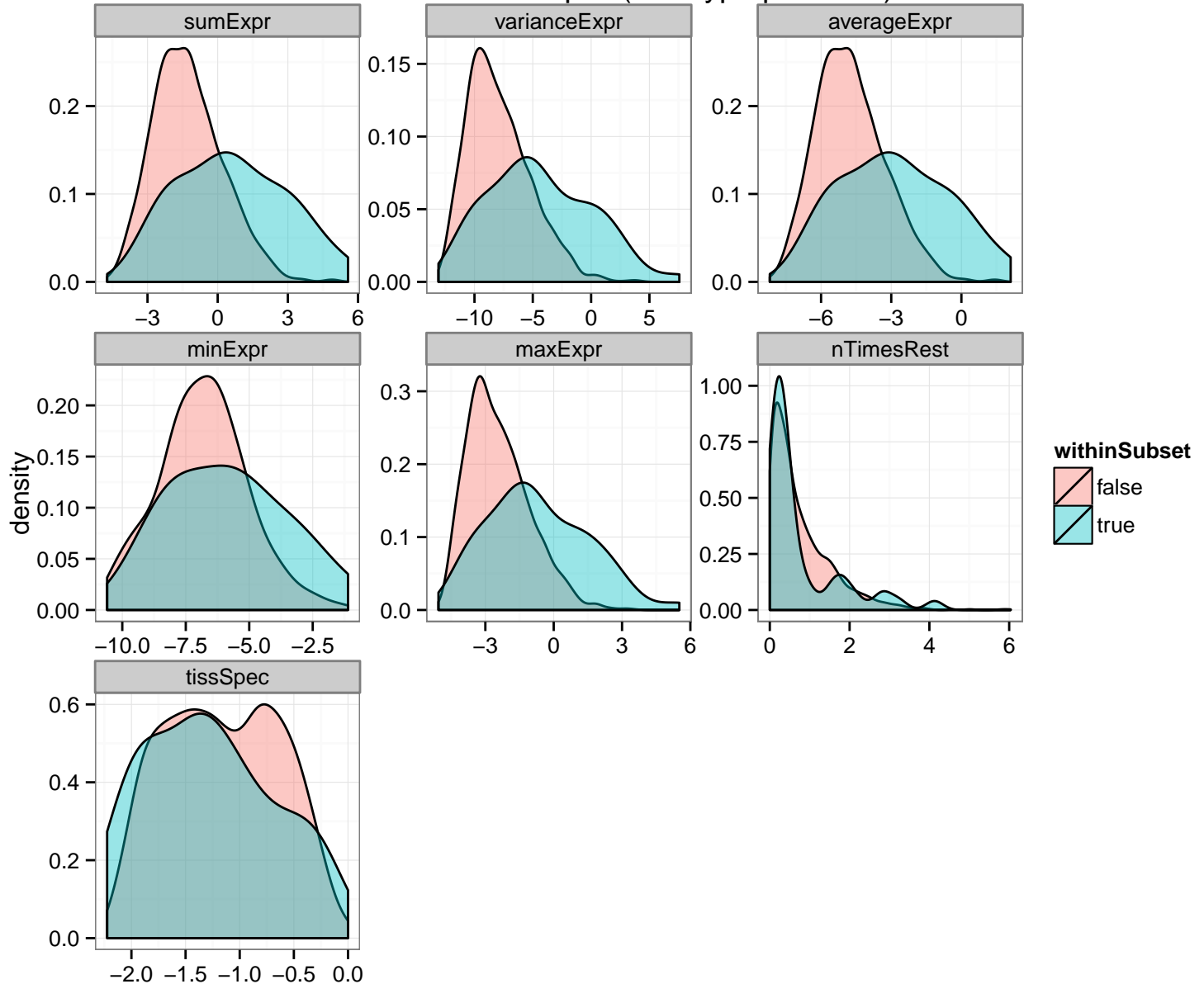


lncRNA group = IDRlessthan0\_1 ::: data cols = bothPullDowns  
( 46 / 1955 ) = (func/total lncRNA) ::: biotype = remove\_antisense  
subset =  
46 lncRNAs out of 1955 total lncRNAs

measures across all samples(cell-type/pulldown)



value of facet label