IncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (81 / 4571) = (func/total lncRNA) ::: biotype = all_biotypes logAbsRank, range = all points 0.3 -0.2 factor(label) density 0.1 -0.0 --7.5 -5.0 -2.5 -10.00.0 log(abs(rank))