lncRNA group = IDRlessthan0\_2 ::: data cols = bothPullDowns (48 / 2413 ) = (func/total IncRNA) ::: biotype = lincProcTrans PageRank, full range 0 300 -200 100 factor(label) 0 0 30 -20 10 -0 0.0 0.5 1.0 1.5 2.0

rank

count