IncRNA group = allLncRNA ::: data cols = Inpa (88 / 6668) = (func/total lncRNA) ::: biotype = all\_biotypes logAbsRank, range = all points 0.25 -0.20 0.15 factor(label) 0.10 0.05 -

density

0.00

-10.0

-7.5

-5.0

log(abs(rank))

-2.5

0.0