lncRNA group = IDRlessthan0_1 ::: data cols = lpa (23 / 902) = (func/total IncRNA) ::: biotype = antisense measures accross all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 60 2.0 2000 1.5 -1500 40 1.0 -1000 20 -0.5 500 0.0 0 40 6 2.0 20 60 9 0.0 0.5 1.0 1.5 3 0 maxExpr minExpr nTimesRest 8 800 -0.6 6 -600 withinSubset density 000 0.4 false 4 · true 0.2 -2 · 200 0 0 -0.0 т 5 10 200 400 0.02 Ò 0.00 0.01 0.03 600 0 tissSpec 3 2 · 1 -

value of facet label

0

0.25

0.50

0.75

1.00