**Notes corresponding to video** <http://tinyurl.com/mv690/video/permanova>









**library(vegan)**

**mrpp(measurements, group=soils, distance="bray")**

**adonis(measurements ~ soils + fert, distance="bray")**

**adonis(measurements ~ soils \* fert, distance="bray")**

**bc=bcdist(measurements)**

**pairwise.perm.manova(bc, soils, p.method="holm")**



