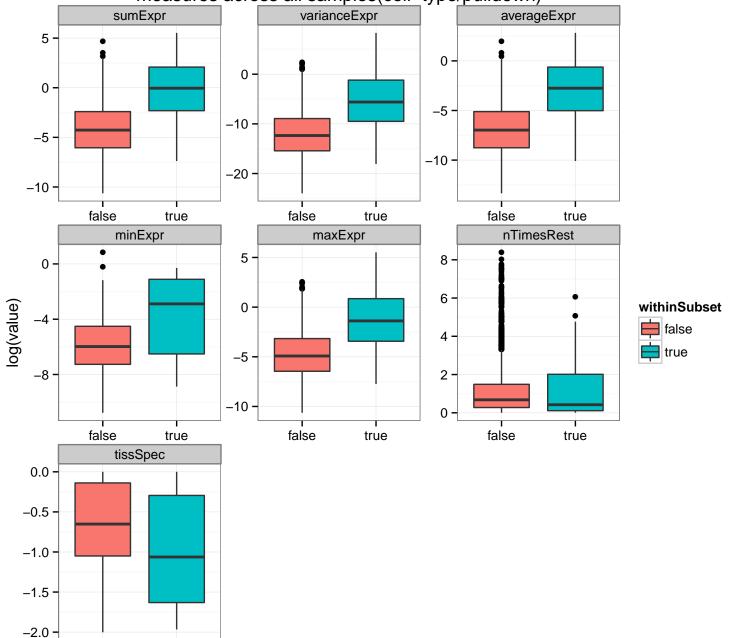
IncRNA group = allLncRNA ::: data cols = Ipa ( 49 / 3453 ) = (func/total IncRNA) ::: biotype = lincProcTrans measures across all samples(cell-type/pulldown)



value of facet label

false

true