lncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (45 / 1554) = (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))