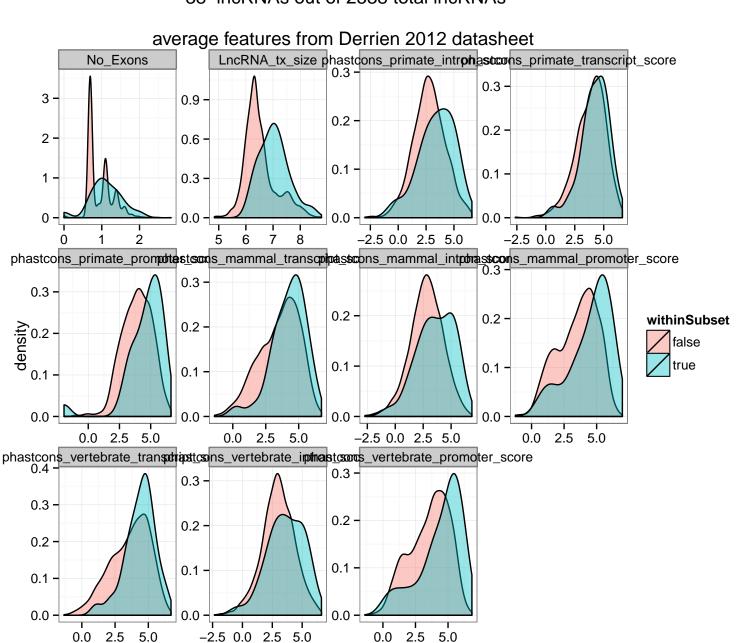
IncRNA group = allLncRNA ::: data cols = bothPullDowns (38 / 2538) = (func/total lncRNA) ::: biotype = antisense subset = 38 lncRNAs out of 2538 total lncRNAs



value of facet label