IncRNA group = IDRlessthan0\_1 ::: data cols = bothPullDowns (42 / 1356) = (func/total lncRNA) ::: biotype = lincProcTrans PageRank, full range 0 60 -40 -20 factor(label) 0 0 1 5 -4 -3 -2 -1 -0 -1.5 0.5 1.0 0.0 rank

count