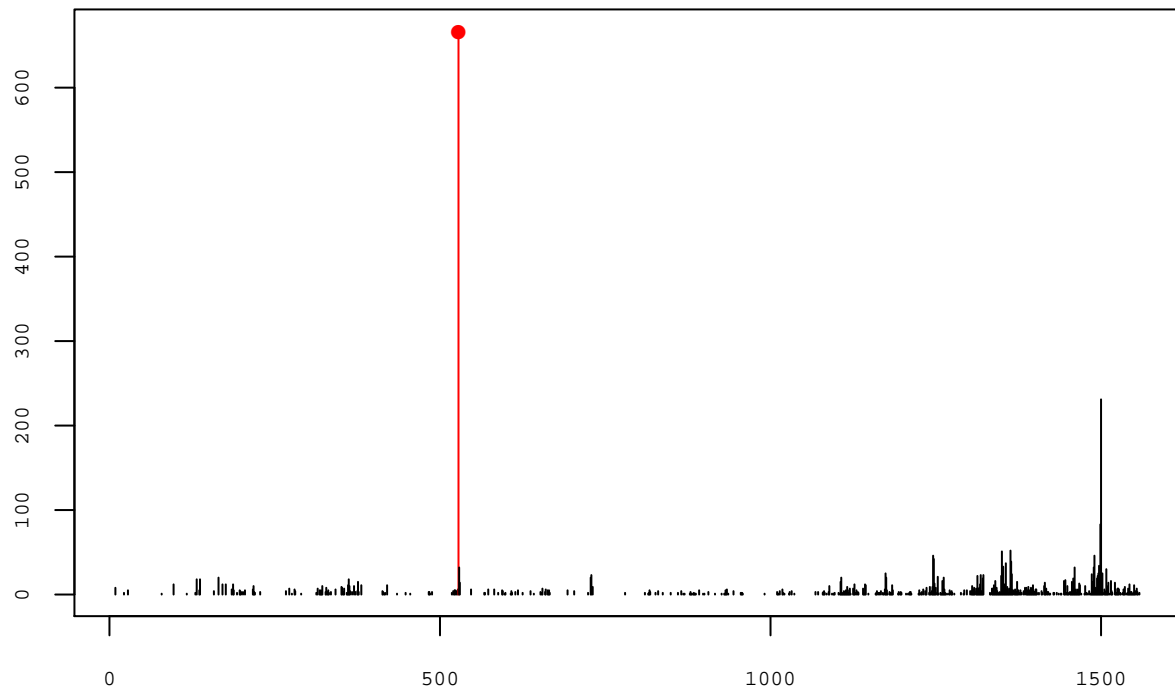


PggD12227_c16_g1_i1

5' GGAGAAAAAGCTCCATTCAATCCAAAGCGCGC '3
| | | | | | | | | | | | | | | | | |
3' TCTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Transcript position

Cleavage site: 528 Tag abundance: 666 Weighted abundance: 666 Category: 0
sRNA abundance: 8438 Alignment score: 3 MFE ratio: 0.705 p-value: 0.008