lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (82 / 5087) = (func/total IncRNA) ::: biotype = all_biotypes PageRank, full range 0 600 -400 200 factor(label) 0 0 30 20 10 -0 0.5 0.0 1.0 1.5 rank

count