lncRNA group = IDRlessthan0\_1 ::: data cols = Inpa (26 / 1119) = (func/total lncRNA) ::: biotype = lincRNA measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.3 0.15 -0.2 0.10 -0.1 0.05 -0.00 0.0 0.0 -5 -**1**0 <u>-</u>2 Ö -2.5 2.5 -6 -4 5.0 0 -8 maxExpr minExpr nTimesRest 0.3 -0.75 withinSubset 0.2 -0.50 false true 0.1 -0.25 0.0 0.00 0.0 -5.0 2.5 2 -7.5 -2.5 -10.0 -2.5-5.0 4 6 0 tissSpec

0.3

0.2

0.1

0.0

0.3

density 0.2

0.0

8.0

0.6

0.4 -

0.2

0.0

-2.0 -1.5 -1.0 -0.5

-5.0