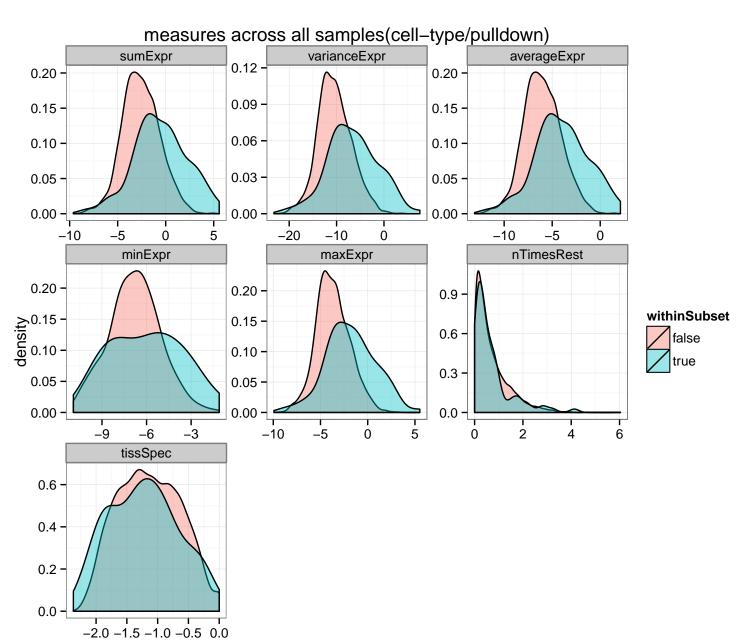
IncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (83 / 5164) = (func/total IncRNA) ::: biotype = all_biotypes subset = 83 IncRNAs out of 5164 total IncRNAs



value of facet label