lncRNA group = IDRlessthan0\_1 ::: data cols = Inpa (23/935) = (func/total IncRNA) ::: biotype = lincRNAlogAbsRank, range = all points factor(label) -7.5 -5.0 -2.5 0.0

log(abs(rank))

0.2 -

density

0.1 -

0.0 -