predict for labels = new.grps kendall 1.00 -0.75 x\_prefix RNAseq\_all\_genes rnaseq\_effector\_path\_vals rnaseq\_metabolic\_mod\_activities O.50 rnaseq\_metabolic\_mod\_genevalues rnaseq\_metabolic\_mod\_nodevalues rnaseq\_path\_vals rnaseq\_signaling\_genes 0.25 -0.00 -

x\_prefix