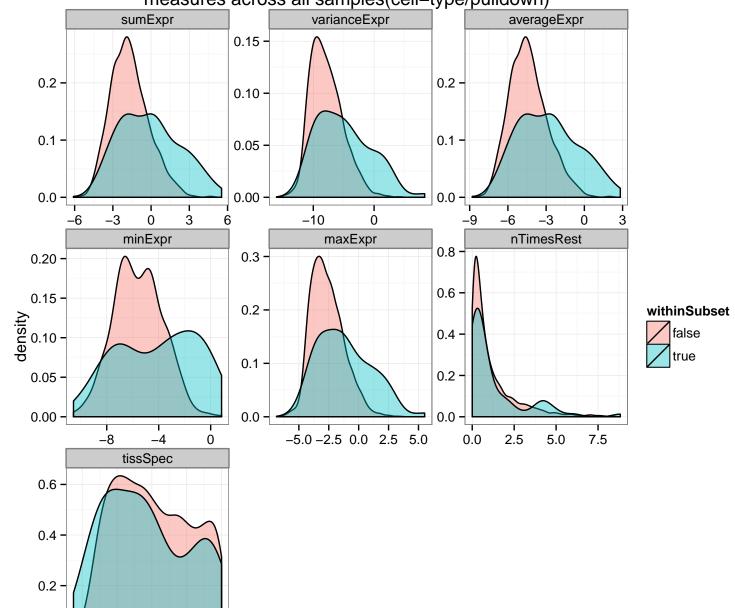
IncRNA group = IDRlessthan0\_1 ::: data cols = Inpa
( 69 / 2559 ) = (func/total IncRNA) ::: biotype = all\_biotypes
 measures across all samples(cell-type/pulldown)



value of facet label

0.0

-2.0 -1.5 -1.0 -0.5