IncRNA group = IDRlessthan0_01 ::: data cols = Ipa (11 / 109) = (func/total IncRNA) ::: biotype = lincRNA measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.3 -0.2 -0.2 0.1 -0.1 -0.0 0.0 т –2 2 -2.5 0.0 -5.0 0 2.5 0 4 maxExpr nTimesRest minExpr 0.6 0.4 withinSubset 0.4 false 0.2 true 0.2 0.0 0.0 2 2 -5.0 6 -2.5 -1 0 4 -7.5 0.0 -2 0 tissSpec

0.3 -

0.2

0.1 -

0.0

0.15 -

density

0.05

0.00

0.6

0.4

0.2

0.0 -

-1.0

-0.5

0.0

-1.5

value of facet label