IncRNA group = IDRlessthan0_01 ::: data cols = lpa (4 / 72) = (func/total IncRNA) ::: biotype = antisense measures accross all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.15 6 2.0 -0.10 1.5 4 1.0 0.05 2 -0.5 0.00 0 0.0 60 40 20 8 2 12 3 0 4 0 4 minExpr maxExpr nTimesRest 12 -0.15 -0.6 withinSubset 8 density 0.10 -0.4 false true 4 -0.05 0.2 -0 -0.0 0.00 1.0 5 0.5 1.5 10 0.0 0 2000 4000 6000 0 tissSpec 2.0 -1.5 1.0 0.5

value of facet label

0.0

0.25

0.50

0.75

1.00