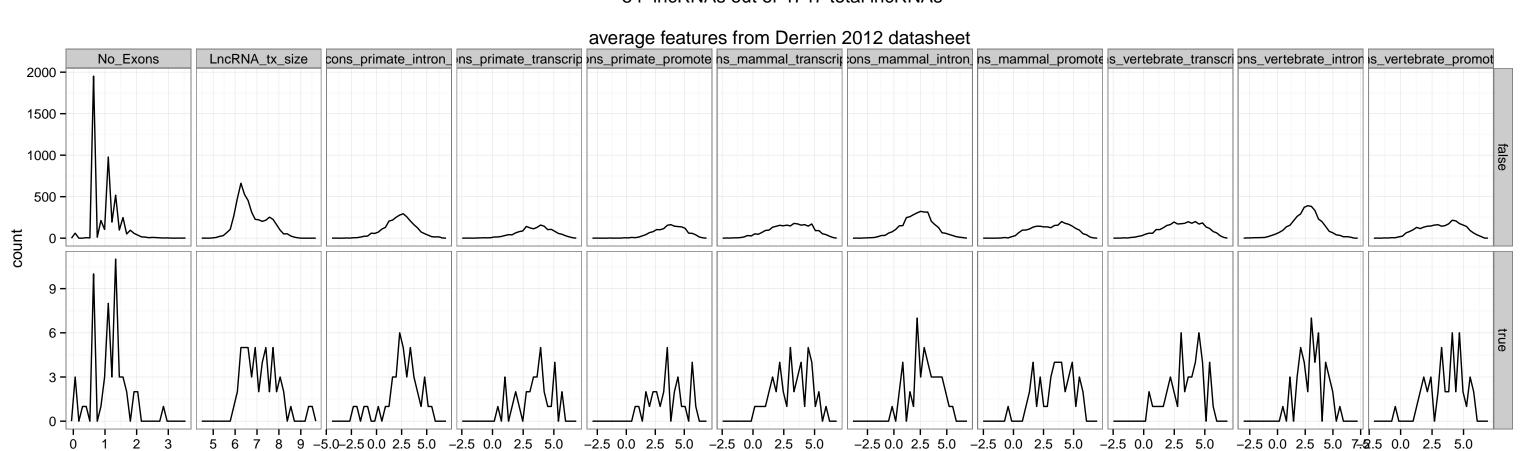
IncRNA group = allLncRNA ::: data cols = lpa (54 / 4747) = (func/total IncRNA) ::: biotype = remove\_antisense subset = 54 IncRNAs out of 4747 total IncRNAs



value of facet label