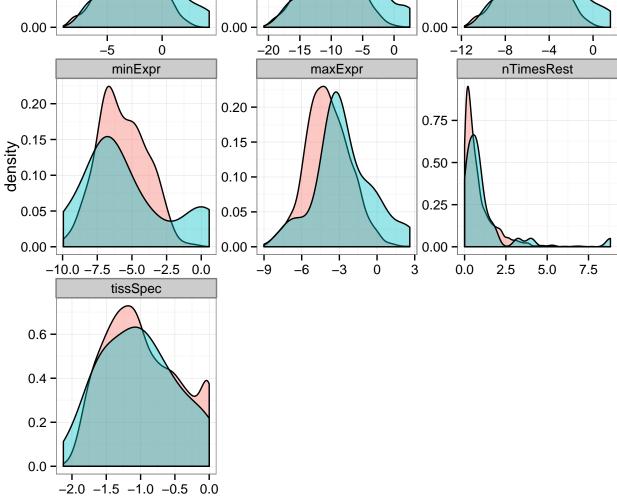
IncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (31/1889) = (func/total IncRNA) ::: biotype = antisense measures across all samples(cell-type/pulldown) varianceExpr sumExpr averageExpr 0.20 0.12 -0.15 0.09 -0.10 0.06 -0.05 0.03 -0.00 0.00 -20 -15 -10 -5 -5 Ó 0 -12 -8 -4 0 minExpr maxExpr nTimesRest



0.20

0.15

0.10

0.05

withinSubset false true

value of facet label