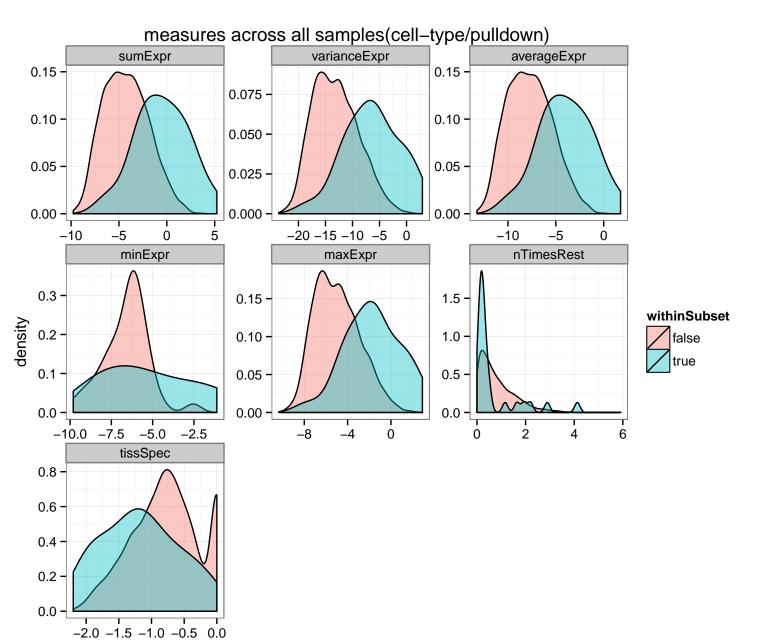
IncRNA group = allLncRNA ::: data cols = bothPullDowns (28 / 2998) = (func/total IncRNA) ::: biotype = lincRNA subset = 28 IncRNAs out of 2998 total IncRNAs



value of facet label