IncRNA group = allLncRNA ::: data cols = lpa (90 / 6451) = (func/total IncRNA) ::: biotype = all\_biotypes 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 0 5 –10 -5 -5 -10 0 -20 -10 minExpr maxExpr nTimesRest 0.15 -0.75 withinSubset 0.10 -0.50 false true 0.05 -0.25 -0.00 0.00 -5 0 5 -4 -8 2.5 5.0 7.5 -10 0.0 tissSpec

0.15 -

0.10 -

0.05

0.00

0.20

0.15 -

0.05

0.00

1.00

0.75 -

0.50

0.25

0.00

-2.0 -1.5 -1.0 -0.5

density