lncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (45 / 1554) = (func/total lncRNA) ::: biotype = remove_antisense logRank, range = median +/- 2 ds0 40 30 -20 -10 conut 2.0 factor(label) 0 1.5 -1.0 -0.5 -0.0 -0.6 0.2 0.4 8.0 1.0 logRank