IncRNA group = IDRlessthan0_01 ::: data cols = Inpa (25 / 242) = (func/total lncRNA) ::: biotype = all_biotypes measures accross all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 12.5 2.5 0.15 -10.0 2.0 7.5 0.10 1.5 5.0 -1.0 -0.05 2.5 -0.5 -0.00 0.0 0.0 10 100 150 200 250 1000 2000 3000 4000 5 15 50 0 minExpr maxExpr nTimesRest 0.6 -0.075 20 0.4 withinSubset 15 0.050 false 10 true 0.2 0.0255 0 -0.0 0.000 0.5 1.0 1.5 2.0 50 100 150 200 250 2000 4000 0.0 0 6000 tissSpec 2.0 1.5 1.0

density

0.5

0.0

0.25

0.50

0.75

1.00

value of facet label