IncRNA group = IDRlessthan0\_2 ::: data cols = bothPullDowns  $(31 \overline{/} 2082) = (func/total IncRNA) ::: biotype = lincRNA$ measures across all samples(cell-type/pulldown) varianceExpr sumExpr averageExpr 0.12 -0.20 0.150.08 -0.10 0.04 -0.05 0.00 0.00 -5 -20 -15 -10 -5 -5 -<del>1</del>0 -10 0 5 0 0 minExpr nTimesRest maxExpr 0.25 1.00 -0.20 -0.75 withinSubset 0.15 false 0.50 0.10 true 0.25 0.05 -0.00 0.00 -4 2 -10.0 -7.5 -5.0 Ö -2.5-8 6 0 tissSpec

0.20

0.15

0.10

0.05

0.00

0.3 -

0.2

0.1

0.0

0.8 -

0.6 -

0.4 -

0.2 -

0.0

-2.0 -1.5 -1.0 -0.5 0.0

density

value of facet label