lncRNA group = IDRlessthan0_2 ::: data cols = Inpa (30 / 1652) = (func/total IncRNA) ::: biotype = antisense measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.20 -0.100 -0.15 0.075 -0.10 -0.050 0.05 0.025 0.000 0.00 -20 -15 -10 -8 -5 0 0 -12 -4 0 minExpr maxExpr nTimesRest 0.20 -0.9 withinSubset 0.15 -0.6 false 0.10 true 0.3 0.05 0.00 0.0 -3 -9 -6 2 Ö 3 -7 -5 3 -3 0 -11 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