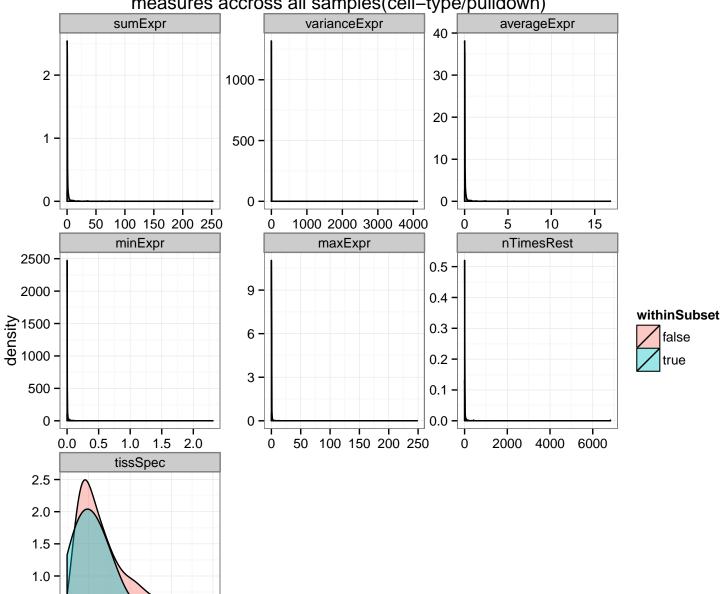
IncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (69 / 2581) = (func/total IncRNA) ::: biotype = all_biotypes measures accross all samples(cell-type/pulldown)



value of facet label

0.5 -

0.0 -

0.50

0.25

0.75

1.00