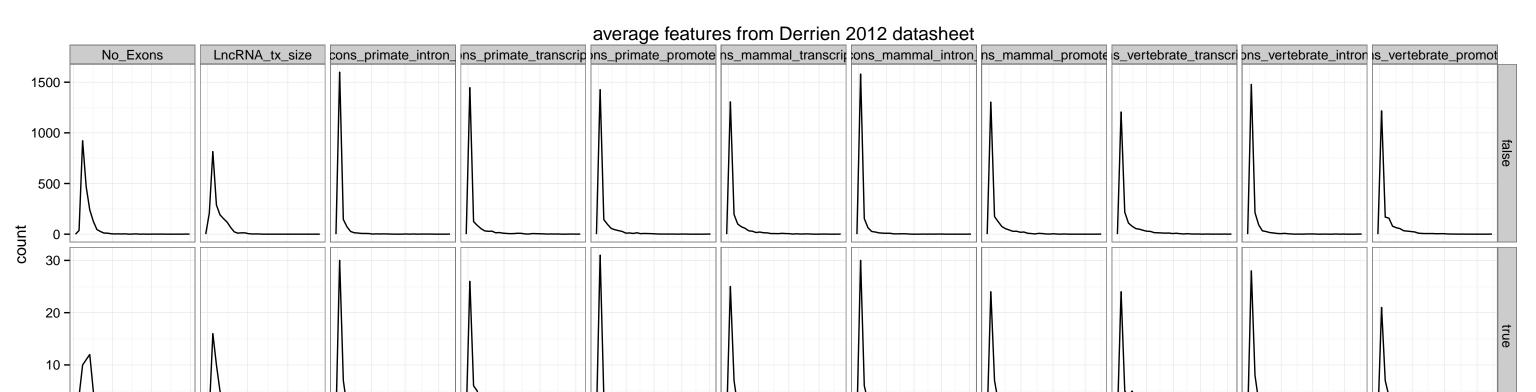lncRNA group = IDRlessthan0_1 ::: data cols = lpa
(46 / 1954) = (func/total lncRNA) ::: biotype = remove_antisense
subset =
46 lncRNAs out of 1954 total lncRNAs

average features from Derrien 2012 datasheet