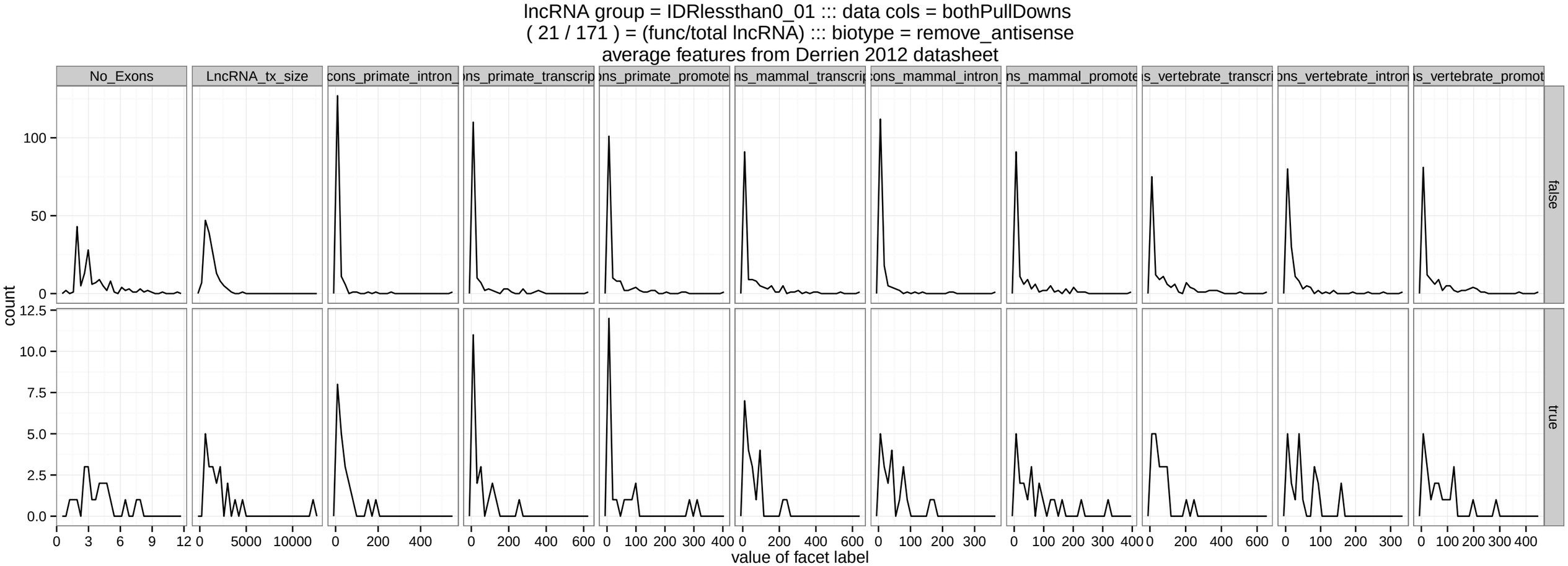
lncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns
(21 / 171) = (func/total lncRNA) ::: biotype = remove_antisense
average features from Derrien 2012 datasheet
No_Exons
LncRNA_tx_size
cons_primate_intron_
ns_primate_transcrip
ons_primate_promote
ns_mammal_transcrip
cons_mammal_intron_
ns_mammal_promote
s_vertebrate_transcri
ons_vertebrate_intron
s_vertebrate_promot
false
true
count
value of facet label