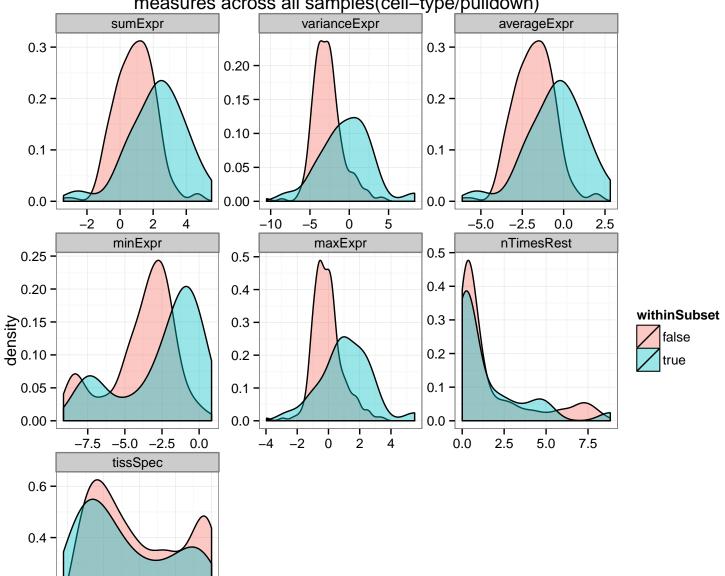
IncRNA group = IDRlessthan0\_01 ::: data cols = lpa ( 25 / 243 ) = (func/total IncRNA) ::: biotype = all\_biotypes measures across all samples(cell-type/pulldown)



value of facet label

0.2

0.0

-2.0 -1.5 -1.0 -0.5