IncRNA group = allLncRNA ::: data cols = Inpa (30 / 2803) = (func/total lncRNA) ::: biotype = lincRNA measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.15 -0.075 -0.10 -0.050 -0.05 0.025 0.000 0.00 <del>-</del>5 <u>-</u>5 -20 -15 -10 -5 Ö 0 0 -10 5 minExpr nTimesRest maxExpr 1.00 -0.15 0.75 withinSubset 0.10 false 0.50 true 0.05 -0.25 -0.00 0.00 -4 -10.0 -7.5 -5.0 Ö -8 2 -2.5 6 0 tissSpec

0.15

0.10 -

0.05

0.00

0.3 -

0.2

0.1

0.0

8.0

0.6

0.4 -

0.2

0.0

-2.0 -1.5 -1.0 -0.5 0.0

density

-10

value of facet label