lncRNA group = IDRlessthan0_2 ::: data cols = lpa (30 / 1675) = (func/total IncRNA) ::: biotype = antisense measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.125 -0.20 -0.100 -0.15 -0.075 -0.10 -0.050 -0.05 0.025 0.000 0.00 -7.5 -5.0 -2.5 0.0 -10 -20 -15 2.5 0 -8 -4 0 maxExpr minExpr nTimesRest 0.25 0.20 -0.9 withinSubset 0.15 -0.6 false 0.10 true 0.3 0.05 0.00 0.0 -3 2 0 <u>-</u>9 -7 -6 3 -5 -3 4 6 -11 0 tissSpec

0.20 -

0.15 -

0.10 -

0.05

0.00

0.20

density 0.10 -

0.05

0.00

0.6 -

0.4 -

0.2

0.0

-2.0 -1.5 -1.0 -0.5 0.0

value of facet label