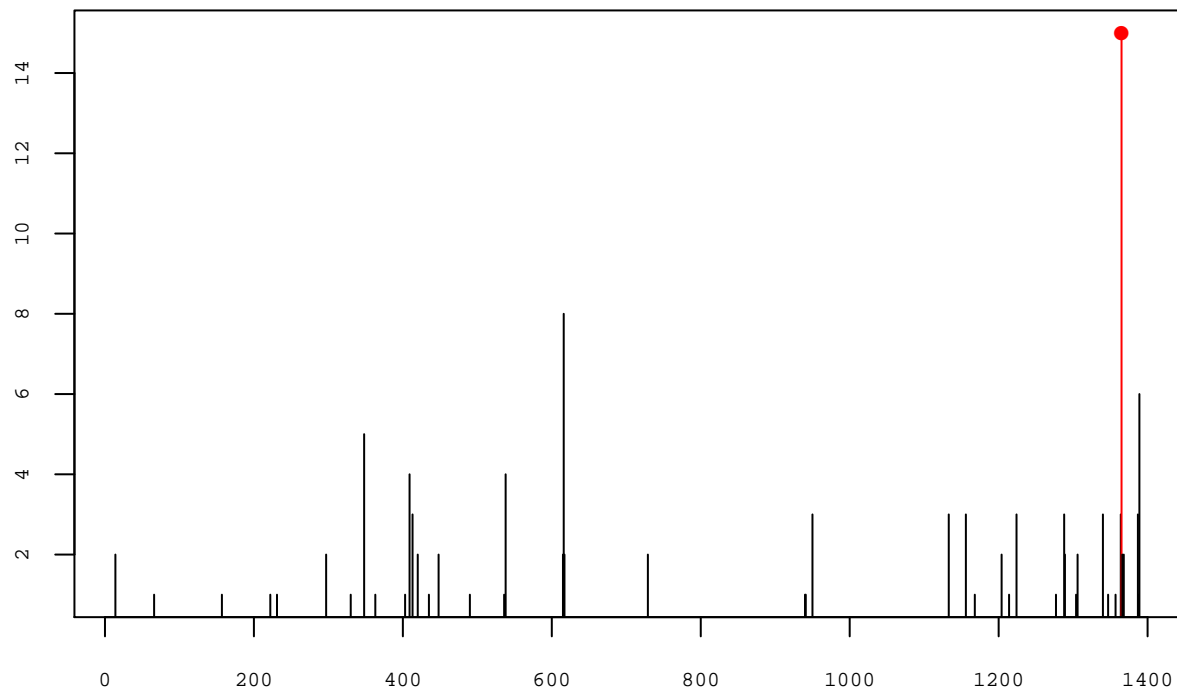


PggD12466_c24_g1_i1

5' CGTTGTTTCGTGAACAGCTCGTGTTCGGCTCGT '3
|||||
3' ACAAGCACTTGTTCGAGCACAAC '5

Fragment Abundance



Transcript position

Cleavage site: 1365

Tag abundance: 15

Weighted abundance: 0.29

Category: 0

sRNA abundance: 13

Alignment score: 2

MFE ratio: 0.93

p-value: 0.028