IncRNA group = allLncRNA ::: data cols = bothPullDowns (27 / 2546) = (func/total lncRNA) ::: biotype = lincRNA logAbsRank, range = all points 0.25 -0.20 0.15 factor(label) 0.10 -0.05 -0.00 --3 -9 -6 0

log(abs(rank))

density