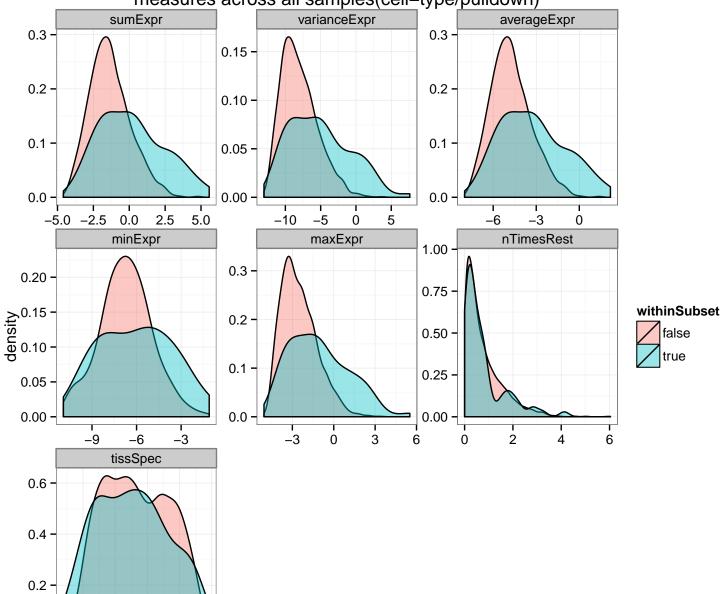
IncRNA group = IDRlessthan0_1 ::: data cols = Inpa
(69 / 2559) = (func/total IncRNA) ::: biotype = all_biotypes
measures across all samples(cell-type/pulldown)

sumExpr
varianceExpr
averageExpr
0.3



value of facet label

0.0

-2.0 -1.5 -1.0 -0.5 0.0