IncRNA group = IDRlessthan0_2 ::: data cols = lpa (30 / 1675) = (func/total IncRNA) ::: biotype = antisense measures accross all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 12000 150 9000 -100 6000 -50 3000 -0 0 40 60 0.5 1.5 20 3 6 9 1.0 2.0 0.0 0 minExpr maxExpr nTimesRest 15 -0.6 10 withinSubset 0.4 false true 5 0.2 -0.0 0 5 0.01 0.02 200 400 0.03 10 600 0 0 tissSpec

4 -

2

1000

500

0.00

3

2 -

1 -

0

0.25

0.50

0.75

1.00

density

0

value of facet label