lncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (25 / 243) = (func/total lncRNA) ::: biotype = all_biotypes PageRank, range = median +/- 2 ds 0 9 6 3 -2.0 factor(label) 0 1 1.5 -1.0 0.5 0.0 -8.0 1.2 1.6 0.4 rank