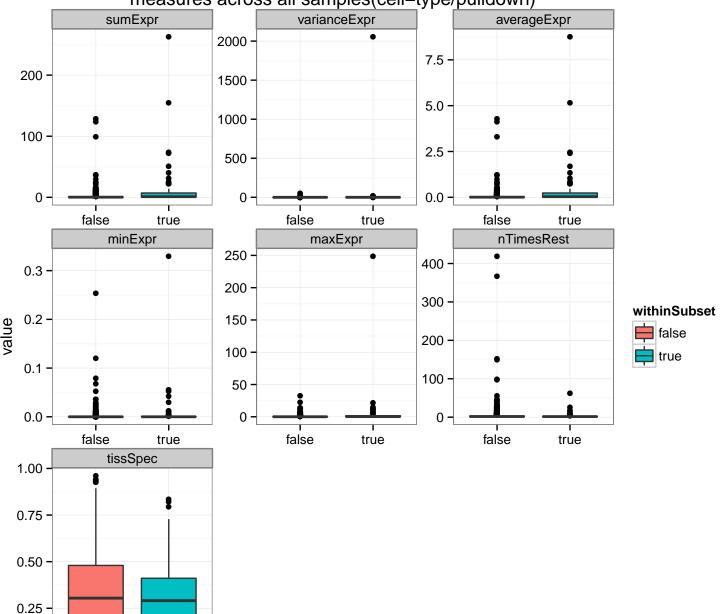
IncRNA group = IDRlessthan0_1 ::: data cols = Inpa
(69 / 2559) = (func/total IncRNA) ::: biotype = all_biotypes
 measures across all samples(cell-type/pulldown)



value of facet label

false

true