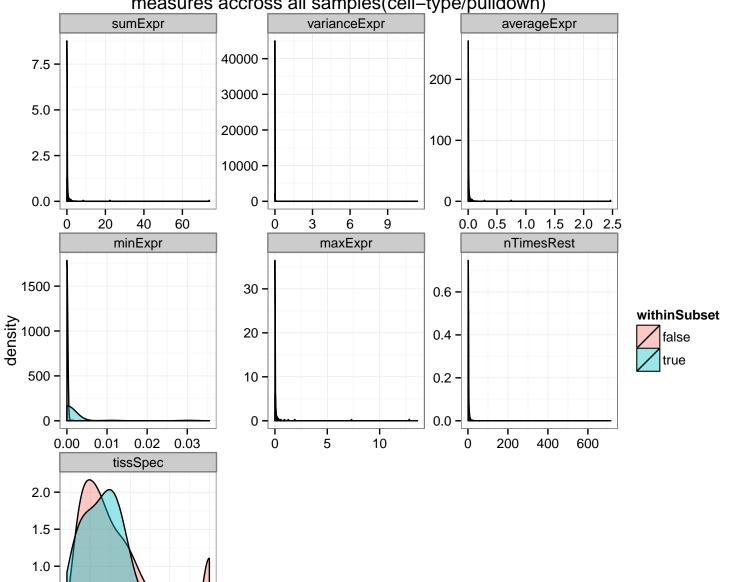
IncRNA group = allLncRNA ::: data cols = bothPullDowns (38 / 2538) = (func/total IncRNA) ::: biotype = antisense measures accross all samples(cell-type/pulldown)



0.5

0.0 -

0.25

0.50

0.75

1.00