lncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (21 / 146) = (func/total lncRNA) ::: biotype = lincProcTrans PageRank, range = median +/- 2 ds 6 4 2 factor(label) 0 count 0 1 1.00 -0.75 -0.50 0.25 0.00 1.2 1.6 0.4 8.0 rank