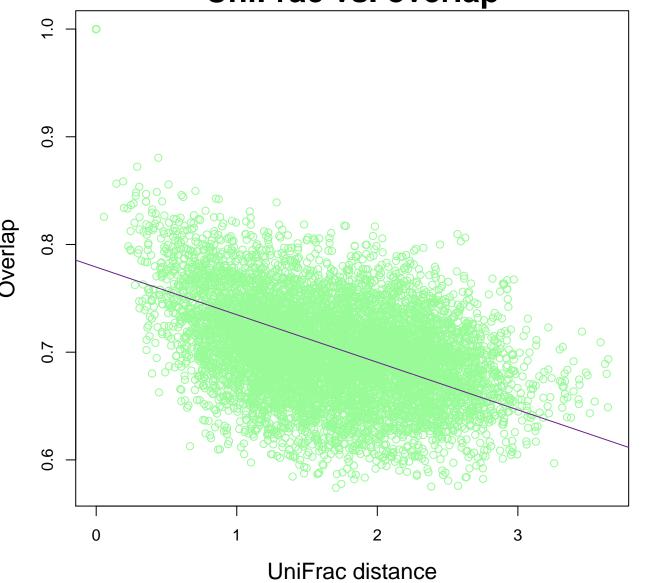
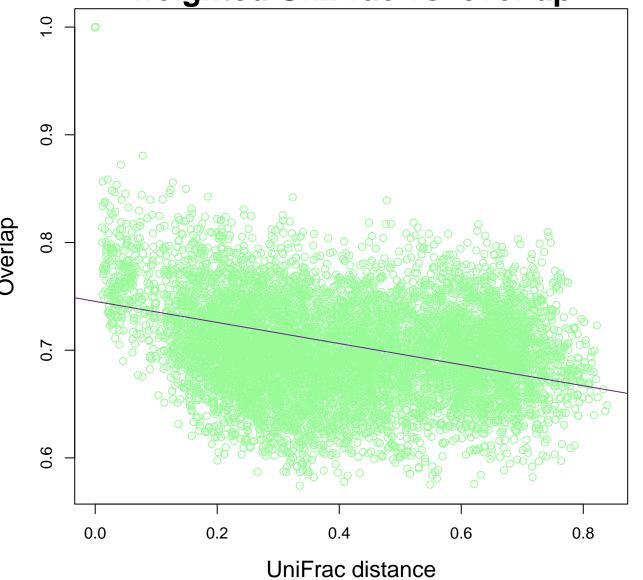
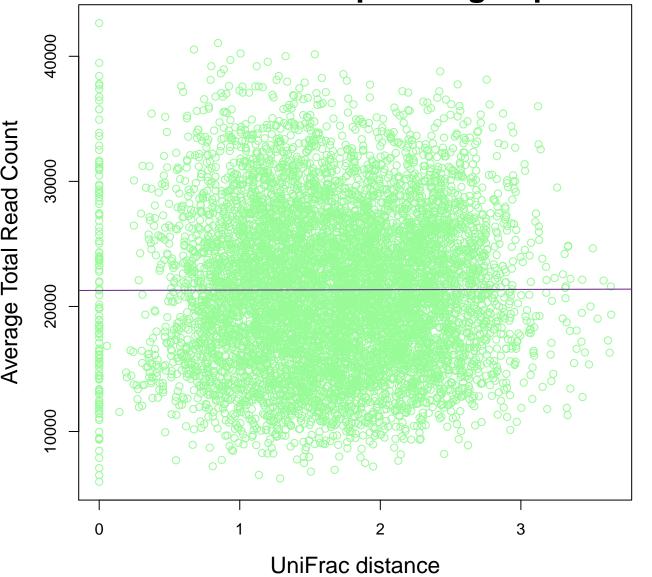
CLR transform weighted UniFrac vs. overlap



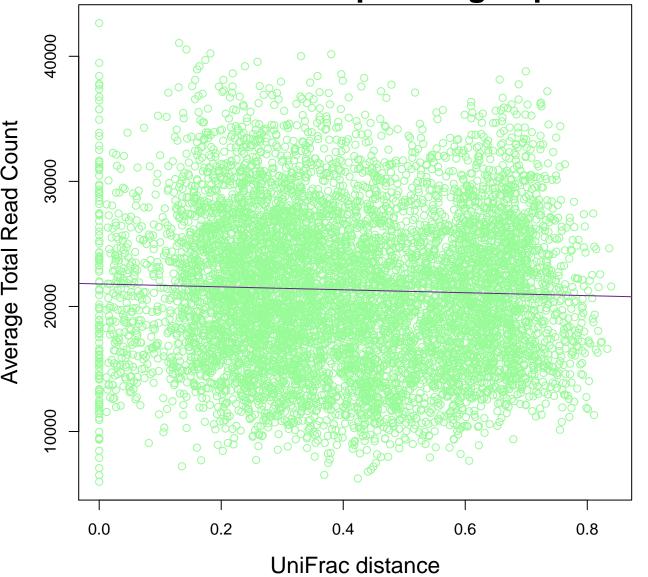
Proportional abundance weighted UniFrac vs. overlap



CLR transform weighted UniFrac vs. sequencing depth



Proportional abundance weighted UniFrac vs. sequencing depth



CLR transformed weighted UniFrac principle coordinates analysis L. crispatus _. iners bv -1.5-1.0-0.50.0 0.5 1.0 1.5 2.0 First Component 0.414 variance explained

Second Component 0.121 variance explained

0.5

Proportional abundance weighted UniFrac principle coordinates analysis 0.3 Second Component 0.115 variance explained L. crispatus . iners -0.20.0 0.2 0.4 First Component 0.634 variance explained