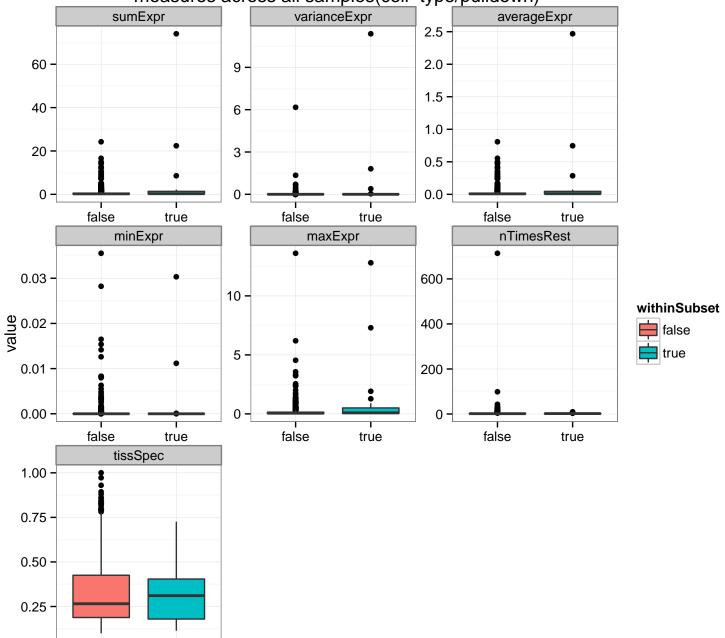
IncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (23 / 902) = (func/total IncRNA) ::: biotype = antisense measures across all samples(cell-type/pulldown)



value of facet label

false

true