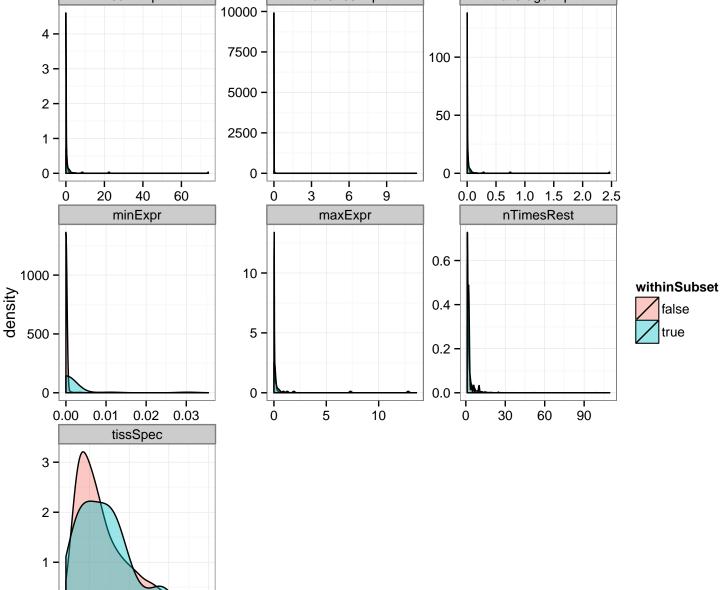
IncRNA group = IDRIessthan0_2 ::: data cols = Inpa (31 / 1841) = (func/total IncRNA) ::: biotype = antisense measures accross all samples(cell-type/pulldown) sumExpr varianceExpr averageExpr 10000 -7500 -100 5000 50 2500 -0 0 0.5 1.5 2.0 20 40 60 3 6 9 0.0 1.0 0



value of facet label

0

0.25

0.50

0.75

1.00