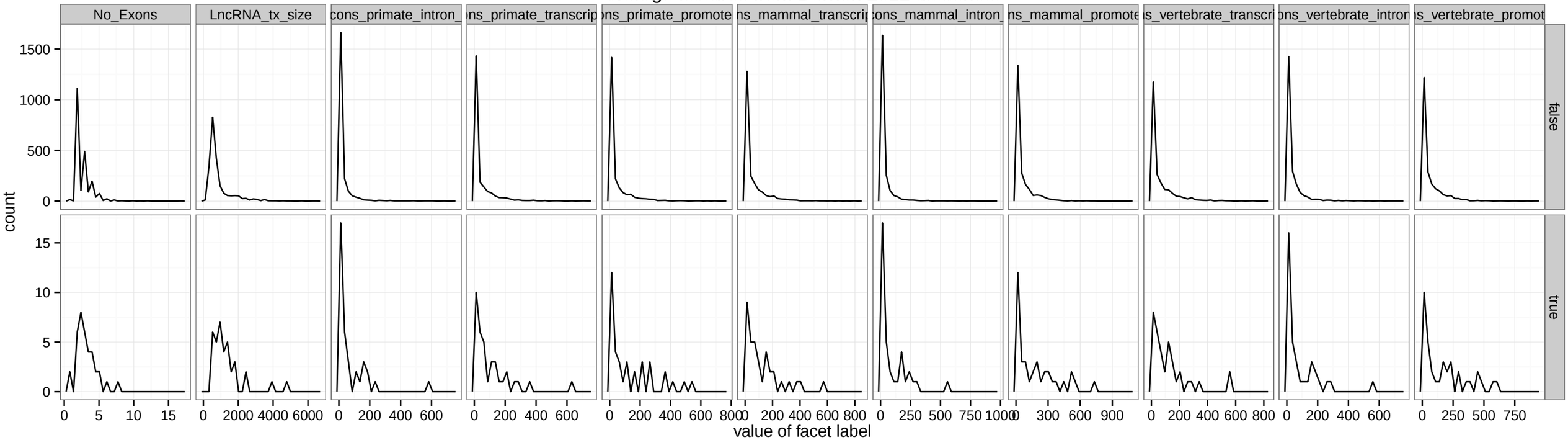
lncRNA group = allLncRNA ::: data cols = lnpa
(36 / 2219) = (func/total lncRNA) ::: biotype = antisense
subset =
36 lncRNAs out of 2219 total lncRNAs

average features from Derrien 2012 datasheet