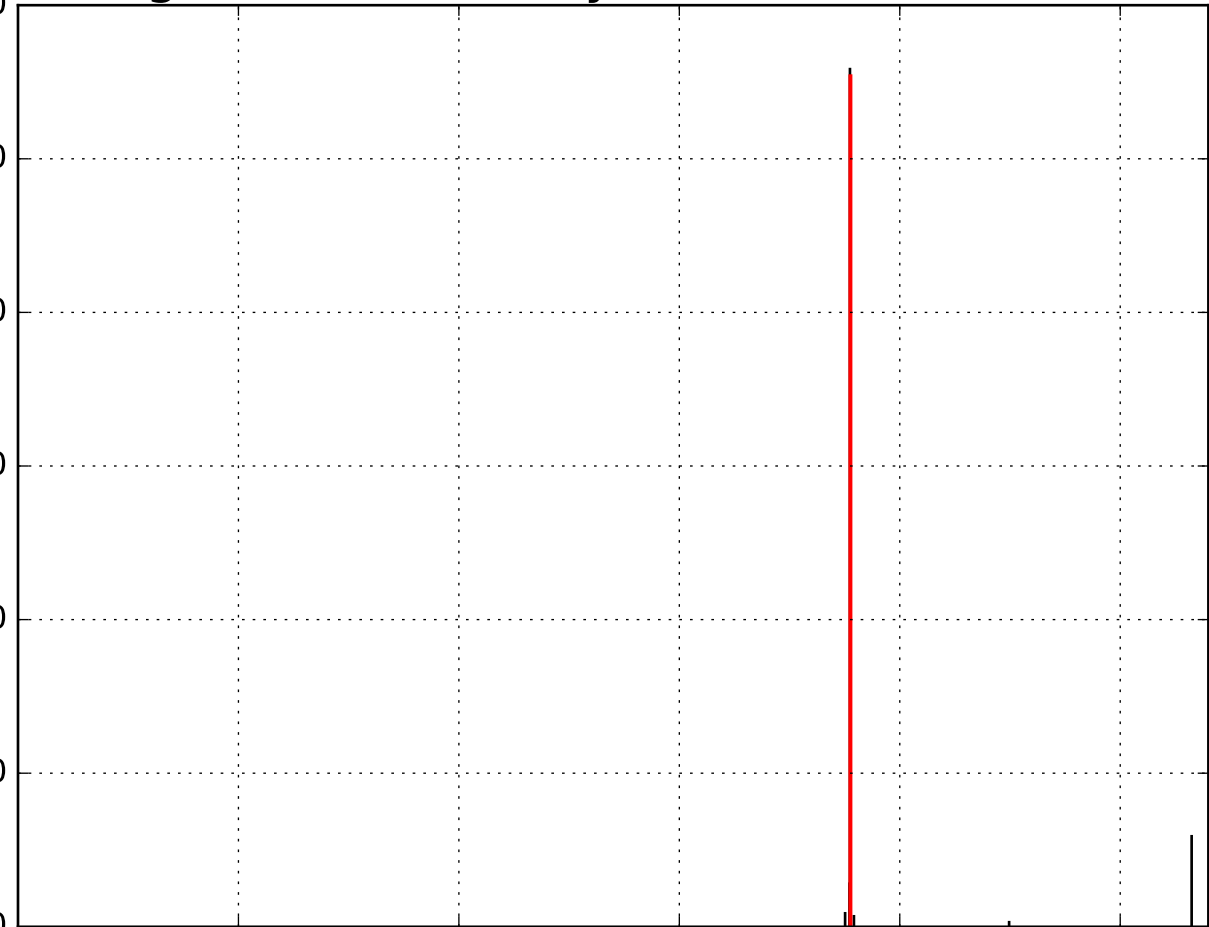


gma-miR156r/Glyma.05G019000.1

Degradome Abundance (TP10M)

120
100
80
60
40
20
0



Nucleotide position (nt)