IncRNA group = IDRlessthan0\_2 ::: data cols = Inpa (47 / 2308) = (func/total lncRNA) ::: biotype = lincProcTrans logAbsRank, range = all points 0.20 0.15 factor(label) 0.10 0.05 0.00

<u>-3</u>

log(abs(rank))

0

-6

density