IncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (47 / 2369) = (func/total IncRNA) ::: biotype = lincProcTrans logAbsRank, range = all points 0.3 -0.2 factor(label) 0.1 0.0 -10.0-7.5 -2.5 -5.00.0 log(abs(rank))

density