lncRNA group = IDRlessthan0_1 ::: data cols = Inpa (23 / 892) = (func/total IncRNA) ::: biotype = antisense measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.3 0.3 0.15 -0.2 -0.2 0.10 -0.1 -0.1 0.05 -0.0 0.00 0.0 0.0 -5 ò -2.5 2.5 **-10** -4 -2 -6 0 minExpr maxExpr nTimesRest 1.00 -0.20 0.3 -0.75 density 0.10 withinSubset 0.2 false 0.50 true 0.1 -0.25 0.05 0.00 0.0 0.00 0.0 3 <u>-</u>9 _.7 -2.5 2 4 -5 2.5 -3 -5.0-11 0 tissSpec 0.8 -0.6 -0.4 -

value of facet label

0.2 -

0.0

-2.0 -1.5 -1.0 -0.5 0.0