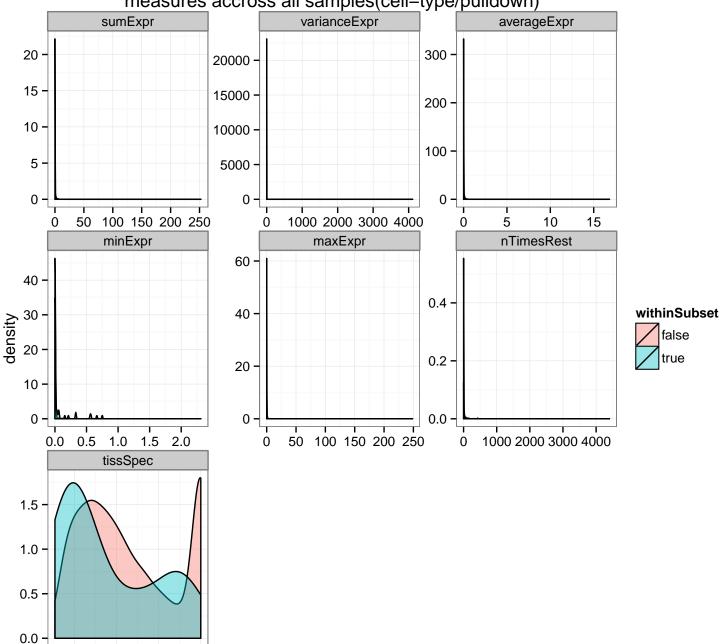
IncRNA group = allLncRNA ::: data cols = lpa (52 / 3884) = (func/total lncRNA) ::: biotype = remove_antisense measures accross all samples(cell-type/pulldown)



value of facet label

0.25

0.50

0.75

1.00