IncRNA group = IDRlessthan0_01 ::: data cols = Inpa (25 / 242) = (func/total lncRNA) ::: biotype = all_biotypes measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.3 -0.20 0.2 0.15 0.10 0.1 -0.05 0.00 0.0 0 2 0 -2 5 -2.5 4 _10 -5 -5.0 0.0 2.5 minExpr nTimesRest maxExpr 0.5 0.5 0.4 0.4 withinSubset 0.3 0.3 false 0.2 0.2 true 0.1 0.1 0.0 2.5 0 2 -7.5 -5.0 -2.5 -2 5.0 4 7.5 0.0 0.0 tissSpec

0.3 -

0.2

0.1 -

0.0

0.25

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