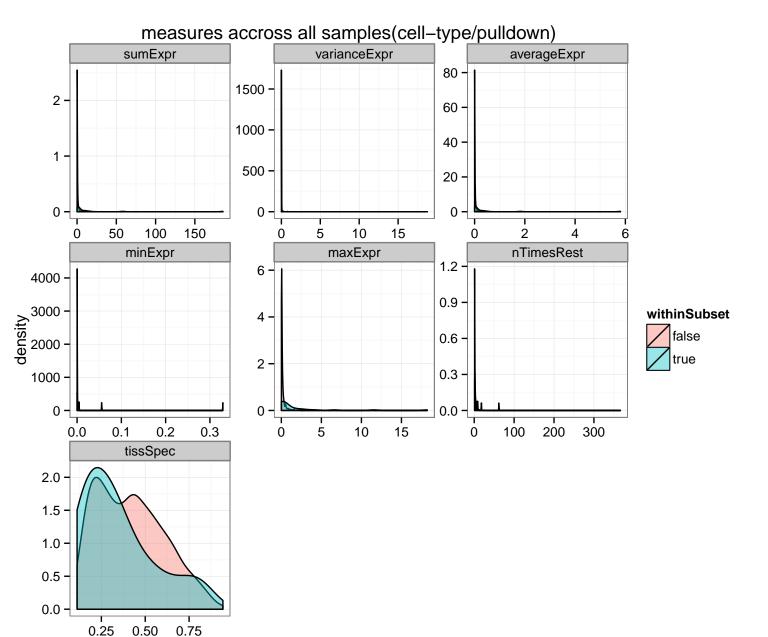# lncRNA group = IDRlessthan0_1 ::: data cols = lnpa
(23 / 935) = (func/total lncRNA) ::: biotype = lincRNA
subset =
23 lncRNAs out of 935 total lncRNAs

measures accross all samples(cell−type/pulldown)

value of facet label