

gma-miR156o/Glyma.05G019000.1

Degradome Abundance (TP10M)

120

100

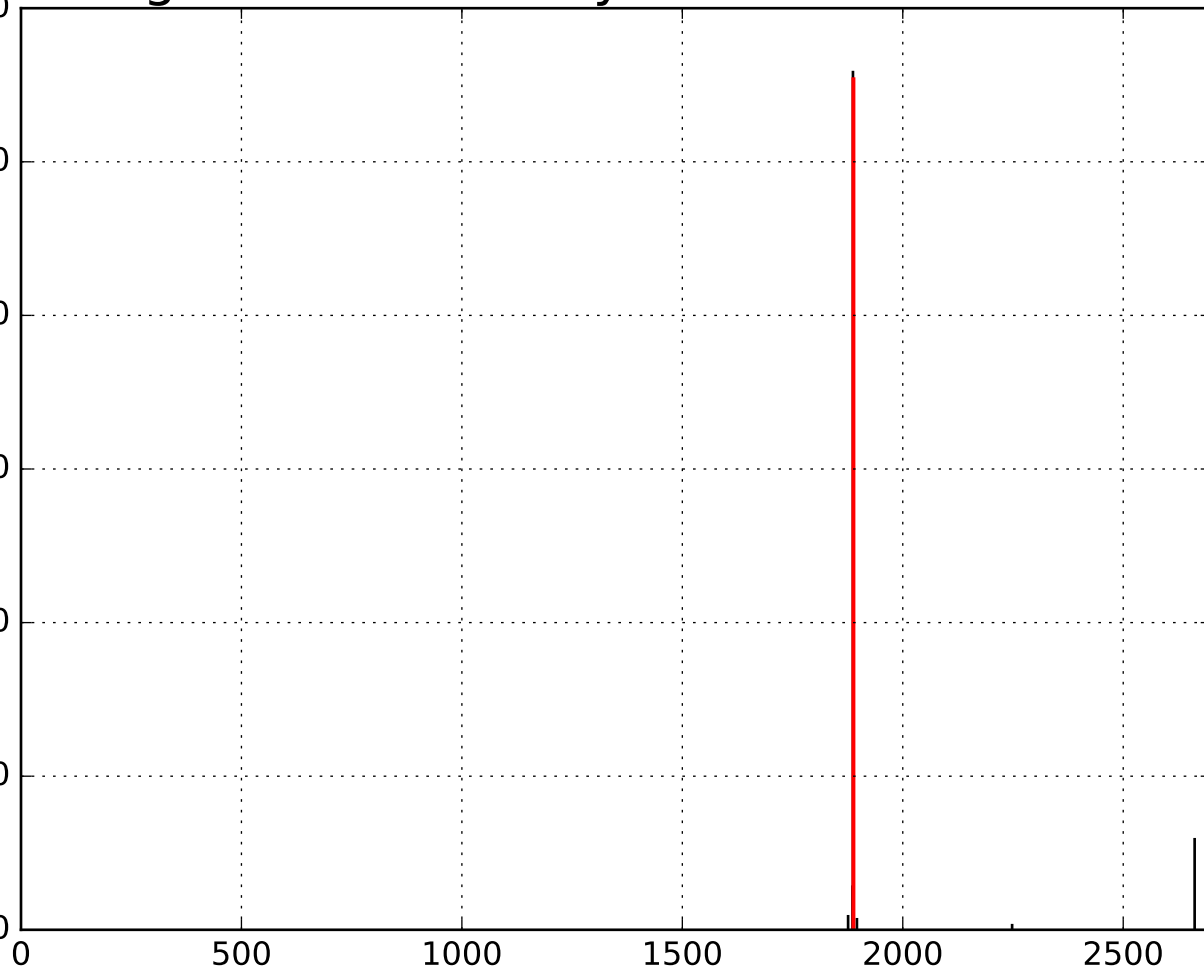
80

60

40

20

0



Nucleotide position (nt)