lncRNA group = IDRlessthan0_2 ::: data cols = lpa (82 / 4574) = (func/total lncRNA) ::: biotype = all_biotypes measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.125 0.20 -0.100 -0.15 0.075 -0.10 -0.050 -0.05 -0.025 -0.000 0.00 0 -10 . –4 Ö -8 -5 -20 -12 0 5 maxExpr minExpr nTimesRest 1.00 -0.20 0.75 withinSubset 0.15 false 0.50 -0.10 true 0.25 -0.05 0.00 0.00 -5 0 т 5 2 -6 -3 _9 6 -10 0 4 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