IncRNA group = allLncRNA ::: data cols = lpa (37 / 2359) = (func/total IncRNA) ::: biotype = antisense measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.15 0.075 0.10 -0.050 -0.05 0.025 0.000 0.00 0 и -5 -5 -20 -15 -10 -5 -10 0 5 minExpr nTimesRest maxExpr 0.15 -0.9 withinSubset 0.10 -0.6 false true 0.05 0.3 0.00 0.0 -4 -9 0 2 -7 -5 -3 -8 4 6 0 tissSpec

0.15 -

0.10 -

0.05

0.00

0.20

density 0.10 -

0.05

0.00

0.8 -

0.6 -

0.4 -

0.2

0.0

-2.0 -1.5 -1.0 -0.5 0.0

-11

-10

value of facet label