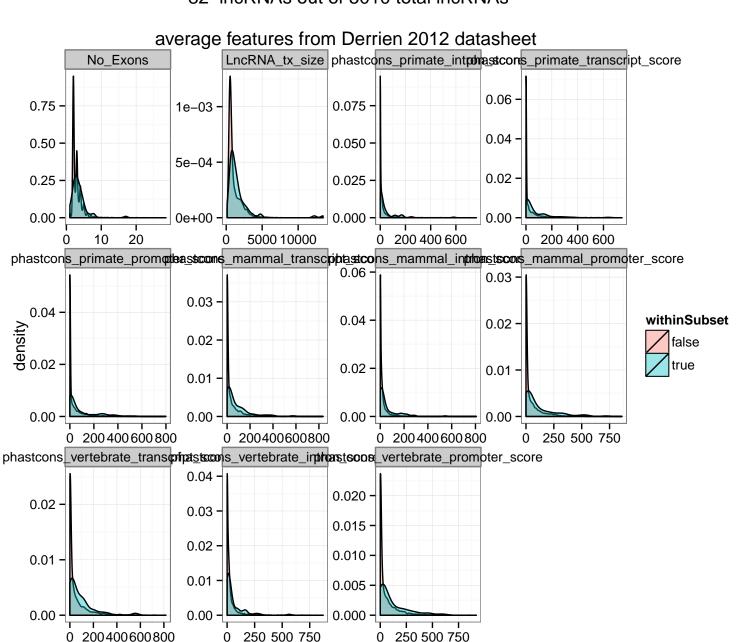lncRNA group = IDRlessthan0_2 ::: data cols = lnpa
(82 / 5010) = (func/total lncRNA) ::: biotype = all_biotypes
subset =
82 lncRNAs out of 5010 total lncRNAs

average features from Derrien 2012 datasheet