IncRNA group = IDRIessthan0_01 ::: data cols = Inpa (11 / 109) = (func/total IncRNA) ::: biotype = lincRNA 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 0.0 т –2 2 -2.5 0.0 -5.0 0 2.5 0 4 maxExpr nTimesRest minExpr 0.6 0.15 -0.4 withinSubset density 0.4 false 0.2 true 0.05 0.2 0.00 0.0 0.0 2 2 -5.0 6 -2.5 -1 0 4 -7.5 0.0 -2 0 tissSpec 0.6 0.4 0.2

value of facet label

0.0 -

-1.0

-0.5

0.0

-1.5