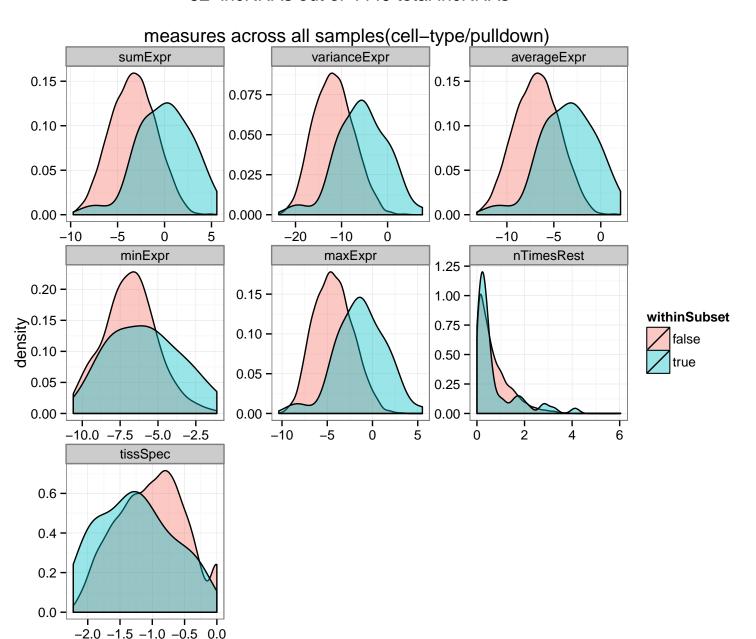
IncRNA group = allLncRNA ::: data cols = Inpa (52 / 4449) = (func/total IncRNA) ::: biotype = remove_antisense subset = 52 IncRNAs out of 4449 total IncRNAs



value of facet label