lncRNA group = IDRlessthan0\_2 ::: data cols = bothPullDowns (52 / 3409) = (func/total lncRNA) ::: biotype = remove\_antisense logRank, range = median +/- 2 ds 200 100 factor(label) 0 count 0 1 25 20 -15 -10 -5 -0 -0.25 0.50 0.00 0.75 logRank