lncRNA group = IDRlessthan0_2 ::: data cols = lpa (31 / 2051) = (func/total lncRNA) ::: biotype = lincRNA measures across all samples(cell-type/pulldown) varianceExpr sumExpr averageExpr 0.12 -0.20 0.15 0.08 -0.10 -0.10 0.04 -0.05 0.00 0.00 0 -20 -15 -10 -5 <u>-4</u> Ó -5 5 0 -8 minExpr nTimesRest maxExpr 0.25 -0.3 -1.00 -0.20 -0.75 withinSubset 0.15 false 0.50 0.10 true 0.25 0.05 -0.00 0.00 -3 0 2 -10.0 -7.5 -5.0 -6 3 -2.56 0 tissSpec 0.4 -

0.20

0.15

0.05

0.00

0.2

0.1

0.0

8.0

0.6

0.2

0.0

-2.0 -1.5 -1.0 -0.5 0.0

density

value of facet label