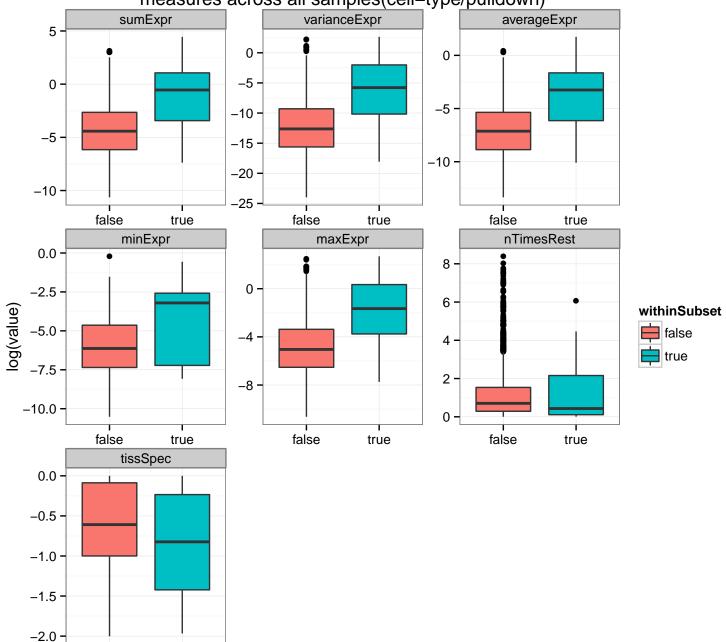
IncRNA group = allLncRNA ::: data cols = lpa (32 / 3068) = (func/total lncRNA) ::: biotype = lincRNA measures across all samples(cell-type/pulldown)



value of facet label

false

true