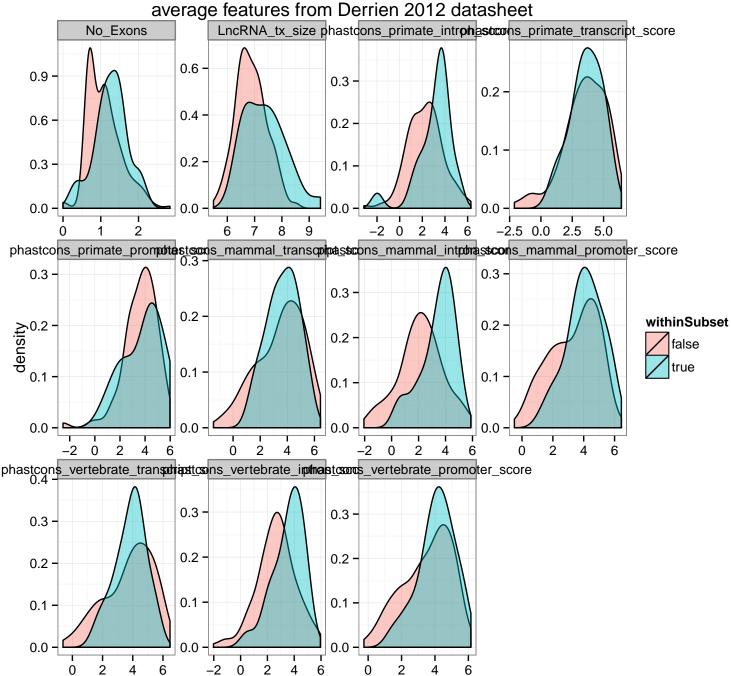
IncRNA group = IDRlessthan0_01 ::: data cols = Inpa (25 / 242) = (func/total IncRNA) ::: biotype = all_biotypes average features from Derrien 2012 datasheet



value of facet label