lncRNA group = IDRlessthan0\_2 ::: data cols = bothPullDowns (27 / 1771) = (func/total IncRNA) ::: biotype = lincRNAPageRank, full range 0 150 100 50 factor(label) 0 0 10 -5 0 0.5

1.0

rank

1.5

0.0