IncRNA group = IDRlessthan0\_01 ::: data cols = lpa (11/109) = (func/total IncRNA) ::: biotype = lincRNAmeasures accross all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 6 7.5 -4 5.0 -2 2.5 -0.0 0 100 150 3 50 10 2 5 5 0 0 4 minExpr maxExpr nTimesRest 0.8 -0.75 0.6 withinSubset 0.50 false 0.4 true 0.25 -0.2 -0.00 0.0 5 0.2 0.3 10 100 15 200 0.1 0 0 300 tissSpec

0.20 -

0.15

0.10

0.05

0.00

3000 -

density 2000

1000 -

0

2.0

1.5

1.0

0.5 -

0.0 -

0.25

0.50

0.75

0.0

0

value of facet label