lncRNA group = IDRlessthan0_2 ::: data cols = lpa
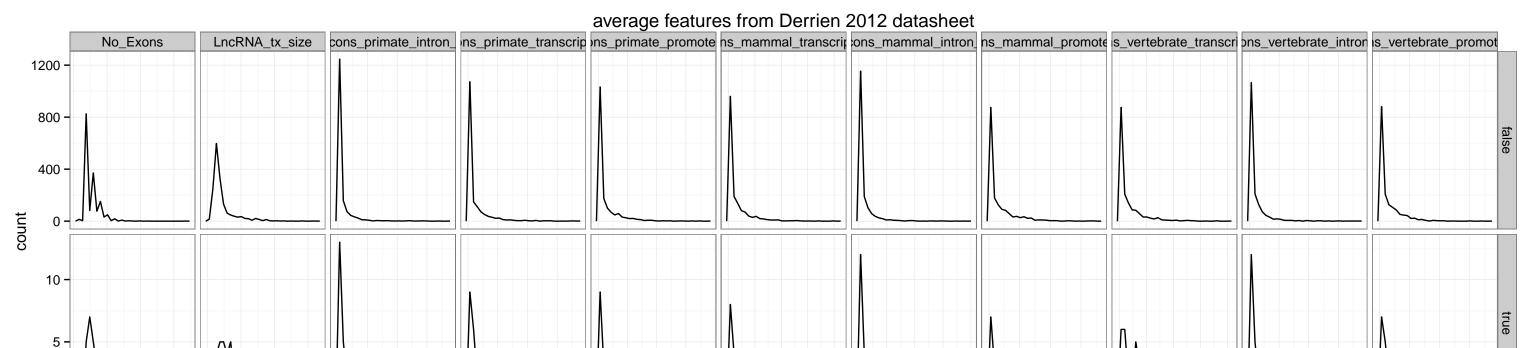(30 / 1675) = (func/total lncRNA) ::: biotype = antisense
subset =
30 lncRNAs out of 1675 total lncRNAs

average features from Derrien 2012 datasheet