lncRNA group = IDRlessthan0_1 ::: data cols = Inpa (69 / 2559) = (func/total lncRNA) ::: biotype = all_biotypes logAbsRank, range = all points 0.3 -0.2 factor(label) 0.1 -0.0 --7.5 -5.0 -2.5 0.0

log(abs(rank))

density