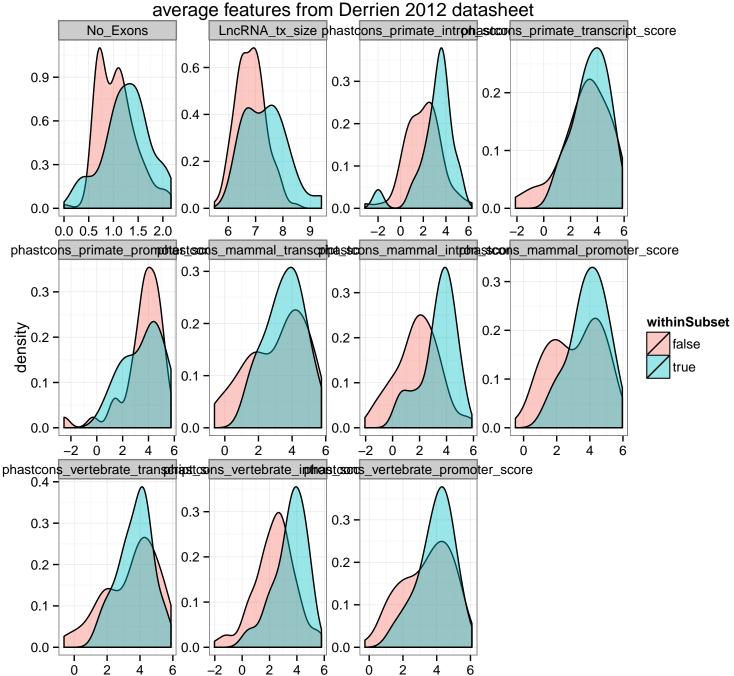
IncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (21 / 146) = (func/total IncRNA) ::: biotype = lincProcTrans average features from Derrien 2012 datasheet



value of facet label