lncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (42 / 1363) = (func/total lncRNA) ::: biotype = lincProcTrans PageRank, range = (-0.1, 1.1)3 · 2 · factor(label) density 1 -0 0.0 0.4 0.8 rank