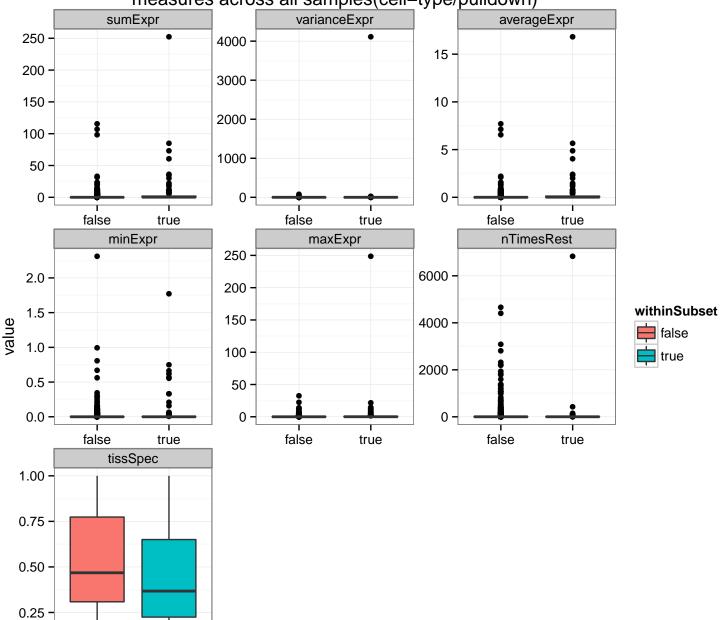
IncRNA group = allLncRNA ::: data cols = lpa (90 / 6451) = (func/total lncRNA) ::: biotype = all_biotypes measures across all samples(cell-type/pulldown)



value of facet label

false

true