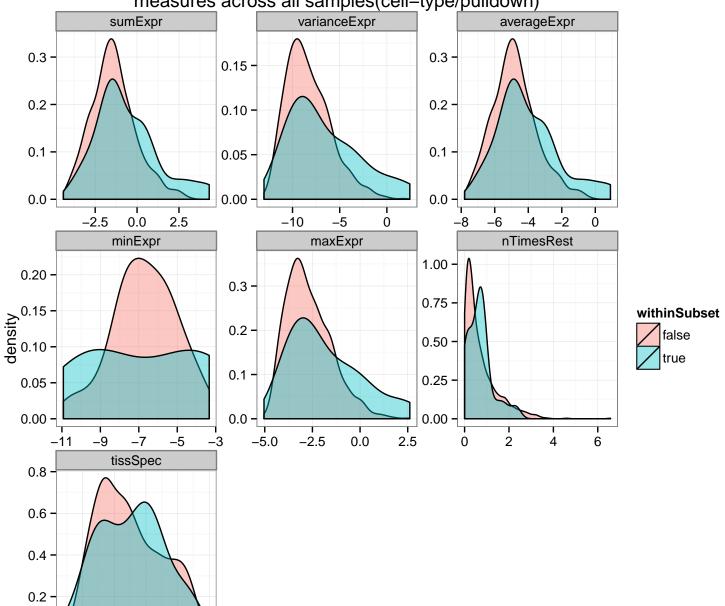
IncRNA group = IDRlessthan0\_1 ::: data cols = bothPullDowns
( 23 / 902 ) = (func/total IncRNA) ::: biotype = antisense
measures across all samples(cell–type/pulldown)

sumExpr varianceExpr averageExpr



value of facet label

0.0

-2.0 -1.5 -1.0 -0.5 0.0