lncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (45 / 1547) = (func/total lncRNA) ::: biotype = remove_antisense logRank, range = median +/- 2 ds0 40 -30 -20 -10 -2.0 7 factor(label) 0 1.5 -1.0 -0.5 -0.0 -0.6 8.0 0.2 0.4 1.0 logRank