lncRNA group = IDRlessthan0_1 ::: data cols = lpa (69 / 2573) = (func/total lncRNA) ::: biotype = all_biotypes logAbsRank, range = all points 0.4 0.3 factor(label) density 0.1

-2

log(abs(rank))

0

0.0 -

-8

-6