lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (17/331) = (func/total lncRNA) ::: biotype = processed_transcript PageRank, full range 40 20 factor(label) 0 0 no 12.5 0 10.0 -7.5 5.0 2.5 0.0 0.5 0.0 1.0 1.5 2.0

rank