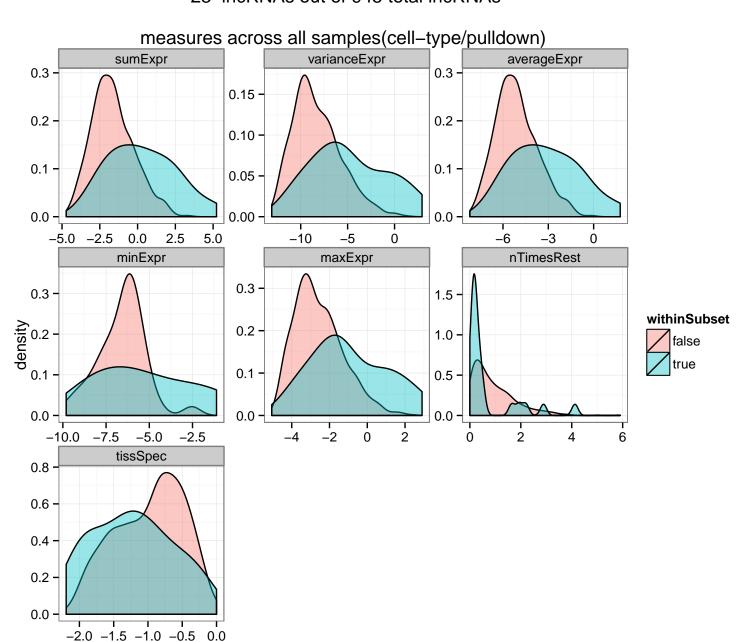
## IncRNA group = IDRlessthan0\_1 ::: data cols = bothPullDowns (23 / 945) = (func/total IncRNA) ::: biotype = lincRNA subset = 23 IncRNAs out of 945 total IncRNAs



value of facet label