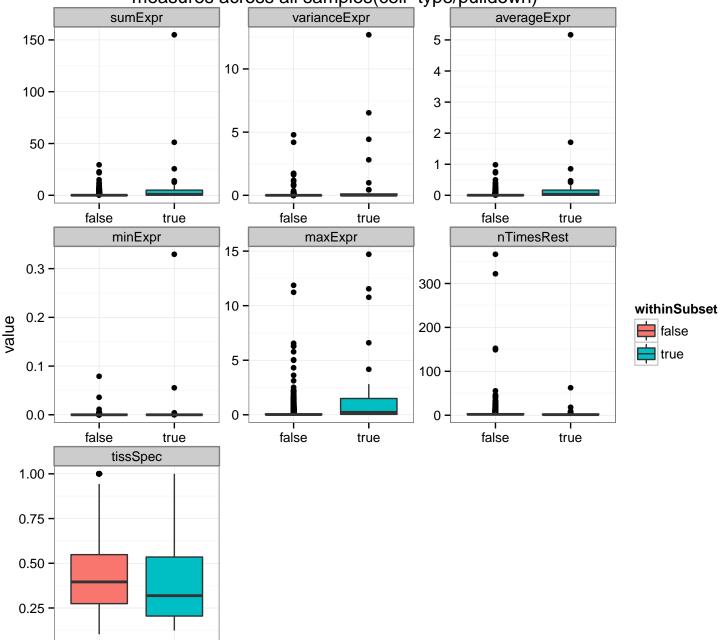
IncRNA group = IDRlessthan0\_2 ::: data cols = bothPullDowns (31 / 2082) = (func/total IncRNA) ::: biotype = lincRNA measures across all samples(cell-type/pulldown)



value of facet label

false

true