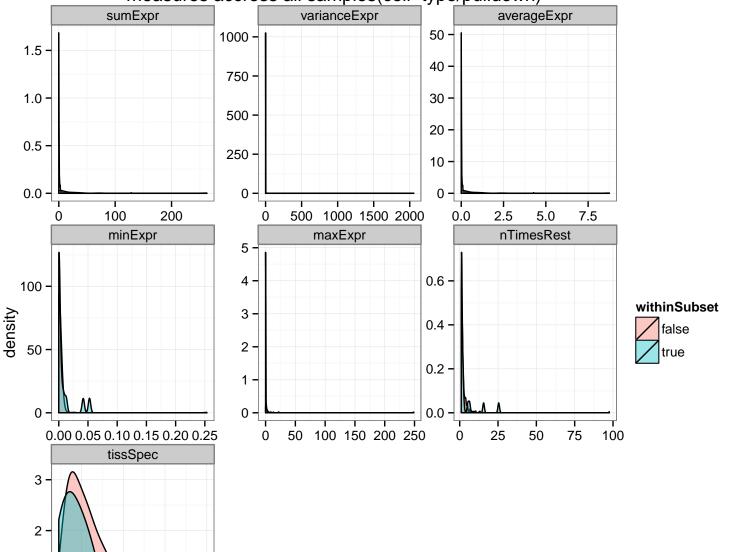
IncRNA group = IDRlessthan0\_2 ::: data cols = lpa ( 17 / 328 ) = (func/total IncRNA) ::: biotype = processed\_transcript measures\_accross all samples(cell-type/pulldown)



1 ·

0

0.25

0.50

0.75

1.00