lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (47 / 2369) = (func/total IncRNA) ::: biotype = lincProcTrans PageRank, full range 0 300 200 100 factor(label) 0 0 20 10 -0 0.5 0.0 1.0 1.5 rank