Effect of diversity distance = bray; transformation = sqrt 0.2 -0.2 -PCoA Axis 2 (15.6 %) (24.2%)Axis **PCoA** -0.2 -

0.2

-0.4

LacZ

ancestor

single

clone

4 clones

24 clones

PCoA Axis 1 (24.2 %)

0.0

-0.2

-0.3 **-**

-0.4