lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (81 / 4571) = (func/total lncRNA) ::: biotype = all_biotypes logRank, range = median +/- 2 ds 0 300 -200 100 factor(label) 0 count 0 1 30 -20 10 0 0.75 0.25 0.50 0.00 logRank