lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (83/5164) = (func/total lncRNA) ::: biotype = all_biotypes PageRank, full range 600 400 200 factor(label) 0 0 40 -30 -20 10 0 0.5 1.0 1.5 0.0 2.0

rank

count