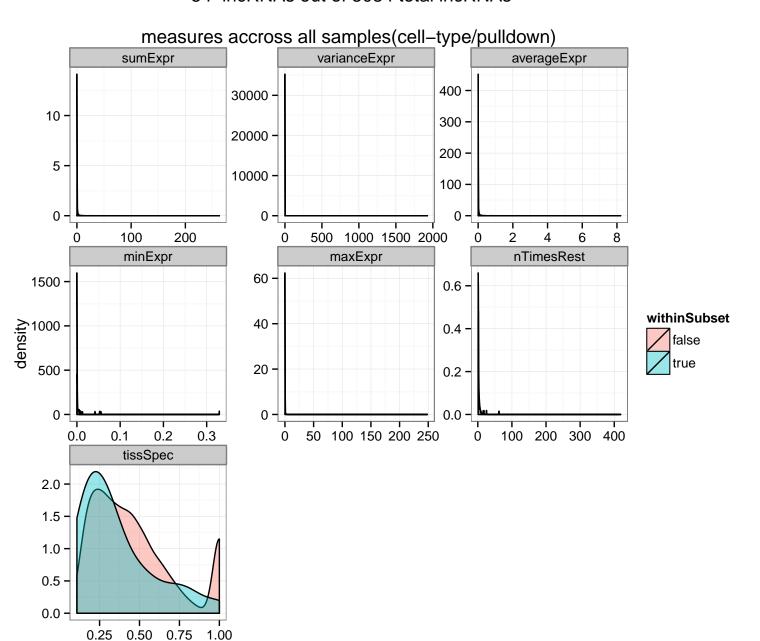
IncRNA group = allLncRNA ::: data cols = bothPullDowns (54 / 5084) = (func/total IncRNA) ::: biotype = remove_antisense subset = 54 IncRNAs out of 5084 total IncRNAs



value of facet label