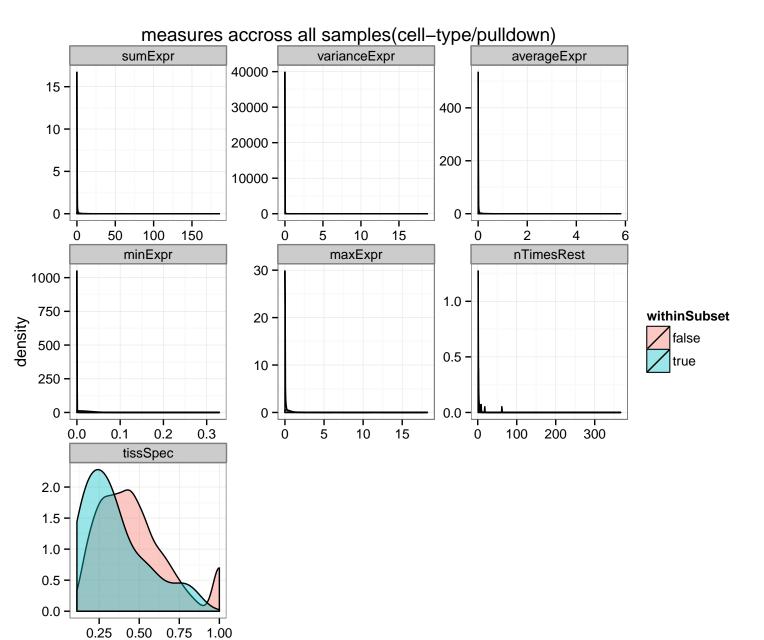
IncRNA group = allLncRNA ::: data cols = Inpa (27 / 2505) = (func/total IncRNA) ::: biotype = lincRNA subset = 27 IncRNAs out of 2505 total IncRNAs



value of facet label