lncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (25 / 277) = (func/total lncRNA) ::: biotype = all_biotypes PageRank, range = (-0.1, 1.1)2.0 1.5 factor(label) density 0.1 0.5 -0.0 -0.4 0.0 8.0 rank