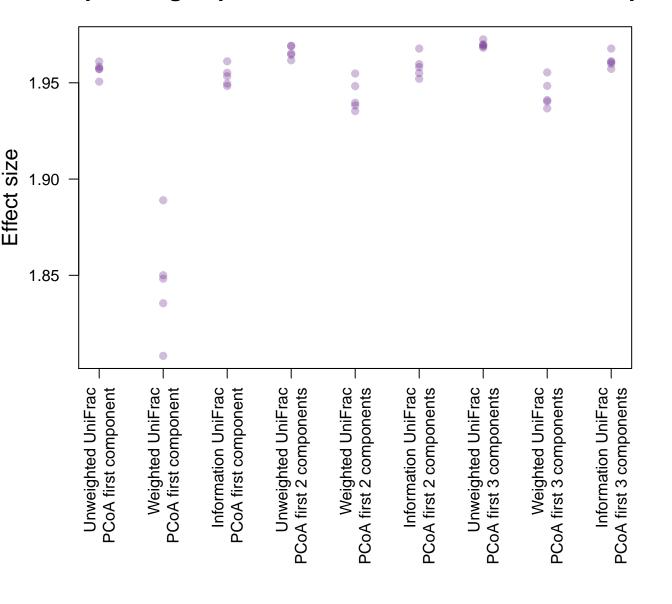
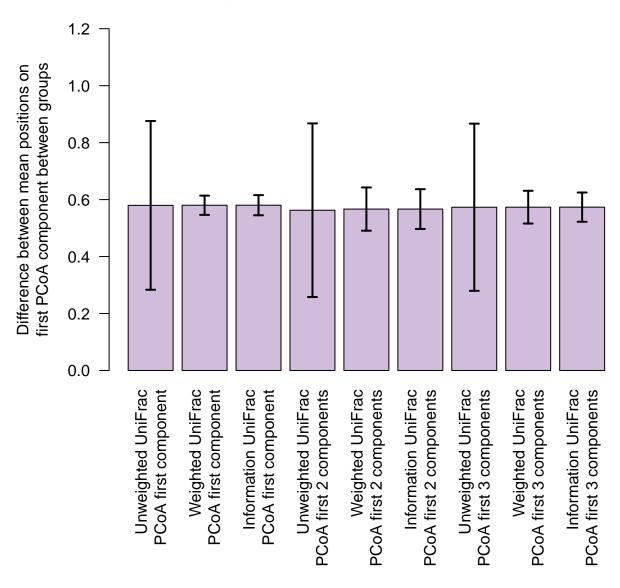
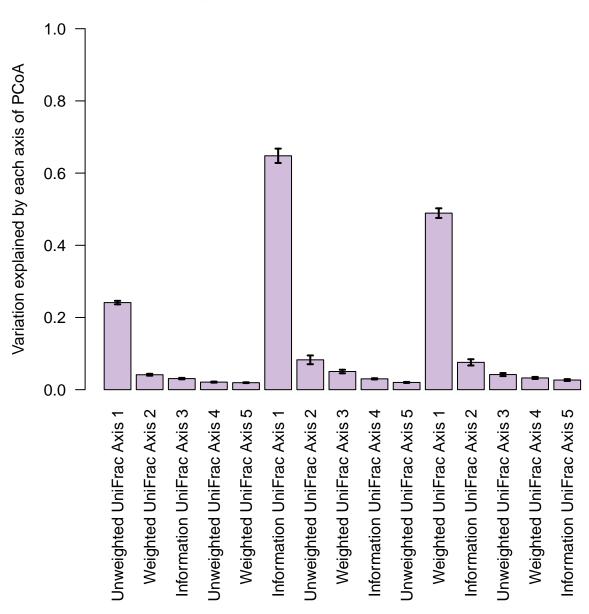
# Sequencing depth < 3000 vs. 3000-6000 reads/sample



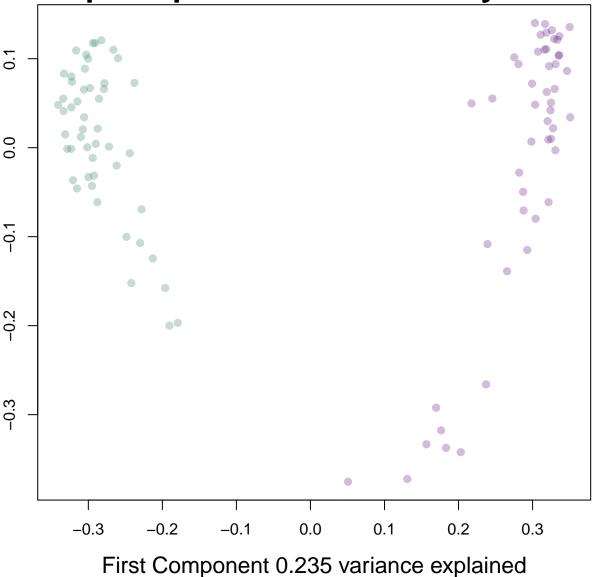
## Sequencing depth < 3000 vs. 3000-6000 reads/sample



## Sequencing depth < 3000 vs. 3000-6000 reads/sample



Unweighted UniFrac principal coordinates analysis



Second Component 0.043 variance explained

**Weighted UniFrac** principal coordinates analysis -0.4-0.20.2 0.0 0.4

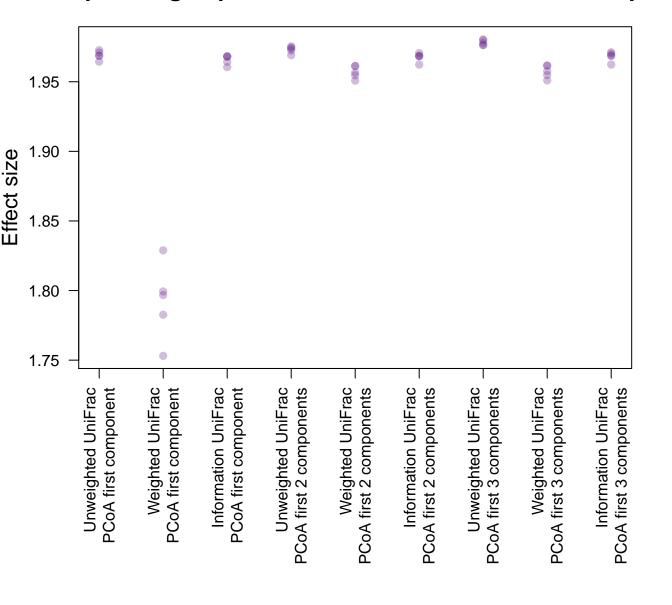
First Component 0.634 variance explained

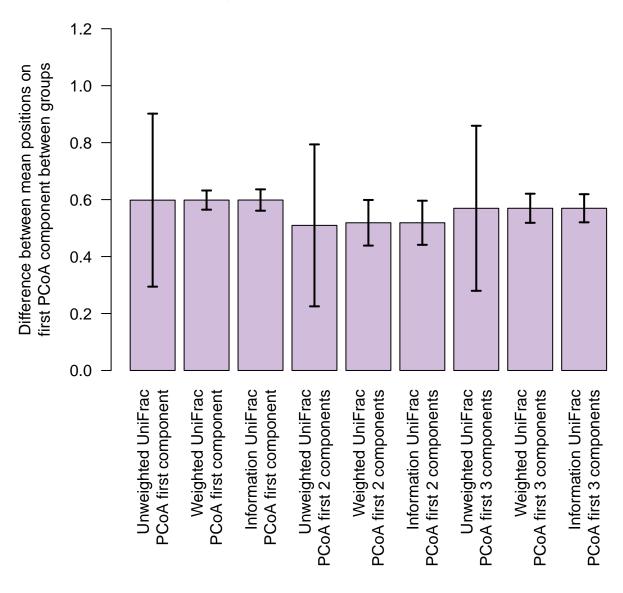
Second Component 0.096 variance explained

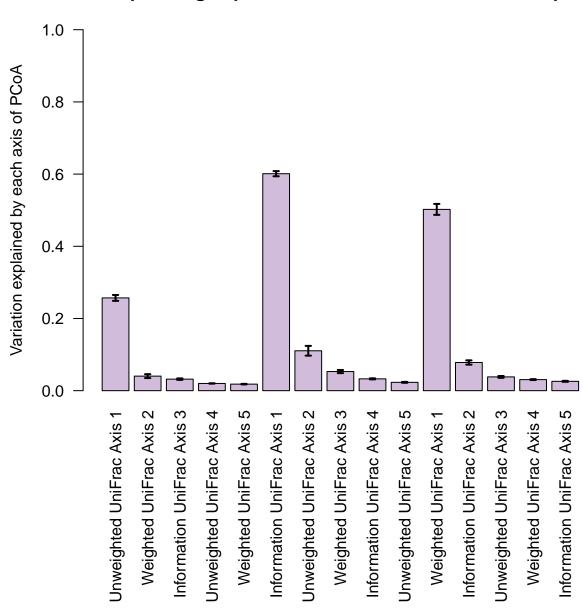
principal coordinates analysis Second Component 0.087 variance explained 0.1 -0.20.0 0.2 0.4

First Component 0.469 variance explained

**Information UniFrac** 







**Unweighted UniFrac** principal coordinates analysis -0.3-0.2-0.10.0 0.1 0.2 0.3

First Component 0.258 variance explained

Second Component 0.041 variance explained

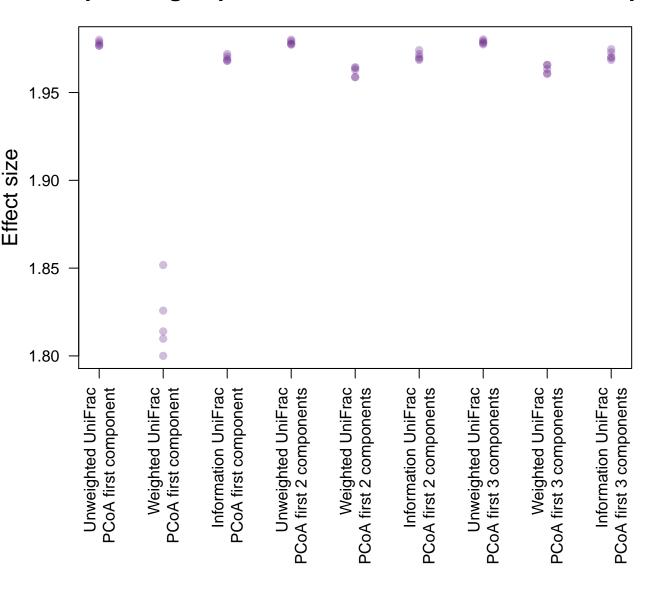
**Weighted UniFrac** principal coordinates analysis Second Component 0.111 variance explained 0.1 -0.20.0 0.2 0.4 First Component 0.6 variance explained

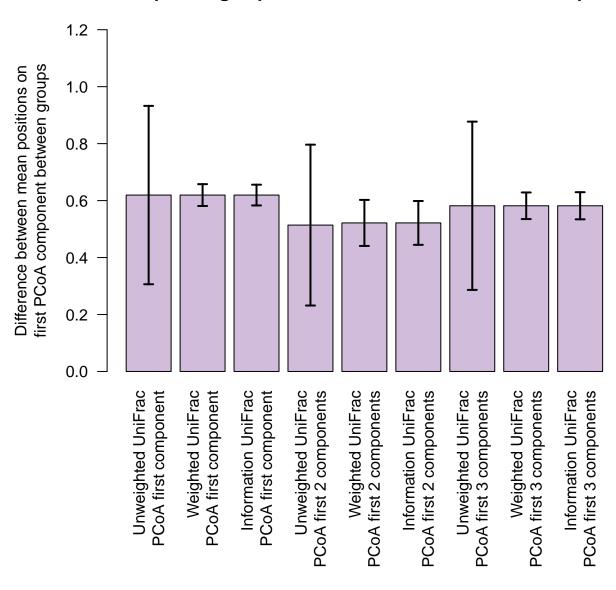
**Information UniFrac** principal coordinates analysis -0.3-0.2-0.10.0 0.1 0.2 0.3 0.4

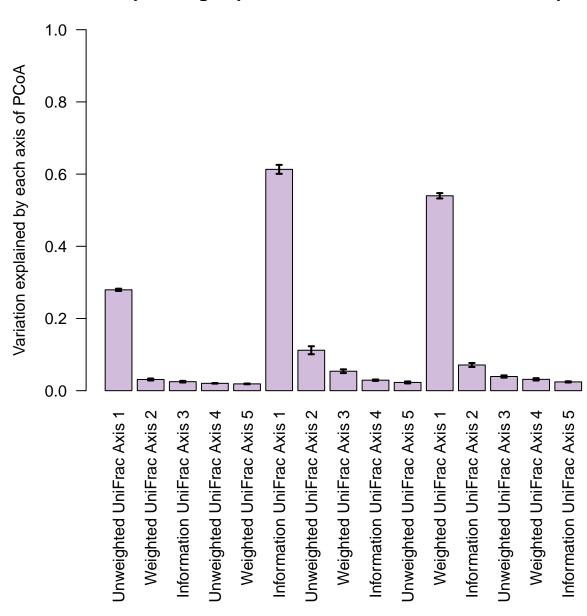
First Component 0.507 variance explained

Second Component 0.077 variance explained

0.1







**Unweighted UniFrac** principal coordinates analysis 0.2 -0.20.0 0.2 0.4 First Component 0.282 variance explained

Second Component 0.034 variance explained

**Weighted UniFrac** principal coordinates analysis -0.20.0 0.2 0.4

Second Component 0.121 variance explained

First Component 0.602 variance explained

**Information UniFrac** principal coordinates analysis 0.2 -0.20.0 0.4 0.2

First Component 0.545 variance explained

Second Component 0.069 variance explained