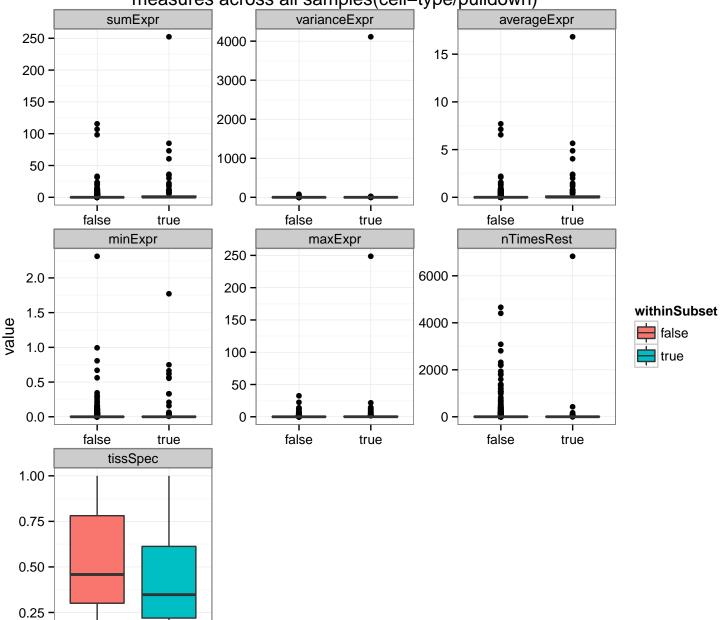
IncRNA group = allLncRNA ::: data cols = Inpa (87 / 6034) = (func/total IncRNA) ::: biotype = all_biotypes measures across all samples(cell-type/pulldown)



value of facet label

false

true