lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (53 / 3463) = (func/total lncRNA) ::: biotype = remove_antisense logRank, range = median +/- 2 ds 300 -200 -100 factor(label) 0 0 1 20 10 -0 0.50 0.00 0.25 0.75 logRank