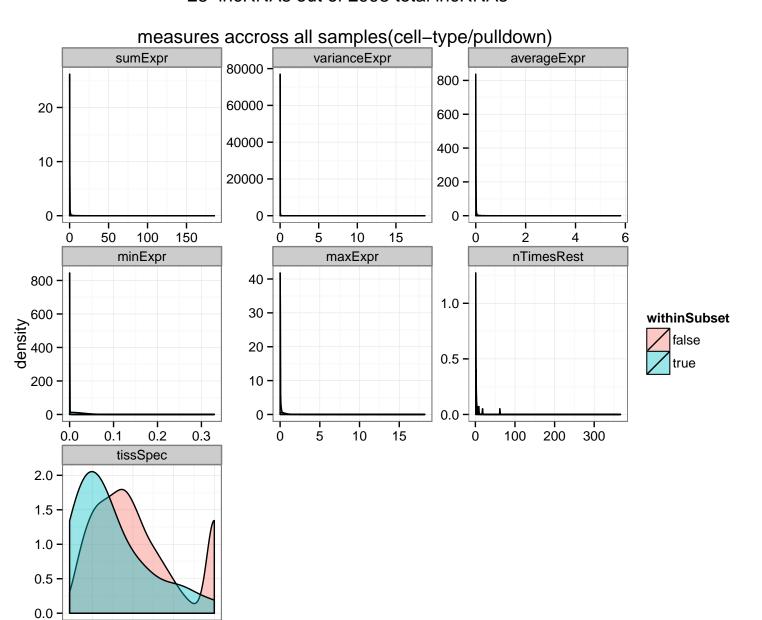
## IncRNA group = allLncRNA ::: data cols = bothPullDowns ( 28 / 2998 ) = (func/total IncRNA) ::: biotype = lincRNA subset = 28 IncRNAs out of 2998 total IncRNAs



value of facet label

0.25

0.50

0.75

1.00