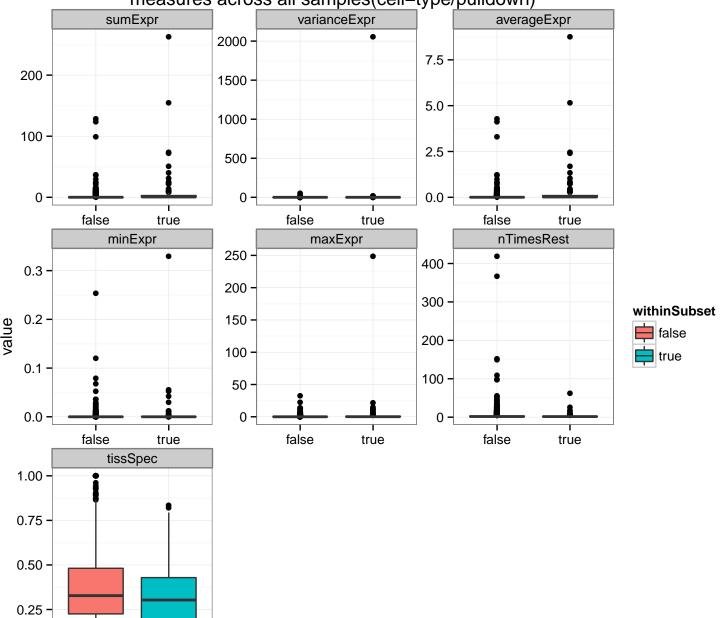
IncRNA group = IDRlessthan0\_2 ::: data cols = Inpa
( 81 / 4501 ) = (func/total IncRNA) ::: biotype = all\_biotypes
 measures across all samples(cell-type/pulldown)



value of facet label

false

true