lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (50 / 2707) = (func/total lncRNA) ::: biotype = remove_antisense logRank, range = median +/- 2 ds 200 -150 -100 -50 factor(label) 0 count 0 1 25 20 -15 -10 -5 -0 -0.75 0.00 0.25 0.50 logRank