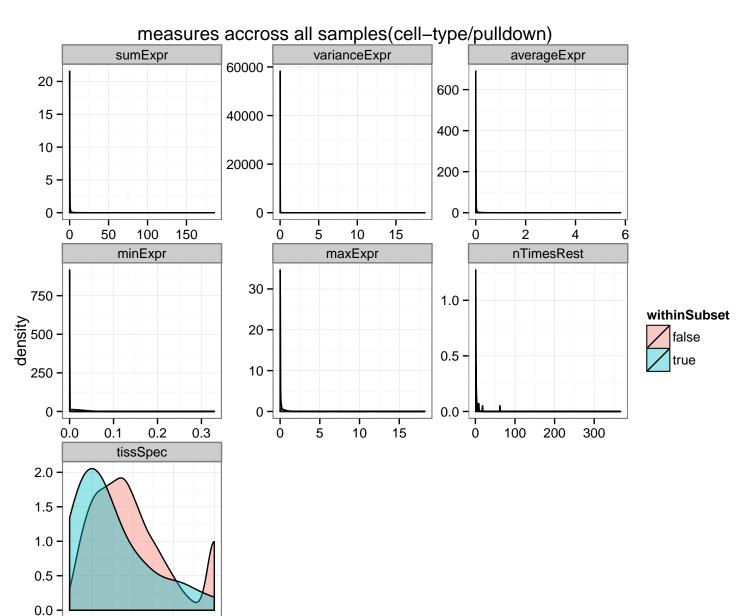
## IncRNA group = allLncRNA ::: data cols = Ipa ( 28 / 2756 ) = (func/total IncRNA) ::: biotype = lincRNA subset = 28 IncRNAs out of 2756 total IncRNAs



value of facet label

0.25

0.50

0.75

1.00