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 0.12 -0.20 0.09 0.15 0.06 -0.10 0.03 -0.05 0.00 0.00 -10 0 и -5 -5 Ö **-10** 0 -10 -20 5 minExpr maxExpr nTimesRest 1.00 -0.20 -0.75 withinSubset 0.15 false 0.50 0.10 true 0.25 0.05 -0.00 0.00 2 0 -6 <u>-</u>3 -5 <u>-</u>9 5 4 -10 0 tissSpec 0.4 -

0.20

0.15

0.10

0.05

0.00

0.20

density 0.10

0.05

0.00

0.6 -

0.2

0.0

-2.0 -1.5 -1.0 -0.5 0.0

value of facet label