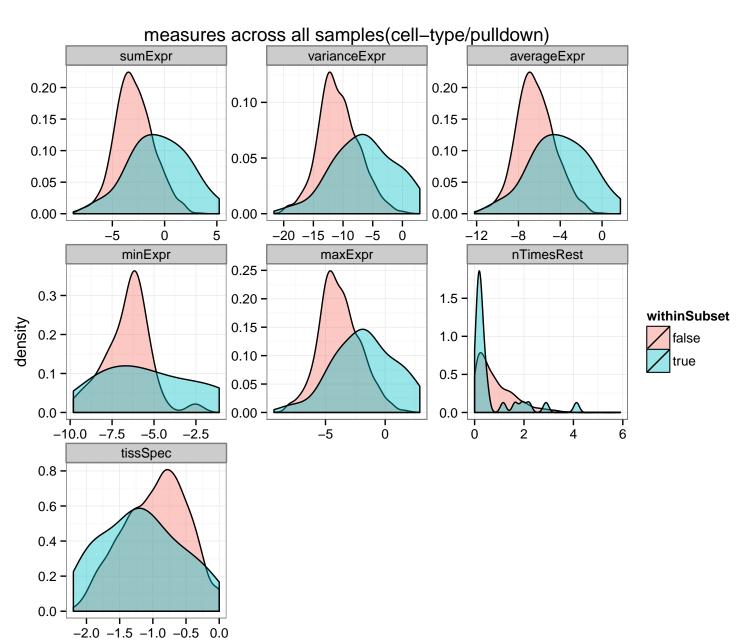
IncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (28 / 1813) = (func/total IncRNA) ::: biotype = lincRNA subset = 28 IncRNAs out of 1813 total IncRNAs



value of facet label