IncRNA group = IDRlessthan0_01 ::: data cols = lpa (25 / 243) = (func/total IncRNA) ::: biotype = all_biotypes expression accross all samples(cell-type/pulldown) false withinSubset false true

2.5

2.0

1.5 -

1.0 -

0.5 -

0.0 -

0.25

0.50

0.75

1.00

JSD for each gene

0.25

0.50

0.75

1.00

density