lncRNA group = IDRlessthan0\_1 ::: data cols = lpa (69 / 2856) = (func/total lncRNA) ::: biotype = all\_biotypes logAbsRank, range = all points 0.4 -0.3 factor(label) density 0.2 0.1 -

-2

log(abs(rank))

0

0.0 -

-8

-6