lncRNA group = IDRlessthan0_2 ::: data cols = lpa (48 / 2379) = (func/total lncRNA) ::: biotype = lincProcTrans 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 -4 0 -10 Ö -8 Ö -5 5 -20 -12 minExpr maxExpr nTimesRest 0.20 -0.75 withinSubset 0.15 -0.50 false 0.10 true 0.25 0.05 -0.00 0.00 -5 0 т 5 -10.0 -7.5 -5.0 2 -2.5 -10 0 6 tissSpec

0.20

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