lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (27 / 1771) = (func/total IncRNA) ::: biotype = lincRNAlogRank, range = median +/- 2 ds 0 100 -75 -50 -25 factor(label) 0 0 1 6 4 2 0 -0.25 0.50 0.00 0.75 logRank

count