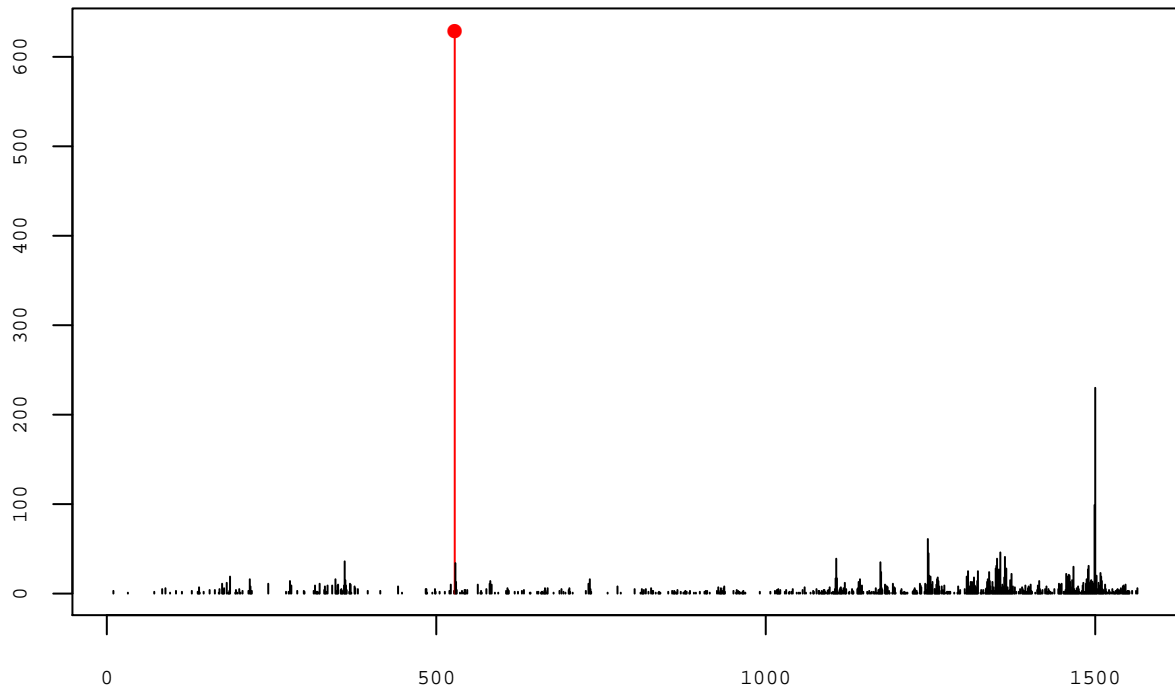


PggD12227_c16_g1_i1

5' GGAGAAAAAGCTCCATTCAATCCAAAGCGCGC '3
 ||| |||
3' CTCGAGGGAAGTTAGGTTC '5

Fragment Abundance



Transcript position

Cleavage site: 528 Tag abundance: 629 Weighted abundance: 629 Category: 0
sRNA abundance: 18 Alignment score: 4 MFE ratio: 0.705 p-value: 0.013