IncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (4/72) = (func/total IncRNA) ::: biotype = antisense expression accross all samples(cell-type/pulldown) false withinSubset false 0.25 0.50 0.75 1.00 0.25 0.50 0.75

JSD for each gene

1.00

2.5

2.0

1.5 -

1.0 -

0.5 -

0.0 -

density