lncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (46 / 1955) = (func/total lncRNA) ::: biotype = remove_antisense logAbsRank, range = all points 0.4 -0.3 factor(label) density 0.2 0.1 -0.0 --8 -6 -2 0 log(abs(rank))