Effect of diversity distance = wunifrac; transformation = log 0.05 0.04 -PCoA Axis 2 (14.2 %) Axis 1 (46.7 %) 0.02 -0.00 -0.00 --0.05 --0.02 **-**-0.04 --0.10 **-**-0.10-0.050.00 LacZ 0.05 single 4 clones 24 clones ancestor clone PCoA Axis 1 (46.7 %)