predict for labels = new.grps kendall 1.00 -0.75 x_prefix microarray_all_genes microarray_effector_path_vals microarray_metabolic_mod_activities o.50 microarray_metabolic_mod_genevalues microarray_metabolic_mod_nodevalues microarray_path_vals microarray_signaling_genes 0.25 -0.00 x_prefix