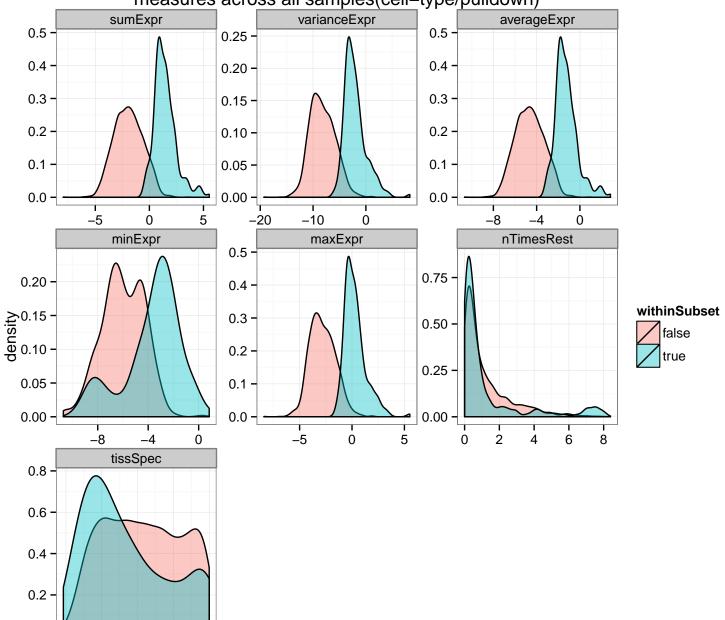
IncRNA group = lincProcTrans ::: data cols = lpa (170 / 1954) = (func/total IncRNA) ::: label = pred by ridgeReg measures across all samples(cell-type/pulldown)



value of facet label

0.0

-2.0 -1.5 -1.0 -0.5

0.0