lncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (46 / 1947) = (func/total lncRNA) ::: biotype = remove_antisense logRank, range = median +/- 2 ds0 40 -20 factor(label) 2.0 -0 1 1.5 -1.0 0.5 0.0 -0.4 0.6 0.8 1.0 0.2 logRank