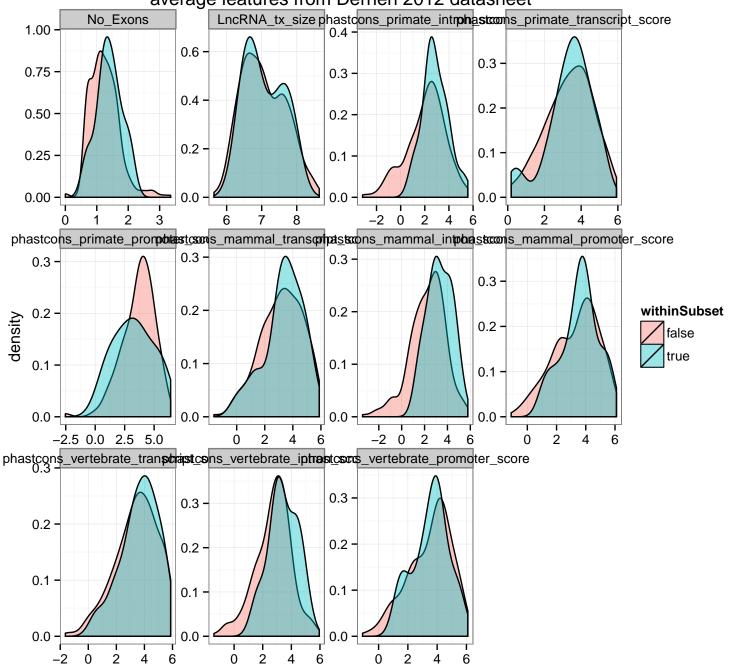
IncRNA group = IDRlessthan0\_2 ::: data cols = Ipa ( 17 / 328 ) = (func/total IncRNA) ::: biotype = processed\_transcript average features from Derrien 2012 datasheet



value of facet label