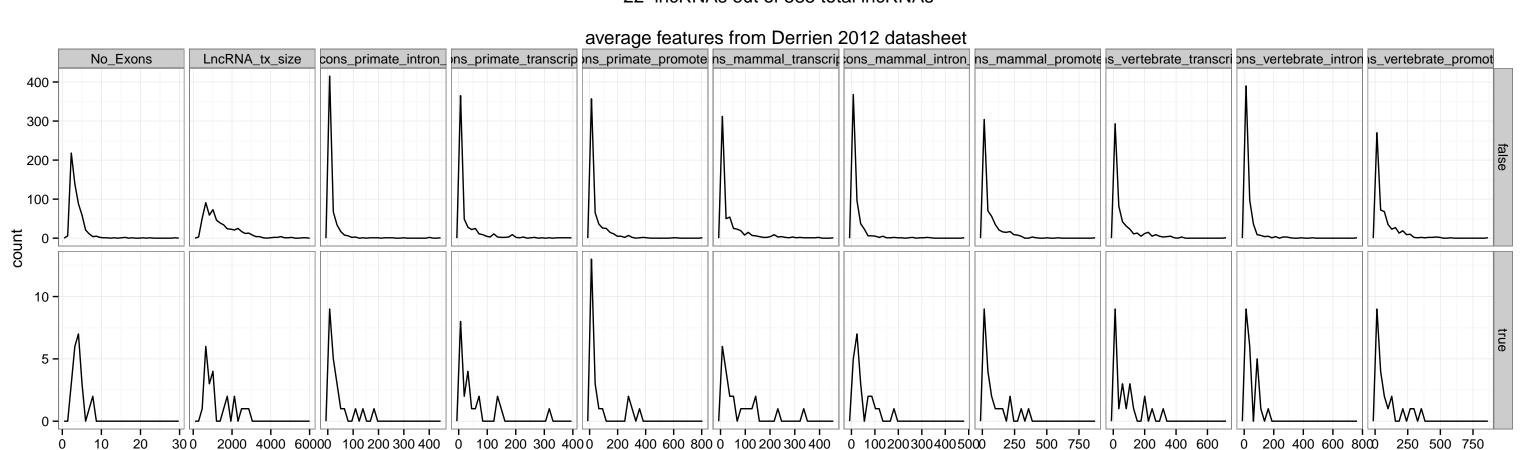
lncRNA group = IDRlessthan0\_1 ::: data cols = Inpa ( 22 / 585 ) = (func/total IncRNA) ::: biotype = processed\_transcript subset = 22 IncRNAs out of 585 total IncRNAs



value of facet label

100 200 300 4000