IncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (4/72) = (func/total IncRNA) ::: biotype = antisensemeasures accross all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 6 3 4 2 2 1 · 0.025 -40 60 1.5 20 6 2.0 9 0.0 0.5 1.0 3 0 0 nTimesRest minExpr maxExpr 0.3 -0.6 withinSubset 0.2 0.4 false true 0.1 0.2 -0.0 0.0 5 10 200 400 0.01 0.02 0.00 0.03 0 600 0 tissSpec

0.125

0.100

0.075

0.050

0.000

900 -

600

300

0

2.5 -

2.0

1.5

1.0 -

0.5 -

0.0 -

0.25

0.50

0.75

1.00

density

value of facet label