IncRNA group = allLncRNA ::: data cols = bothPullDowns (52 / 4177) = (func/total IncRNA) ::: biotype = remove\_antisense PageRank, range = median +/- 2 ds 1250 1000 -750 -500 -250 factor(label) 0 0 40 -30 20 10 0 -0

rank

count