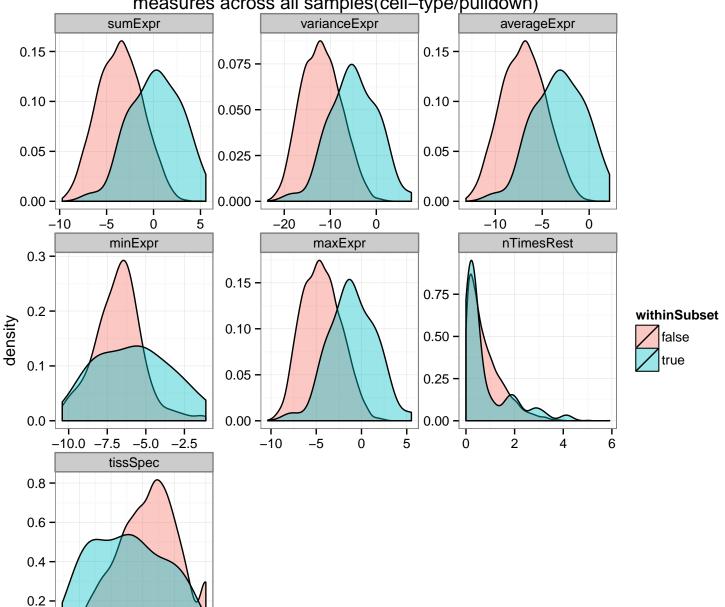
IncRNA group = allLncRNA ::: data cols = Inpa
(47 / 3175) = (func/total IncRNA) ::: biotype = lincProcTrans
measures across all samples(cell-type/pulldown)

sumExpr varianceExpr averageExpr



value of facet label

0.0

-2.0 -1.5 -1.0 -0.5 0.0