IncRNA group = allLncRNA ::: data cols = Inpa (36 / 2219) = (func/total IncRNA) ::: biotype = antisense 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 0 \_<del>.</del>5 -20 -15 -10 -5 -<del>1</del>0 -5 -10 0 5 minExpr nTimesRest maxExpr 0.9 0.15 withinSubset 0.6 -0.10 false true 0.05 0.3 0.00 0.0 -9 2 -7 . –8 Ö -5 -4 3 -3 0 -11 tissSpec

0.15

0.10

0.05

0.00

0.20

density 0.10 -

0.05

0.00

0.8 -

0.6 -

0.4 -

0.2 -

0.0

-2.0 -1.5 -1.0 -0.5 0.0

value of facet label