lncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (42 / 1356) = (func/total IncRNA) ::: biotype = lincProcTrans PageRank, range = median ± -2 ds 0 60 -40 20 factor(label) 0 count 0 1 5 -4 -3 -2 -1 0 -8.0 1.2 1.6 0.0 0.4 rank