IncRNA group = IDRlessthan0_1 ::: data cols = Inpa (26 / 1119) = (func/total IncRNA) ::: biotype = lincRNA expression accross all samples(cell-type/pulldown) false withinSubset false true

0.25

JSD for each gene

0.50

0.75

2.0

1.5

0.5 -

0.0 -

0.25

0.50

0.75