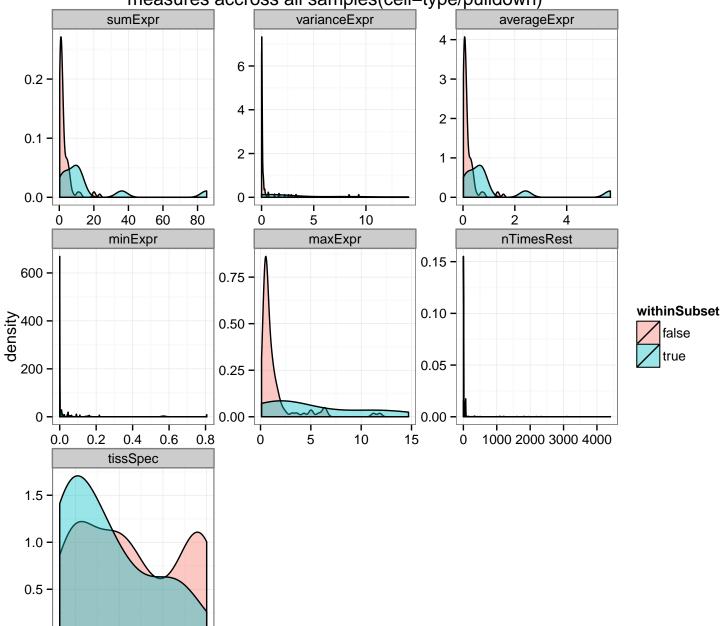
IncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (11 / 109) = (func/total IncRNA) ::: biotype = lincRNA measures accross all samples(cell-type/pulldown)



value of facet label

0.0

0.25

0.50

0.75

1.00