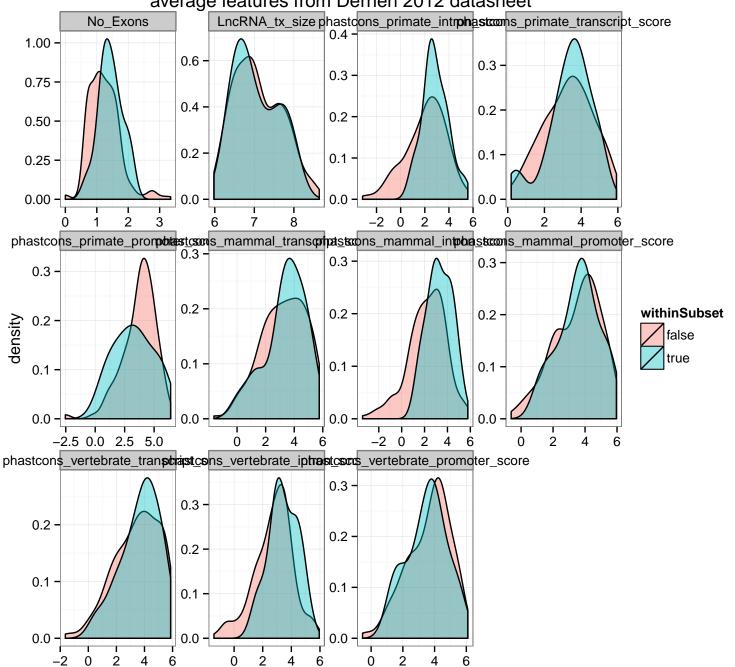
IncRNA group = IDRlessthan0_1 ::: data cols = Inpa (16 / 231) = (func/total IncRNA) ::: biotype = processed_transcript average features from Derrien 2012 datasheet



value of facet label