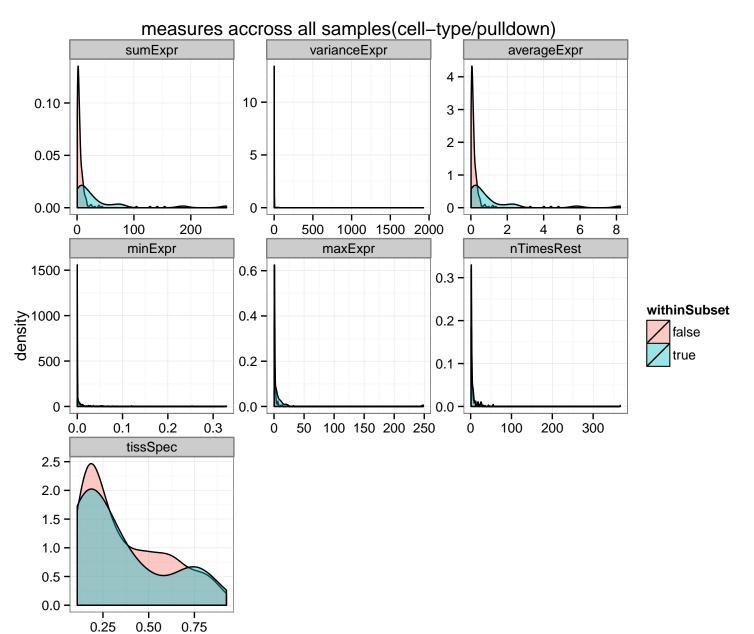# lncRNA group = IDRlessthan0_01 ::: data cols = lpa
## ( 25 / 277 ) = (func/total lncRNA) ::: biotype = all_biotypes
## subset =
## 25  lncRNAs out of 277 total lncRNAs



measures accross all samples(cell−type/pulldown)