lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (47 / 2369) = (func/total IncRNA) ::: biotype = lincProcTrans PageRank, range = median +/- 2 ds 0 300 -200 100 factor(label) 0 0 1 30 20 10 0 0

rank

count