IncRNA group = allLncRNA ::: data cols = lpa (37 / 2359) = (func/total IncRNA) ::: biotype = antisense measures accross all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 200 30000 -150 -20000 -100 10000 -50 -0 0 40 60 0.5 1.5 20 3 6 9 0.0 1.0 2.0 0 nTimesRest minExpr maxExpr 30 -0.6 withinSubset 20 0.4 false true 10 -0.2 0.0 0 5 0.01 0.02 200 400 0.03 10 600 0 0 tissSpec

8

6 -

4 -

2 -

1500

density

500

0

2.0

1.5

1.0

0.5 -

0.0 -

0.25

0.50

0.75

1.00

0.00

0