predict for labels = new.grps kendall linear 0.4 -0.3 x_prefix microarray_all_genes microarray_effector_path_vals microarray_metabolic_mod_activities acc microarray_metabolic_mod_genevalues microarray_metabolic_mod_nodevalues 0.2 microarray_path_vals microarray_signaling_genes 0.1 -0.0 x_prefix