IncRNA group = IDRlessthan0_01 ::: data cols = Inpa (4 / 71) = (func/total IncRNA) ::: biotype = antisense measures accross all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.15 6 -2.0 0.10 1.5 -4 -1.0 -0.05 2 0.5 -0.00 0 0.0 40 60 3 20 8 12 2 0 0 4 0 4 minExpr maxExpr nTimesRest 12.5 0.6 10.0 0.3 withinSubset 7.5 0.4 false 0.2 -5.0 true 0.2 0.1 -2.5 0.0 0.0 0.0 1.0 5 10 0.5 1.5 0.0 0 2000 4000 6000 0 tissSpec 2.0 -1.5 1.0 0.5

density

0.0

0.25

0.50

0.75

1.00

value of facet label