lncRNA group = IDRlessthan0_2 ::: data cols = lpa (31 / 1860) = (func/total IncRNA) ::: biotype = antisense measures across all samples(cell-type/pulldown) varianceExpr sumExpr averageExpr 0.20 0.20 0.12 -0.15 0.15 0.09 -0.10 0.10 0.06 -0.05 0.05 0.03 -0.00 0.00 0.00 ò -20 -15 -10 -5 <u>-4</u> 0 -12 -8 0 minExpr maxExpr nTimesRest 0.20 0.20 -0.75 density 0.10 withinSubset 0.15 -0.50 false 0.10 true 0.25 0.05 0.05 -0.00 0.00 0.00 -3 0 -6 -10.0 -7.5 -5.0 -2.5 0.0 2.5 5.0 3 0.0 7.5 tissSpec 0.6 0.4 -

value of facet label

0.2 -

0.0

-2.0 -1.5 -1.0 -0.5

0.0