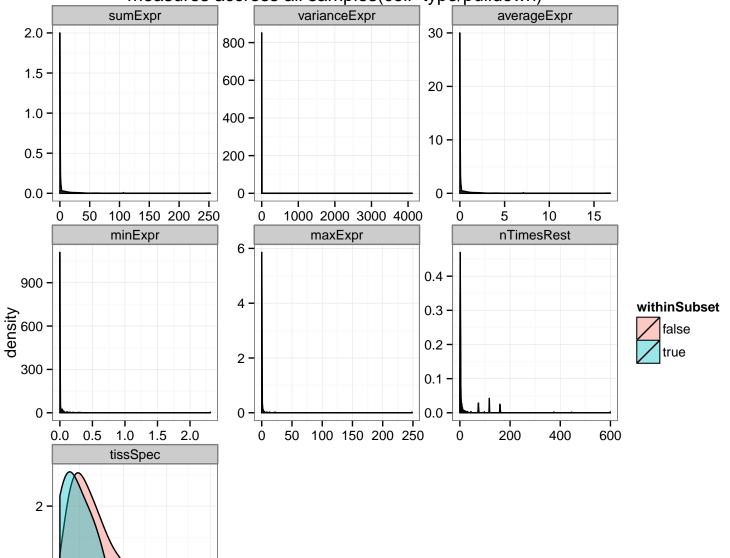
IncRNA group = IDRlessthan0_2 ::: data cols = Inpa (17 / 328) = (func/total IncRNA) ::: biotype = processed_transcript measures accross all samples(cell-type/pulldown)



value of facet label

1 -

0

0.25

0.50

0.75

1.00