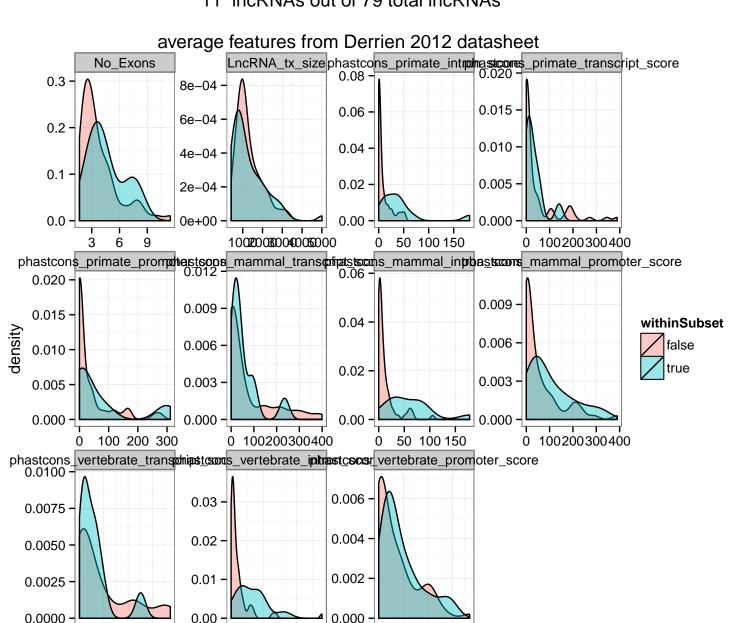
IncRNA group = IDRIessthan0_01 ::: data cols = lpa (11/79) = (func/total lncRNA) ::: biotype = processed_transcript subset = IncRNAs out of 79 total IncRNAs



0.000

value of facet label

0 100 200 300

200

100

0

100 200 300