IncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (42 / 1363) = (func/total lncRNA) ::: biotype = lincProcTrans PageRank, full range 0 60 -40 -20 factor(label) 0 count 0 1 5 -4 -3 -2 -1 0 -0.5 1.0 1.5 0.0 rank