IncRNA group = IDRlessthan0_2 ::: data cols = Inpa (30 / 1996) = (func/total IncRNA) ::: biotype = lincRNA logAbsRank, range = all points 0.20 -0.15 factor(label) 0.10 0.05 0.00 -3 -9 -6 0

log(abs(rank))

density