lncRNA group = IDRlessthan0_01 ::: data cols = lpa (9 / 81) = (func/total IncRNA) ::: biotype = lincRNA logAbsRank, range = all points factor(label) _3 <u>-2</u> -1 log(abs(rank)) 0

0.4 -

0.3

density

0.1 -

0.0 -