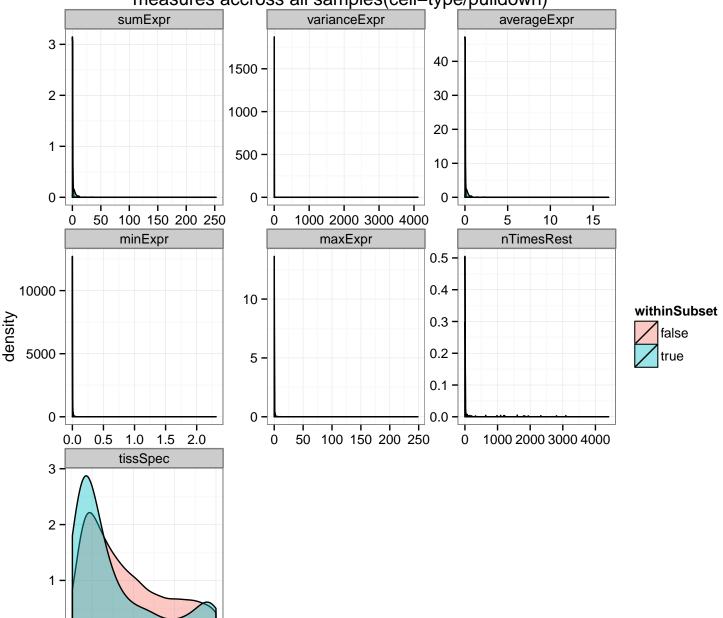
IncRNA group = lincProcTrans ::: data cols = lpa (170 / 1954) = (func/total lncRNA) ::: label = pred by ridgeReg measures accross all samples(cell-type/pulldown)



value of facet label

0

0.25

0.50

0.75

1.00