IncRNA group = allLncRNA ::: data cols = bothPullDowns (39 / 2750) = (func/total IncRNA) ::: biotype = antisense measures across all samples(cell-type/pulldown) varianceExpr sumExpr averageExpr 0.15 -0.075 -0.10 0.050 -0.05 -0.025 0.000 0.00 0 -20 -15 -10 -5 _5 Ö -8 -4 -10 -12 0 minExpr maxExpr nTimesRest 1.00 -0.15 -0.75 withinSubset 0.10 false 0.50 true 0.05 -0.25 -0.00 0.00 Ö -4 5.0 -10.0 -7.5 -5.0 -2.5 -8 0.0 2.5 7.5 tissSpec

0.15 -

0.10

0.05

0.00

0.20

density 0.10 -

0.05

0.00

0.75

0.50 -

0.25

0.00

-2.0 -1.5 -1.0 -0.5 0.0

value of facet label