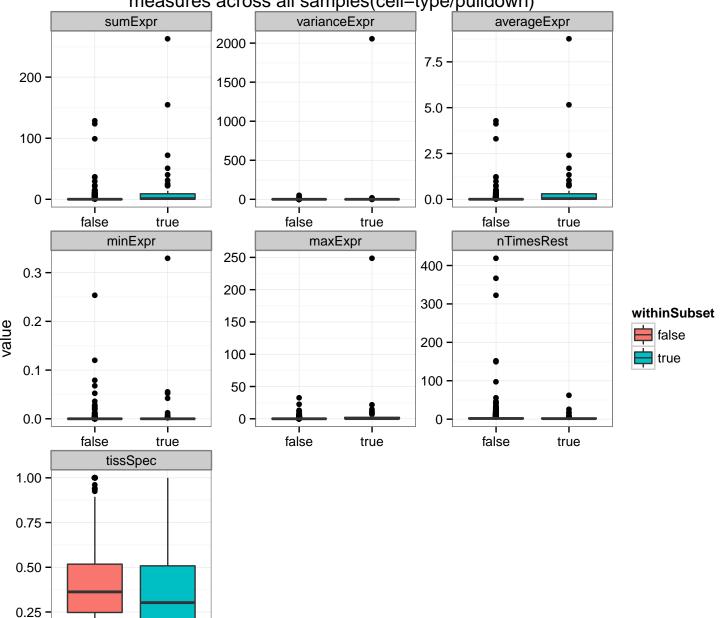
IncRNA group = IDRIessthan0_2 ::: data cols = bothPullDowns (51 / 2754) = (func/total IncRNA) ::: biotype = remove_antisense measures across all samples(cell-type/pulldown)



value of facet label

false

true