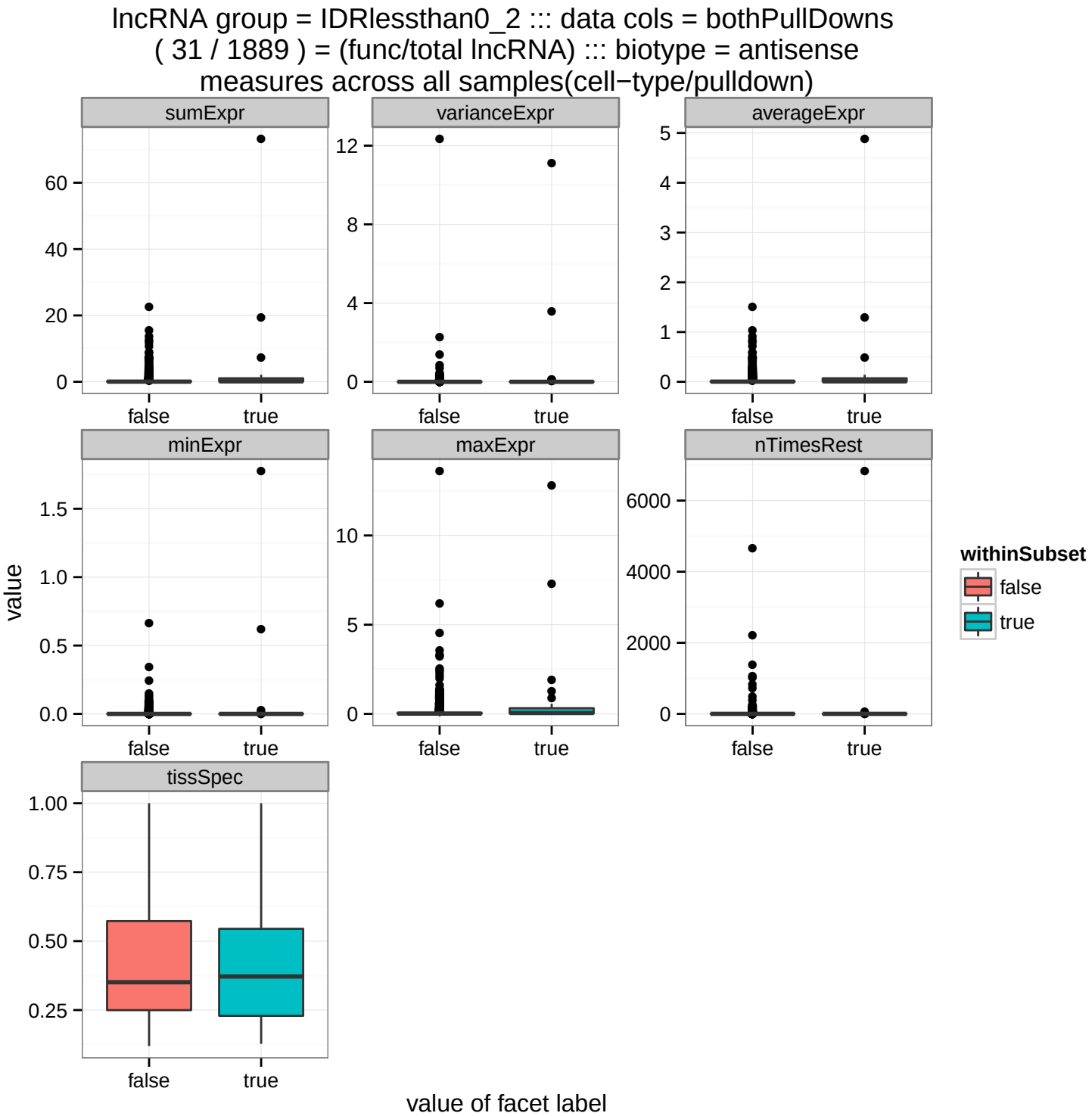
lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns
(31 / 1889) = (func/total lncRNA) ::: biotype = antisense
measures across all samples(cell−type/pulldown)
sumExpr
varianceExpr
averageExpr
minExpr
maxExpr
nTimesRest
tissSpec
value
false
true
withinSubset
value of facet label