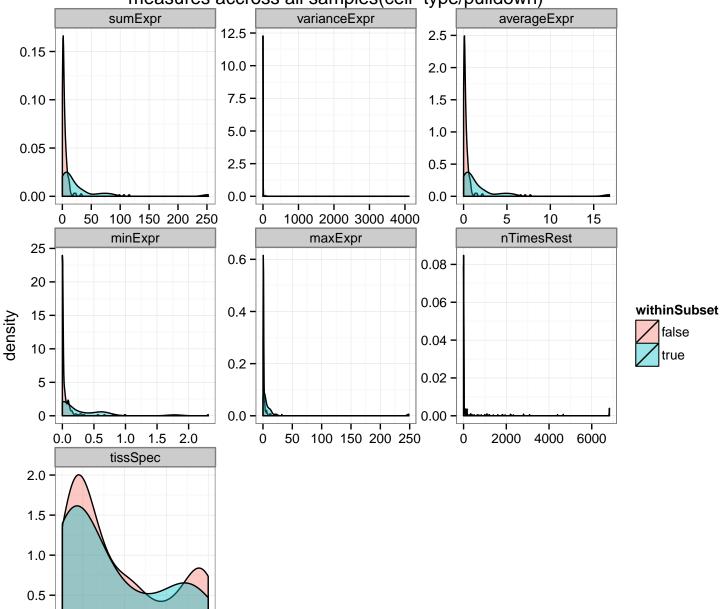
IncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (25 / 243) = (func/total IncRNA) ::: biotype = all_biotypes measures accross all samples(cell-type/pulldown)



value of facet label

0.0

0.50

0.25

0.75

1.00