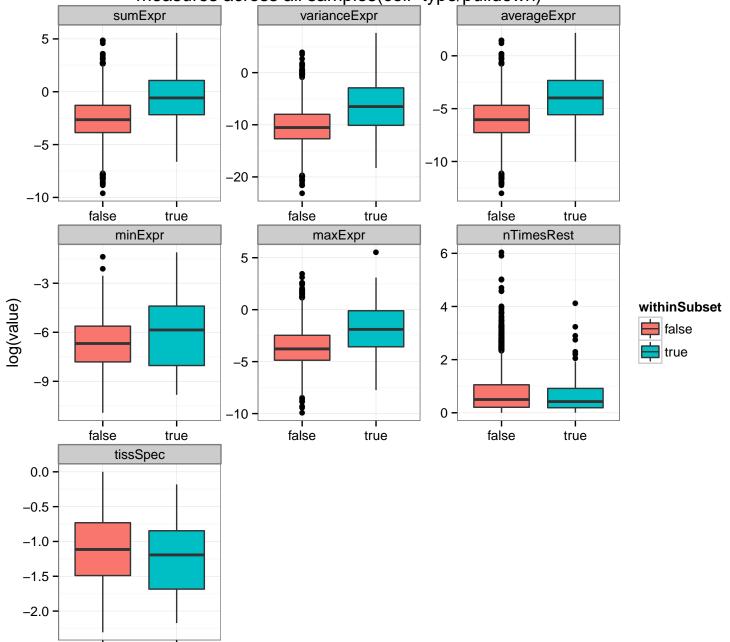
IncRNA group = IDRlessthan0_2 ::: data cols = Inpa
(81 / 4501) = (func/total IncRNA) ::: biotype = all_biotypes
 measures across all samples(cell-type/pulldown)



value of facet label

false

true