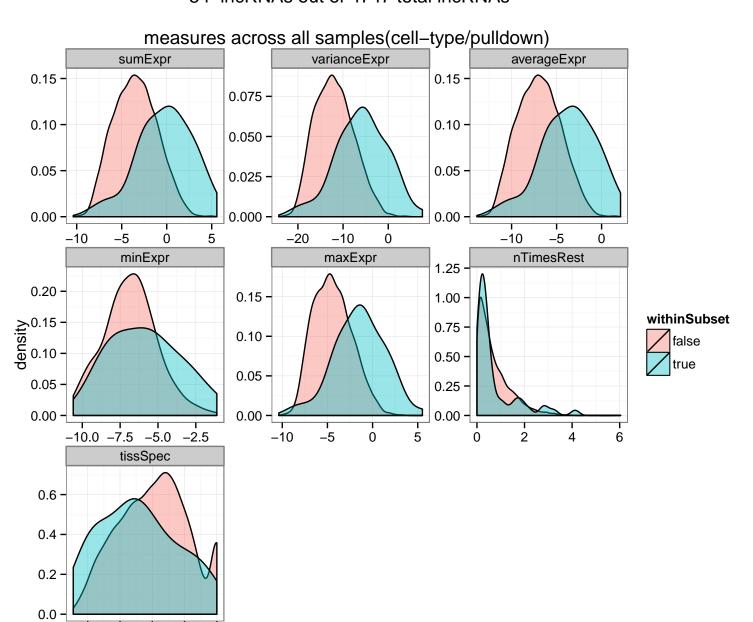
## IncRNA group = allLncRNA ::: data cols = Ipa ( 54 / 4747 ) = (func/total IncRNA) ::: biotype = remove\_antisense subset = 54 IncRNAs out of 4747 total IncRNAs



value of facet label

-2.0 -1.5 -1.0 -0.5 0.0