IncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (5/65) = (func/total IncRNA) ::: biotype = antisensemeasures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.25 0.3 -0.20 0.15 -0.2 0.10 0.1 -0.05 -0.00 0.0 3 2 –2.5 -5.0 0.0 _2 _1 Ö <u>-3</u> 2.5 maxExpr minExpr nTimesRest 0.6 -0.4 withinSubset 0.4 false 0.2 true 0.2 0.0 0.0 0 2 2 _7 <u>-</u>9 -5 -1 4 -3 0 6 tissSpec

0.3 -

0.2 -

0.1 -

0.0

0.25 -

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

value of facet label