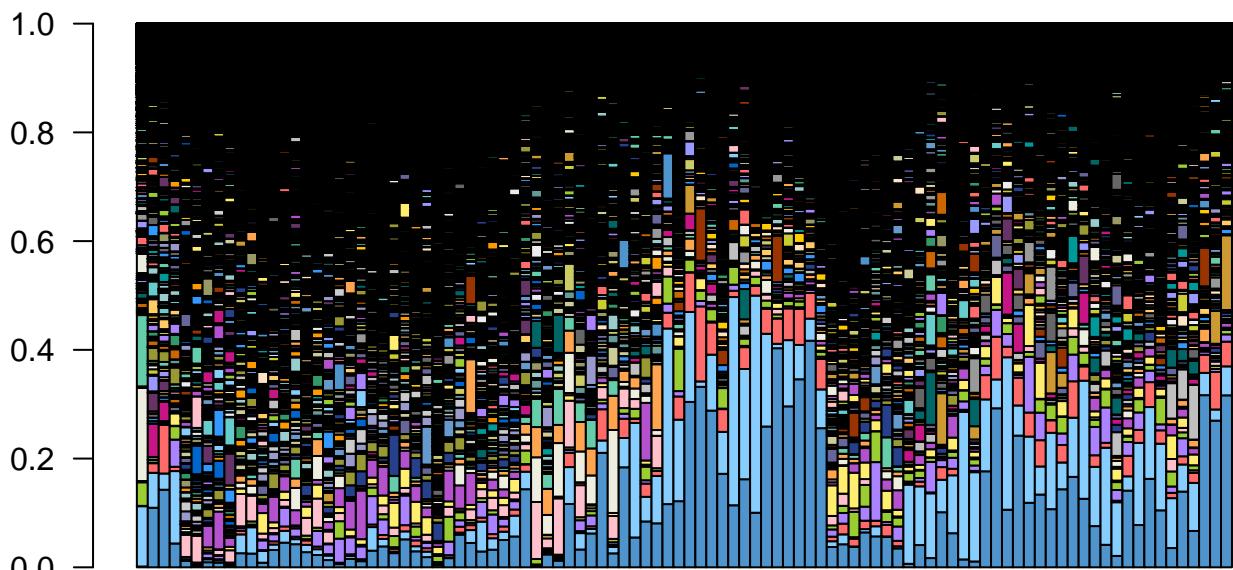
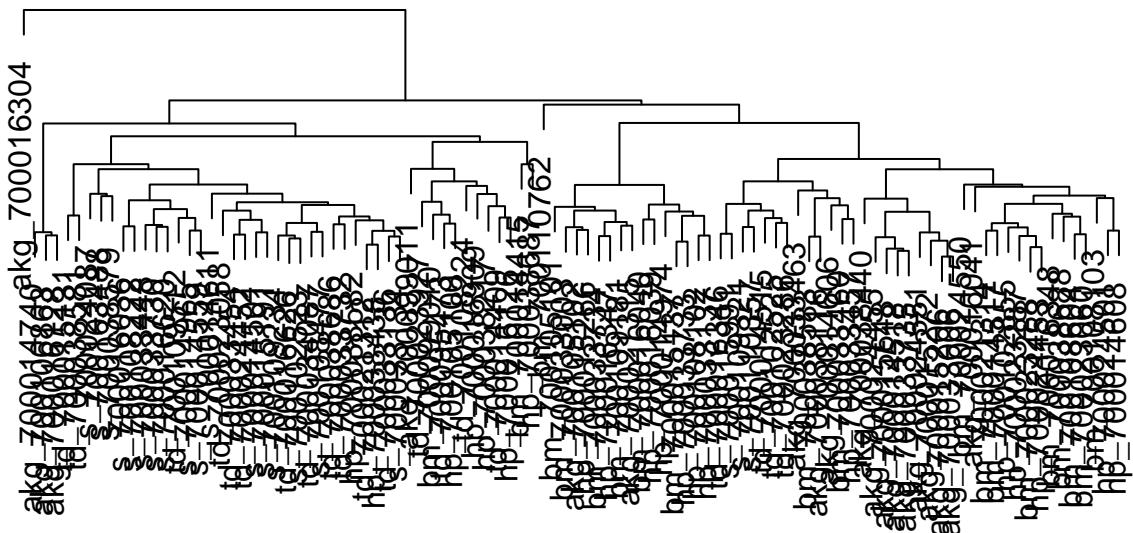
Proportional abundance weighted UniFrac vs. sequencing depth

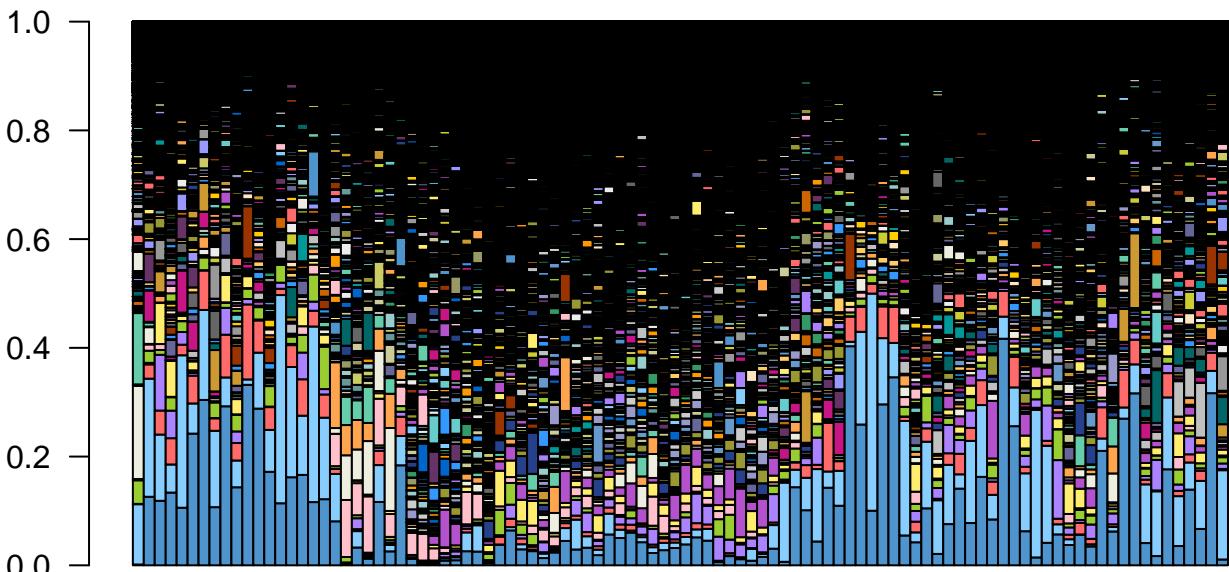
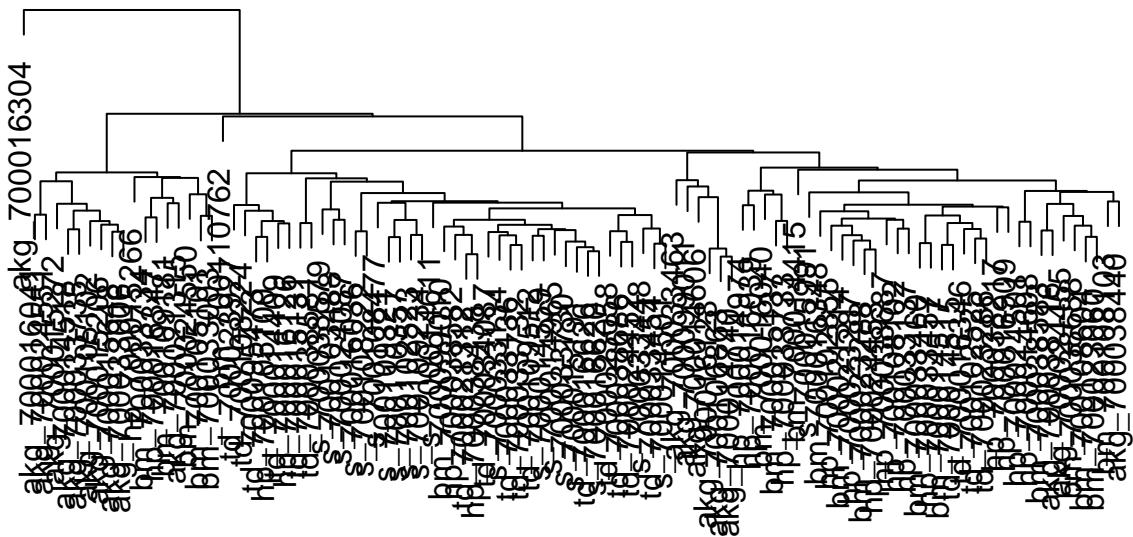


Entropy weighted UniFrac vs. sequencing depth





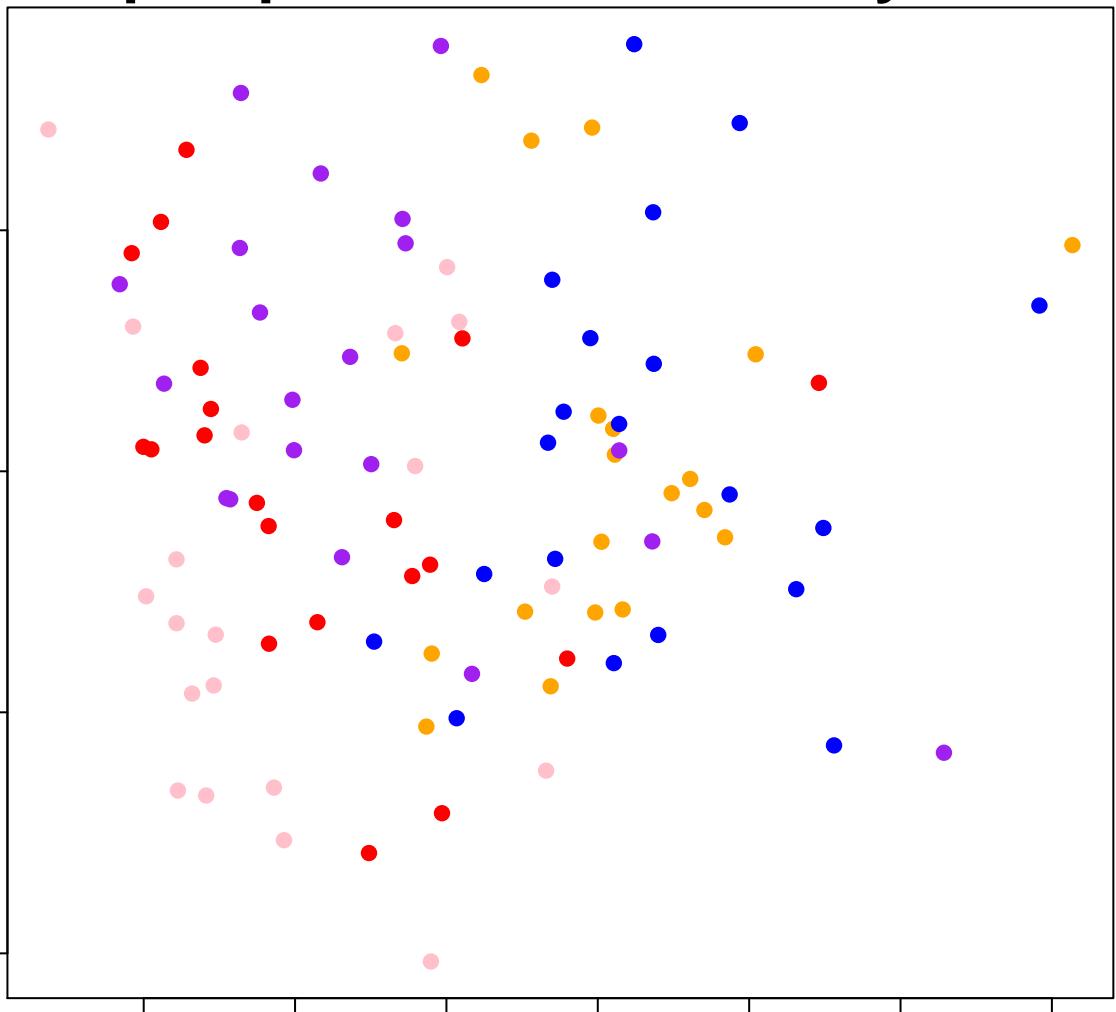




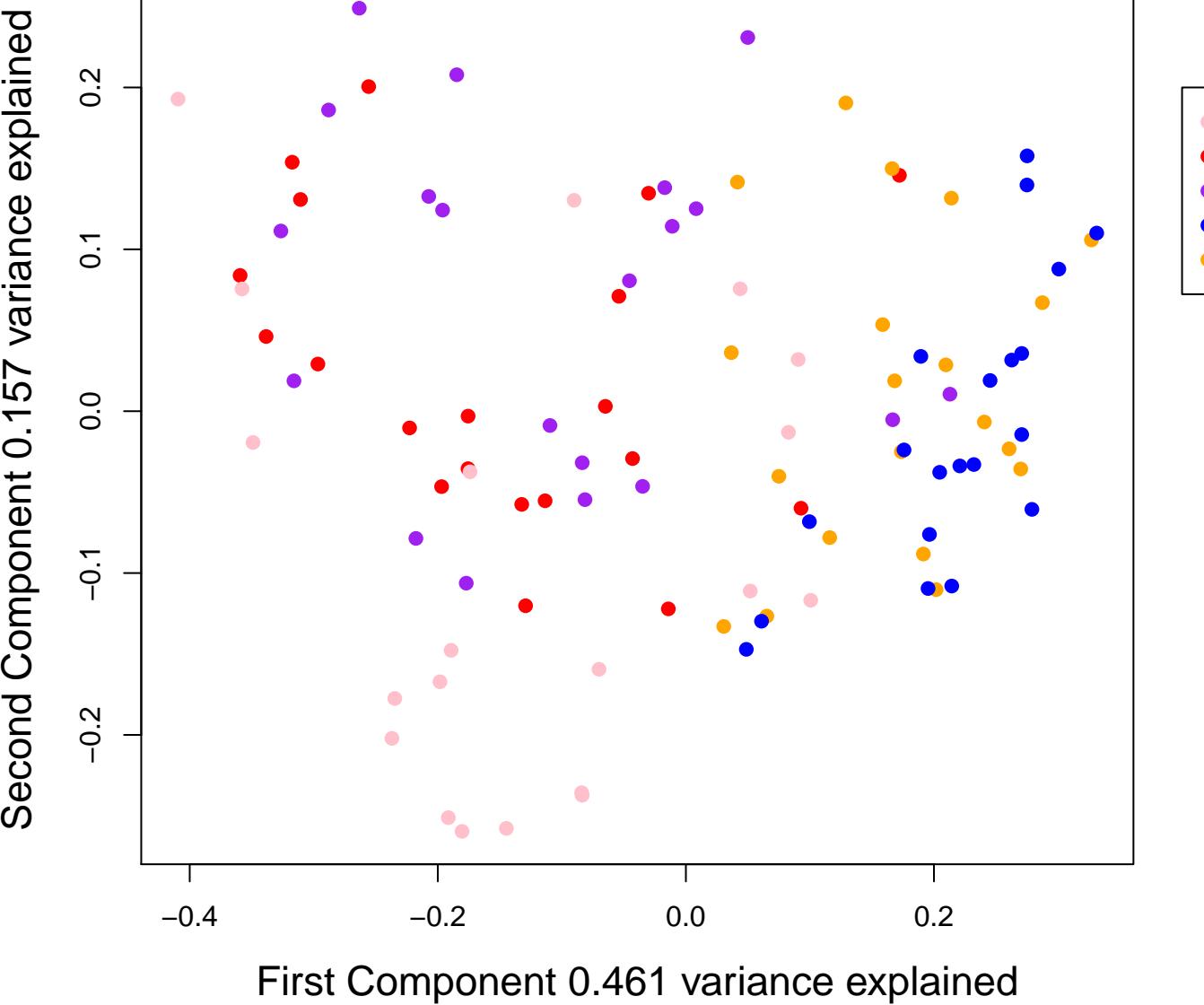
CLR transform weighted UniFrac principal coordinates analysis

Second Component 0.143 variance explained

First Component 0.431 variance explained



Proportional abundance weighted UniFrac principal coordinates analysis



Entropy weighted UniFrac principal coordinates analysis

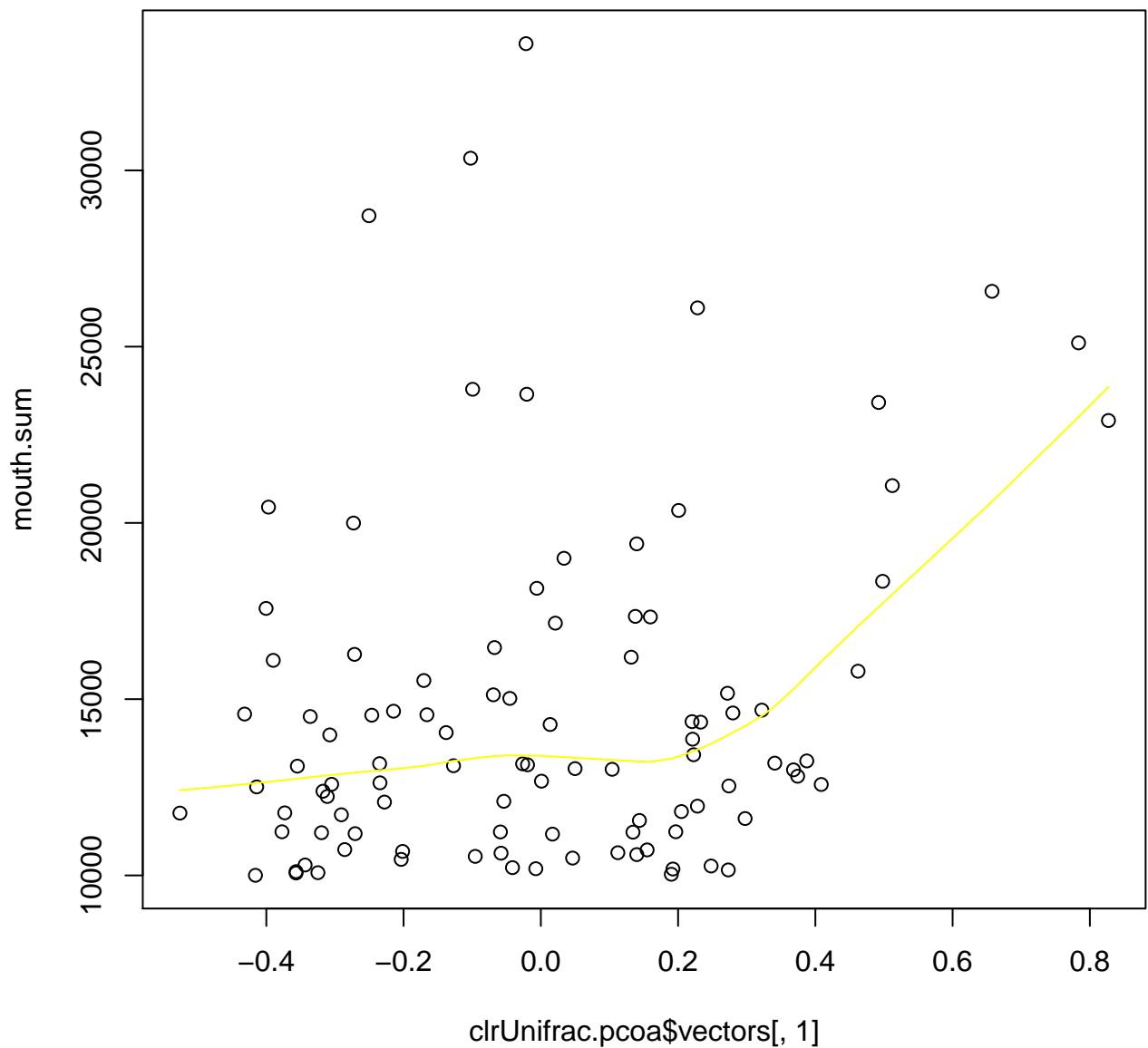
Second Component 0.102 variance explained

-0.4 -0.2 0.0 0.2 0.4

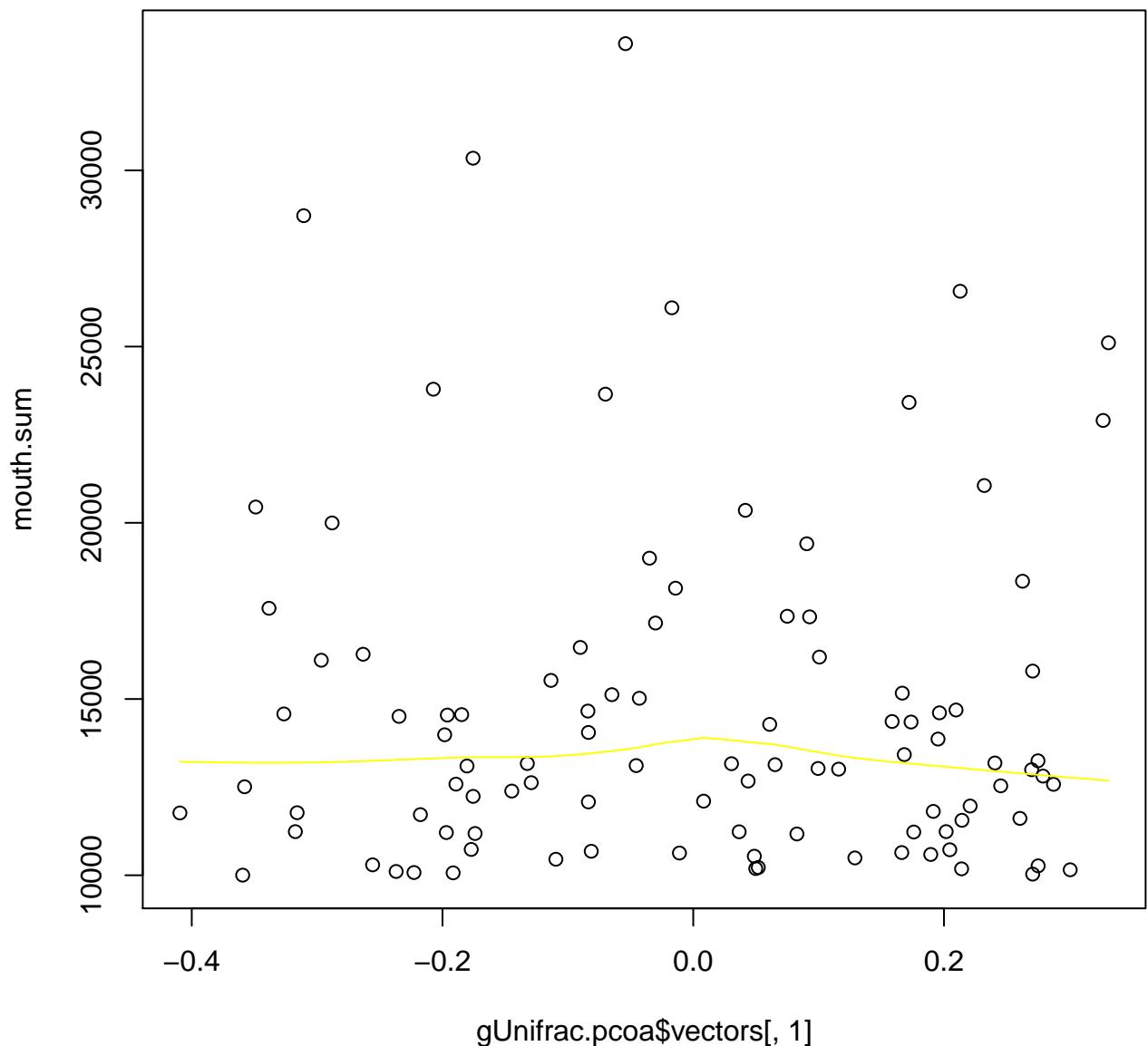
First Component 0.408 variance explained

att
bu
ha
sa
tor

clr combination weights vs avg



gunifrac vs avg



gunifrac vs avg

