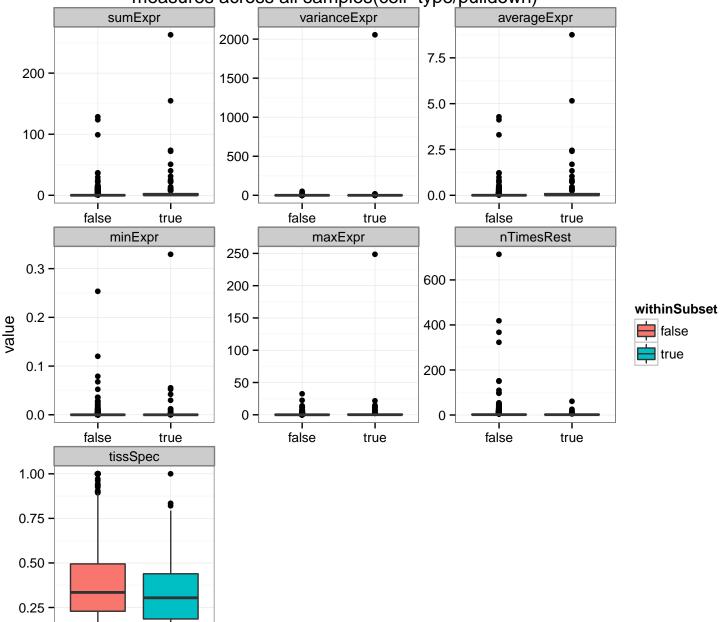
IncRNA group = IDRlessthan0\_2 ::: data cols = bothPullDowns (82 / 4643) = (func/total IncRNA) ::: biotype = all\_biotypes measures across all samples(cell-type/pulldown)



value of facet label

false

true