IncRNA group = IDRlessthan0_2 ::: data cols = Inpa (27 / 1732) = (func/total IncRNA) ::: biotype = lincRNA logAbsRank, range = all points 0.25 -0.20 0.15 factor(label) 0.10 0.05 -0.00 -

-3

log(abs(rank))

0

density

-9

-6