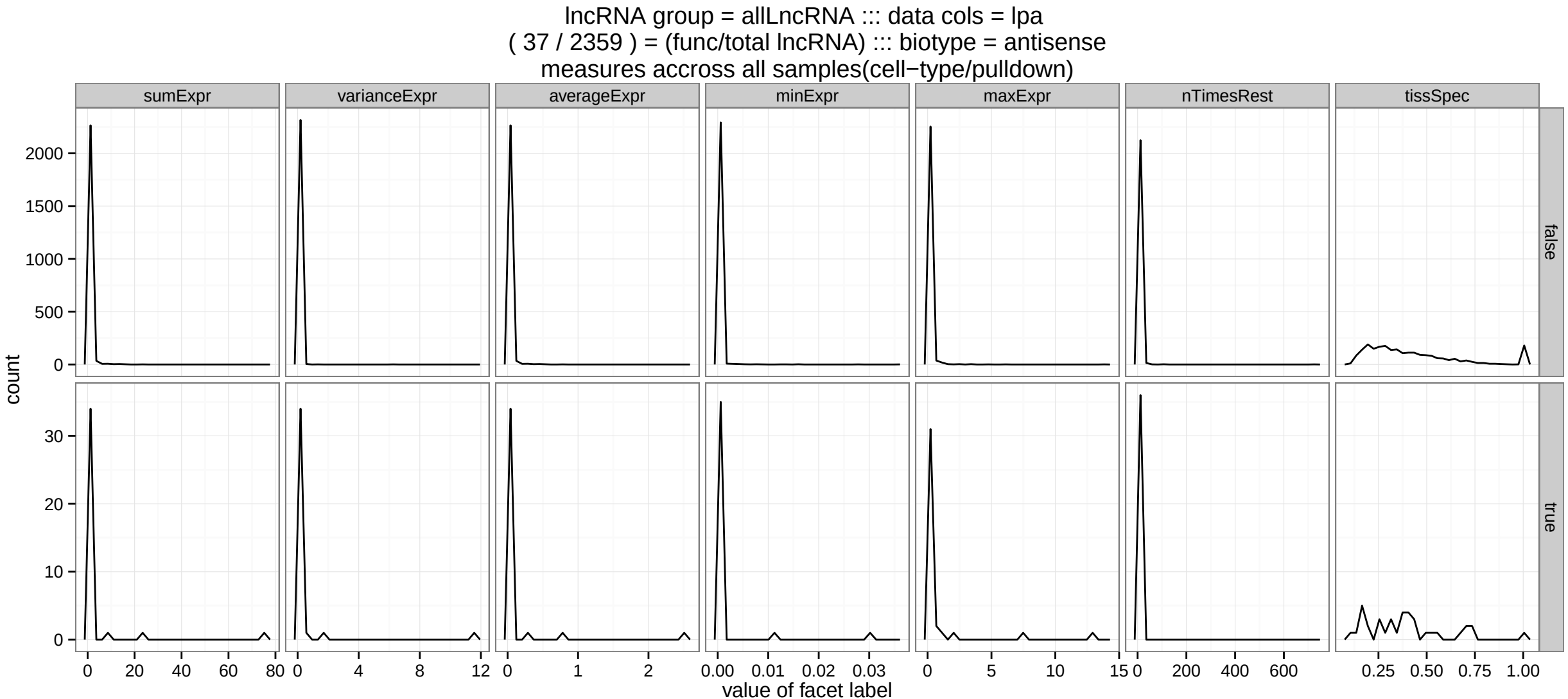
lncRNA group = allLncRNA ::: data cols = lpa
(37 / 2359) = (func/total lncRNA) ::: biotype = antisense
measures accross all samples(cell−type/pulldown)
sumExpr
varianceExpr
averageExpr
minExpr
maxExpr
nTimesRest
tissSpec
false
true
count
value of facet label