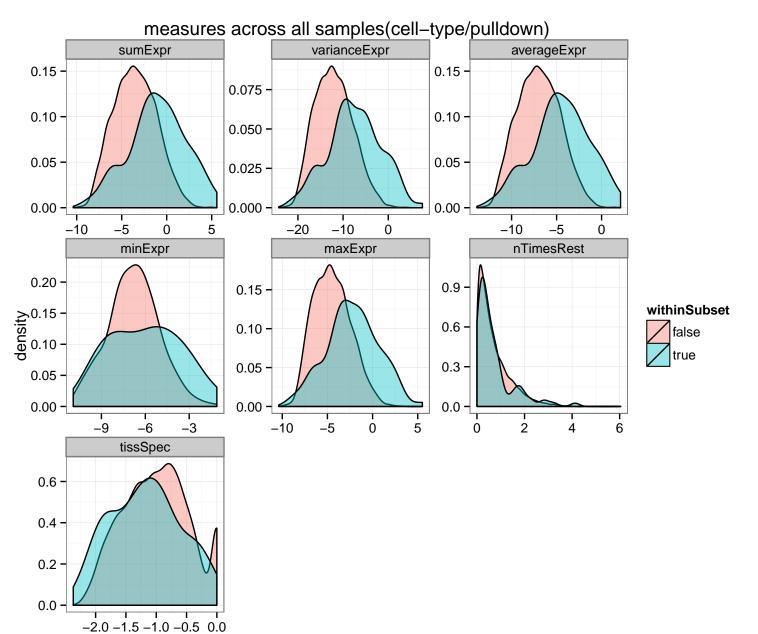
IncRNA group = allLncRNA ::: data cols = lpa (91 / 7106) = (func/total IncRNA) ::: biotype = all_biotypes subset = 91 IncRNAs out of 7106 total IncRNAs



value of facet label