lncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (25 / 243) = (func/total lncRNA) ::: biotype = all_biotypes PageRank, full range 0 10.0 -7.5 5.0 -2.5 factor(label) 0.0 count 0 1 2.0 1.5 -1.0 0.5 0.0 0.8 1.2 0.4 1.6

rank