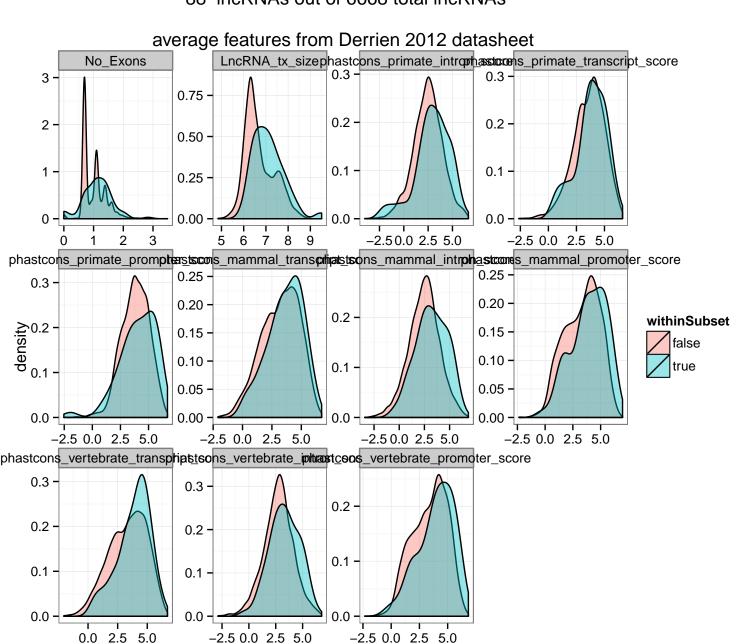
IncRNA group = allLncRNA ::: data cols = Inpa ( 88 / 6668 ) = (func/total IncRNA) ::: biotype = all\_biotypes subset = 88 IncRNAs out of 6668 total IncRNAs



value of facet label