IncRNA group = allLncRNA ::: data cols = Inpa (36 / 2293) = (func/total IncRNA) ::: biotype = antisense logAbsRank, range = all points 0.25 0.20 0.15 factor(label) 0.10 0.05 0.00 -5.0 -2.5 0.0

log(abs(rank))

density