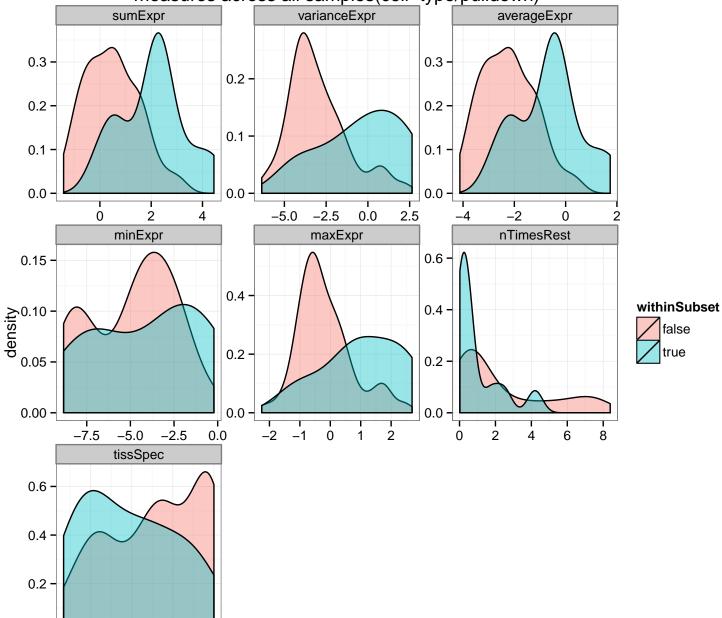
IncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (11 / 109) = (func/total IncRNA) ::: biotype = lincRNA measures across all samples(cell-type/pulldown)



value of facet label

0.0 -

-1.0

-0.5

0.0

-1.5