lncRNA group = IDRlessthan0\_2 ::: data cols = Inpa (30 / 1996) = (func/total lncRNA) ::: biotype = lincRNA measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.125 0.20 0.20 0.100 -0.15 0.15 -0.075 -0.10 0.10 0.050 -0.05 0.05 -0.025 -0.00 0.000 0.00 -5 -20 -15 -10 -5 \_5 **-10** Ö -10 0 5 0 minExpr nTimesRest maxExpr 0.3 -1.00 -0.20 -0.75 -0.2 withinSubset 0.15 false 0.50 -0.10 true 0.1 0.25 0.05 0.0 0.00 0.00 -4 0 -10.0 -7.5 -5.0 2 -2.5-8 6 0 tissSpec 0.8 0.6 0.4 -

density

0.2

0.0

-2.0 -1.5 -1.0 -0.5 0.0

value of facet label