IncRNA group = IDRlessthan0_2 ::: data cols = lpa (31 / 1860) = (func/total IncRNA) ::: biotype = antisense measures accross all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 150 5 10000 -4 -7500 -100 3 5000 2 -50 2500 1 -0 0 60 0.5 1.5 2.0 20 40 3 6 9 0.0 1.0 0 0 minExpr maxExpr nTimesRest 0.6 1000 10 withinSubset 0.4 false true 500 5 0.2 -0.0 0 5 0.01 0.02 200 400 0.03 10 0.00 0 600 0 tissSpec 3 2 ·

density

1 -

0

0.25

0.50

0.75

1.00

value of facet label