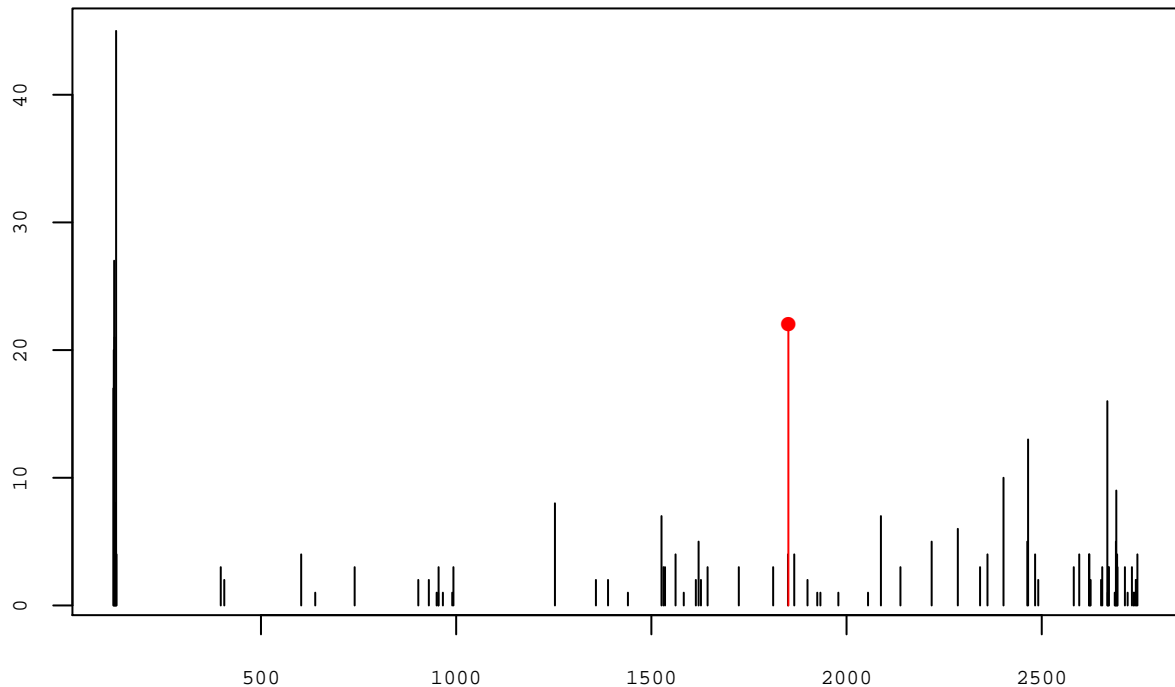


PggD14795_c78_g1_i2

5' AAATACCGAGCTCACTTCAATCCAAGAGAGCC 3'
 ||||| ||||| ||||| |
3' CTCGAGGGAAGTTAGGTTT 5'

Fragment Abundance



Transcript position

Cleavage site: 1851 Tag abundance: 22 Weighted abundance: 5.5 Category: 2
sRNA abundance: 1110 Alignment score: 2.5 MFE ratio: 0.789 p-value: 0.025