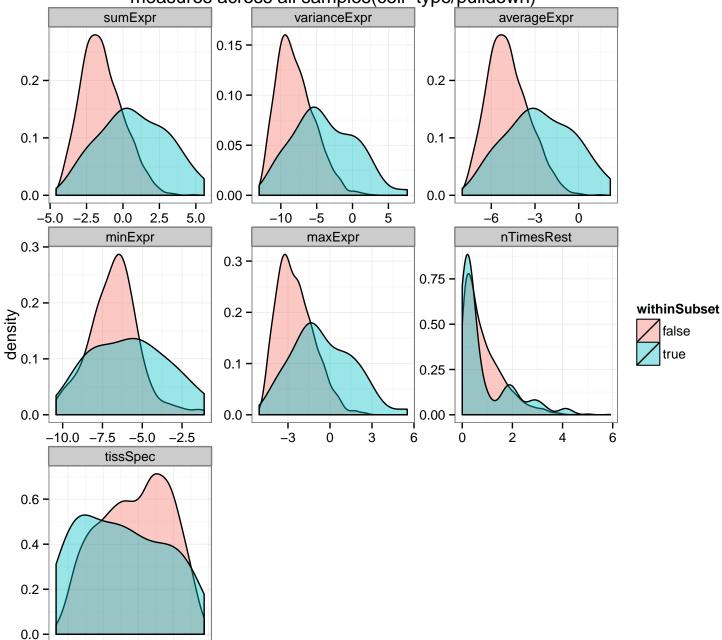
IncRNA group = IDRIessthan0_1 ::: data cols = Inpa
(42 / 1350) = (func/total IncRNA) ::: biotype = lincProcTrans
measures across all samples(cell–type/pulldown)

sumExpr varianceExpr averageExpr



value of facet label

-2.0 -1.5 -1.0 -0.5