lncRNA group = IDRlessthan0\_2 ::: data cols = Inpa (81 / 4501) = (func/total lncRNA) ::: biotype = all\_biotypes measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.20 0.09 -0.15 0.06 -0.10 -0.03 -0.05 0.00 0.00 –10 0 . –4 -5 -20 0 -12 -8 0 5 minExpr maxExpr nTimesRest 0.20 -0.75 0.15 withinSubset 0.50 false 0.10 true 0.25 0.05 -0.00 0.00 0 -4 -5 2.5 -8 5 5.0 7.5 0 -10 0.0 tissSpec

0.20

0.15

0.10

0.05 -

0.00

0.20

0.15 -

0.05

0.00

0.6 -

0.4 -

0.2

0.0

-2.0 -1.5 -1.0 -0.5

density

value of facet label