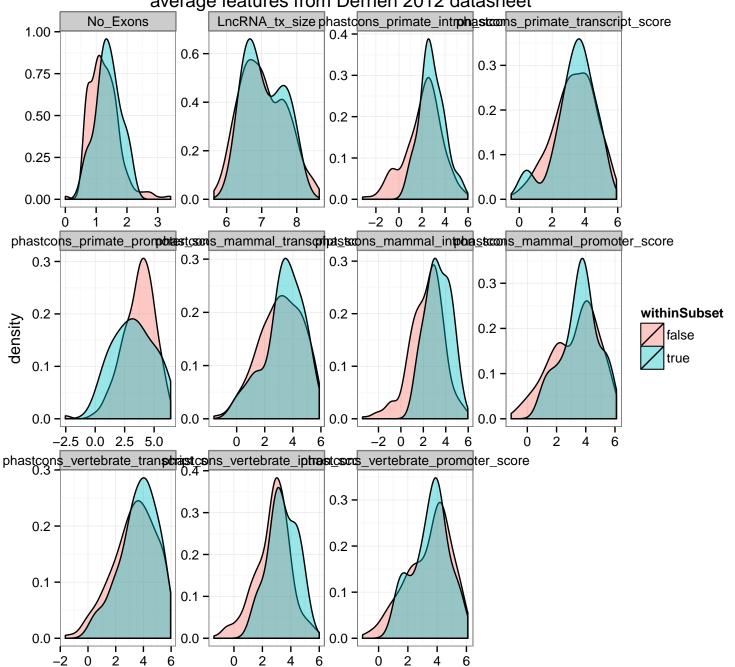
IncRNA group = allLncRNA ::: data cols = Inpa (17 / 372) = (func/total IncRNA) ::: biotype = processed_transcript average features from Derrien 2012 datasheet



value of facet label