lncRNA group = IDRlessthan0_1 ::: data cols = lpa (26 / 1131) = (func/total lncRNA) ::: biotype = lincRNA measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.15 -0.2 -0.2 0.10 -0.1 0.1 0.05 -0.0 0.00 0.0 -5 0 0.0 -3 -10 Ö -5.0 -2.5 2.5 -15 <u>-</u>6 -9 nTimesRest minExpr maxExpr 0.3 0.5 0.20 -0.4 -0.15 -0.2 withinSubset density 0.10 -0.3 false 0.2 true 0.1 0.05 0.1 0.00 0.0 0.0 -5.0 -2.5 0.0 6 -10.0 -7.5 -5.0 -2.52.5 2 8 0.0 0 tissSpec 0.6 -0.4 -0.2

value of facet label

0.0

-2.0 -1.5 -1.0 -0.5