predict for labels = new.grps kendall rbf linear 1.00 -0.75 x\_prefix microarray\_all\_genes microarray\_effector\_path\_vals microarray\_metabolic\_mod\_activities 9 0.50 microarray\_metabolic\_mod\_genevalues microarray\_metabolic\_mod\_nodevalues microarray\_path\_vals microarray\_signaling\_genes 0.25 -0.00 x\_prefix