IncRNA group = IDRlessthan0\_01 ::: data cols = Inpa (4/71) = (func/total IncRNA) ::: biotype = antisense measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.3 0.3 0.2 -0.2 -0.2 -0.1 0.1 0.1 -0.0 0.0 1 –4 2 <u>-2</u> 0 <u>-2</u> -8 Ó 0 -6 -4 minExpr nTimesRest maxExpr 0.3 0.6 -0.4 density 0.1 withinSubset 0.4 false 0.2 true 0.2 -0.0 -0.0 0.0 0 2 -5.0 -2.5 . –2 2.5 5.0 -7.5 0.0 0.0 7.5 tissSpec 0.6 0.4 0.2

value of facet label

0.0 -

-2.0 -1.5 -1.0 -0.5

0.0