IncRNA group = IDRlessthan0\_2 ::: data cols = bothPullDowns (48 / 2413) = (func/total IncRNA) ::: biotype = lincProcTrans logAbsRank, range = all points 0.2 factor(label) 0.1 -0.0 -

-2.5

0.0

2.5

-5.0

log(abs(rank))

density

-7.5

-10.0