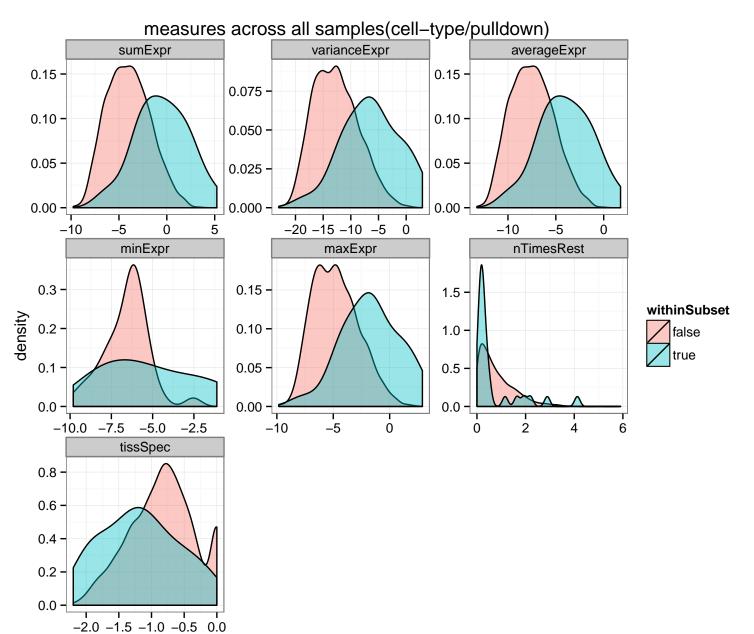
IncRNA group = allLncRNA ::: data cols = Ipa (28 / 2756) = (func/total IncRNA) ::: biotype = lincRNA subset = 28 IncRNAs out of 2756 total IncRNAs



value of facet label