IncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (83 / 5164) = (func/total IncRNA) ::: biotype = all_biotypes logAbsRank, range = all points 0.2 factor(label) density 0.1 -0.0 --5.0 -2.5 -10.0-7.5 0.0 2.5

log(abs(rank))