IncRNA group = IDRlessthan0\_2 ::: data cols = bothPullDowns (51 / 2754) = (func/total lncRNA) ::: biotype = remove\_antisense logRank, range = median +/- 2 ds 250 -200 -150 -100 -50 factor(label) 0 0 1 20 10 0 0.50 0.00 0.25 0.75

logRank