lncRNA group = IDRlessthan0\_1 ::: data cols = bothPullDowns (46 / 1947) = (func/total lncRNA) ::: biotype = remove\_antisense logRank, range = median +/- 2 ds1.5 -1.0 factor(label) density 0.5 -0.0 -0.6 8.0 1.0 0.2 0.4

logRank