IncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (30 / 1701) = (func/total IncRNA) ::: biotype = antisense logAbsRank, range = all points 0.2 factor(label) density 0.1 -0.0 --5.0 -2.5 -7.5 0.0 log(abs(rank))