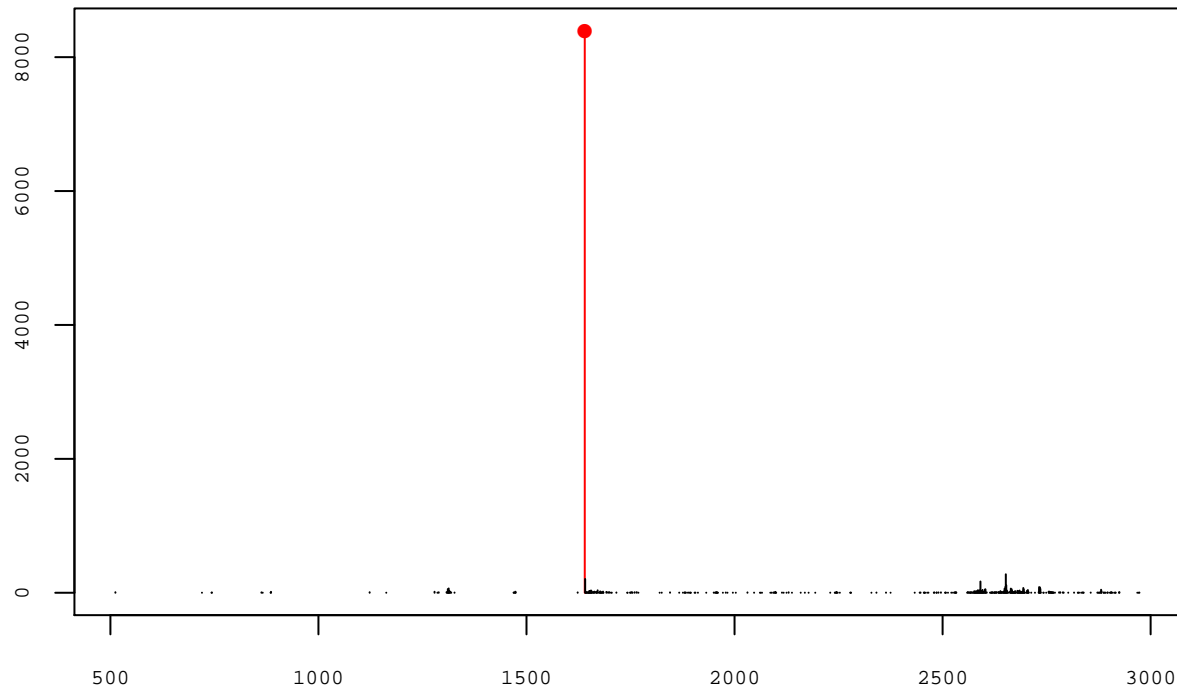


PggD1242_c13_g1_i2

5' GCAGCTAGCATAACAGGGAGCCAGGCAAATTCT 3'
|||||
3' CGTATGTCCTCGGTCCGT 5'

Fragment Abundance



Transcript position

Cleavage site: 1640 Tag abundance: 8392 Weighted abundance: 4196 Category: 0
sRNA abundance: 11 Alignment score: 0 MFE ratio: 1 p-value: 0.001