lncRNA group = allLncRNA ::: data cols = bothPullDowns (38 / 2538) = (func/total IncRNA) ::: biotype = antisense logAbsRank, range = all points 0.20 0.15 factor(label) density 0.10 -0.05 -0.00 --9 -6 -3 0 log(abs(rank))