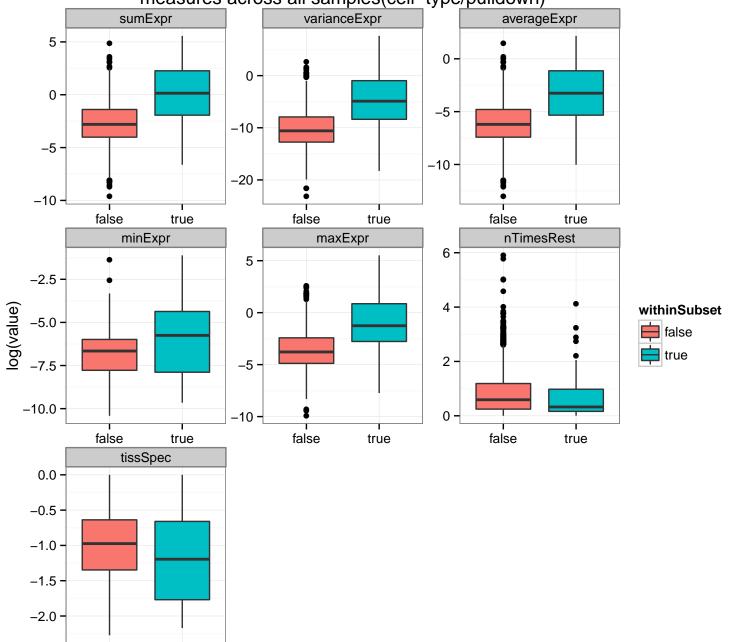
IncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (48 / 2413) = (func/total IncRNA) ::: biotype = lincProcTrans measures across all samples(cell-type/pulldown)



value of facet label

false

true