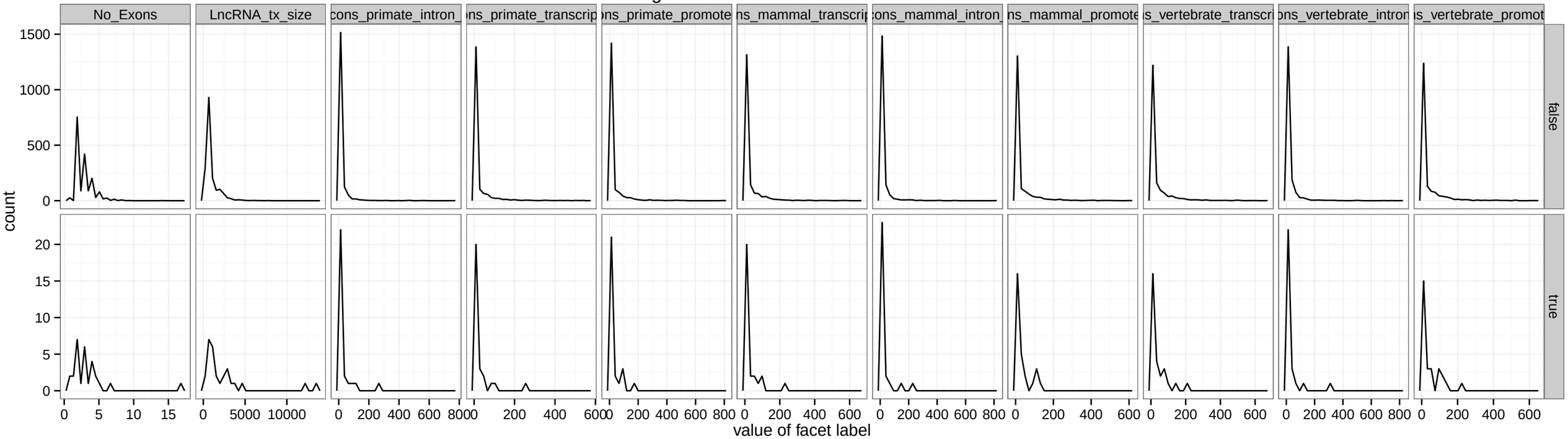
lncRNA group = IDRlessthan0_2 ::: data cols = lpa
(28 / 1784) = (func/total lncRNA) ::: biotype = lincRNA
subset =
28 lncRNAs out of 1784 total lncRNAs

average features from Derrien 2012 datasheet