lncRNA group = IDRlessthan0\_2 ::: data cols = Inpa (47 / 2324) = (func/total lncRNA) ::: biotype = lincProcTrans measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.12 -0.20 0.09 -0.15 0.06 -0.10 0.03 -0.05 0.00 0.00 0 -10 -5 <u>-</u>5 Ö **-10** 0 -10 -20 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 5 0 2 -10.0 -<del>7.5</del> -<del>5.0</del> -5 -2.5 -10 6 0 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

value of facet label