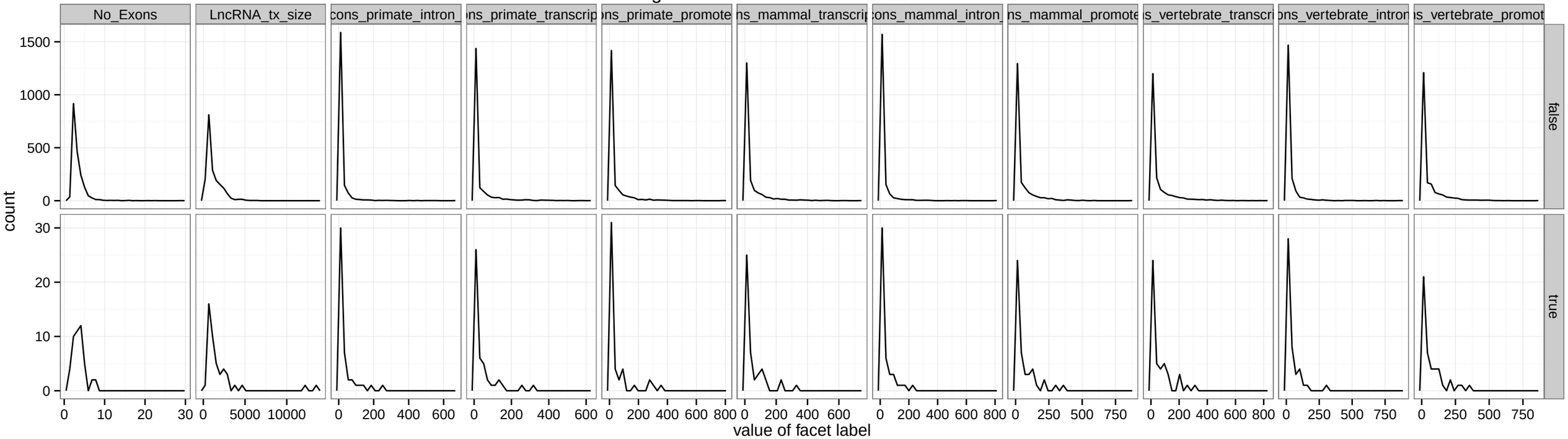
lncRNA group = IDRlessthan0_1 ::: data cols = lnpa
(46 / 1941) = (func/total lncRNA) ::: biotype = remove_antisense
subset =
46 lncRNAs out of 1941 total lncRNAs

average features from Derrien 2012 datasheet