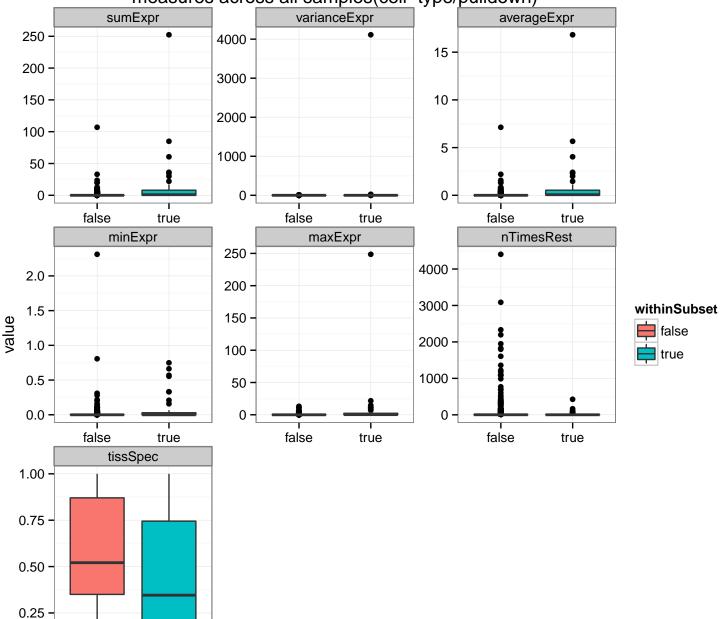
IncRNA group = allLncRNA ::: data cols = lpa (49 / 3453) = (func/total lncRNA) ::: biotype = lincProcTrans measures across all samples(cell-type/pulldown)



value of facet label

false

true