IncRNA group = allLncRNA ::: data cols = Inpa (87 / 6034) = (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 5 0 -10 -5 -5 Ö **-10** -20 minExpr maxExpr nTimesRest 1.00 -0.15 -0.75 withinSubset 0.10 false 0.50 true 0.05 0.25 0.00 0.00 -5 0 5 2 -6 -3 <u>-</u>9 4 -10 0 tissSpec

0.15

0.10

0.05

0.00

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 0.0

-10

value of facet label