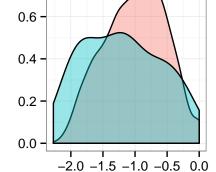
IncRNA group = IDRlessthan0\_2 ::: data cols = bothPullDowns (48 / 2413) = (func/total lncRNA) ::: biotype = lincProcTrans measures across all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 0.12 -0.20 0.09 -0.15 -0.06 -0.10 -0.03 -0.05 0.00 0.00 0 -5 <u>-</u>5 -10 **-10** -10 -20 0 0 5 minExpr nTimesRest maxExpr 0.20 -0.75 withinSubset 0.15 -0.50 false 0.10 true 0.25 0.05 -0.00 0.00 -10.0 -7.5 -5.0 -5 Ó 5 2 -2.5 -10 6 0



tissSpec

0.20

0.15

0.10

0.05

0.00

0.3

0.2

0.1 -

0.0

density

value of facet label