IncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (69 / 2857) = (func/total lncRNA) ::: biotype = all_biotypes PageRank, range = (-0.1, 1.1)2.0 -1.5 factor(label) density 0.5 0.0 0.4 0.0 8.0 rank