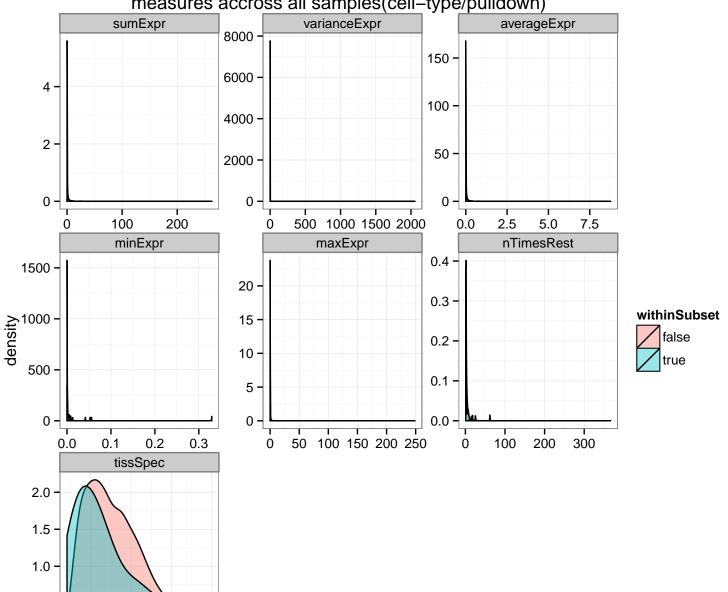
IncRNA group = IDRlessthan0\_2 ::: data cols = bothPullDowns (48 / 2413) = (func/total IncRNA) ::: biotype = lincProcTrans measures accross all samples(cell-type/pulldown)



value of facet label

0.5

0.0 -

0.25

0.50

0.75

1.00