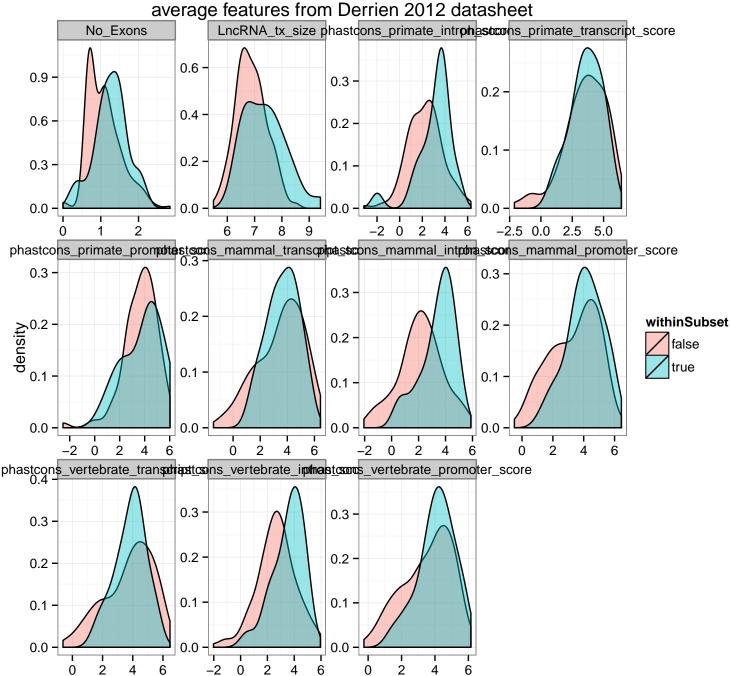
IncRNA group = IDRlessthan0\_01 ::: data cols = bothPullDowns (25 / 243) = (func/total IncRNA) ::: biotype = all\_biotypes average features from Derrien 2012 datasheet



value of facet label