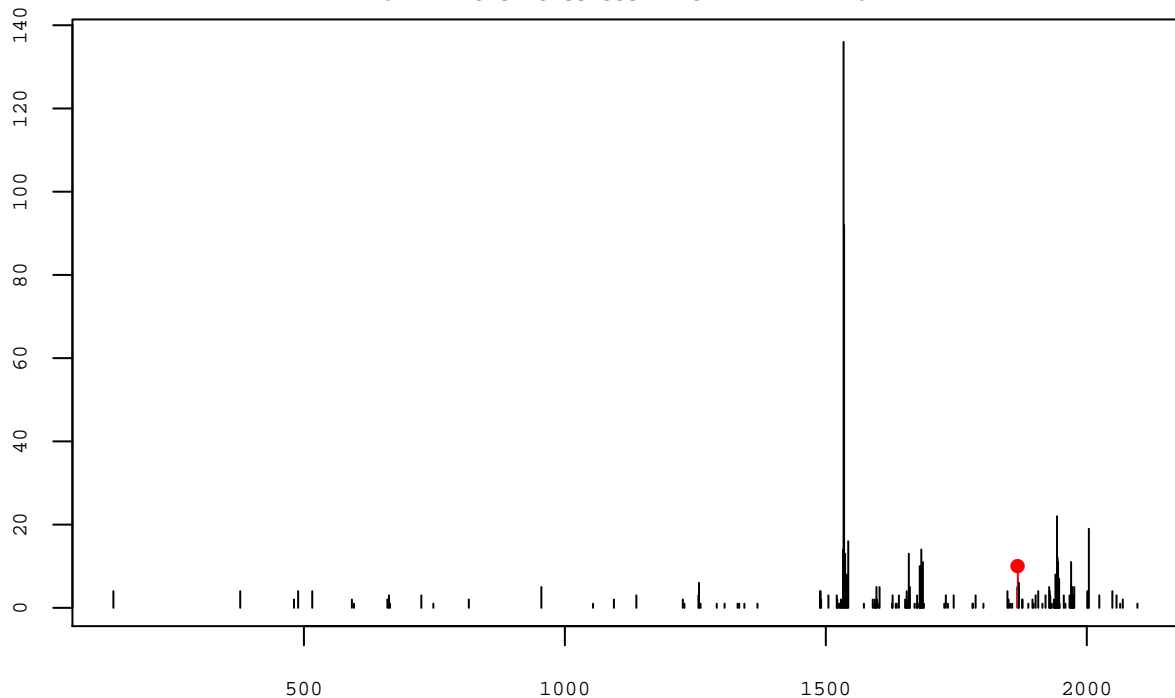


PggD1935_c0_g1_i1

5' CAAACAGGTCAGGTGCATTTTCAAATGTCAC '3
||| |o| ||||| | ||||| |
3' TTGTCTAGTCCACGAAAAGTTT '5

Fragment Abundance



Transcript position

Cleavage site: 1868 Tag abundance: 10 Weighted abundance: 10 Category: 2
sRNA abundance: 6 Alignment score: 2.5 MFE ratio: 0.831 p-value: 0.009