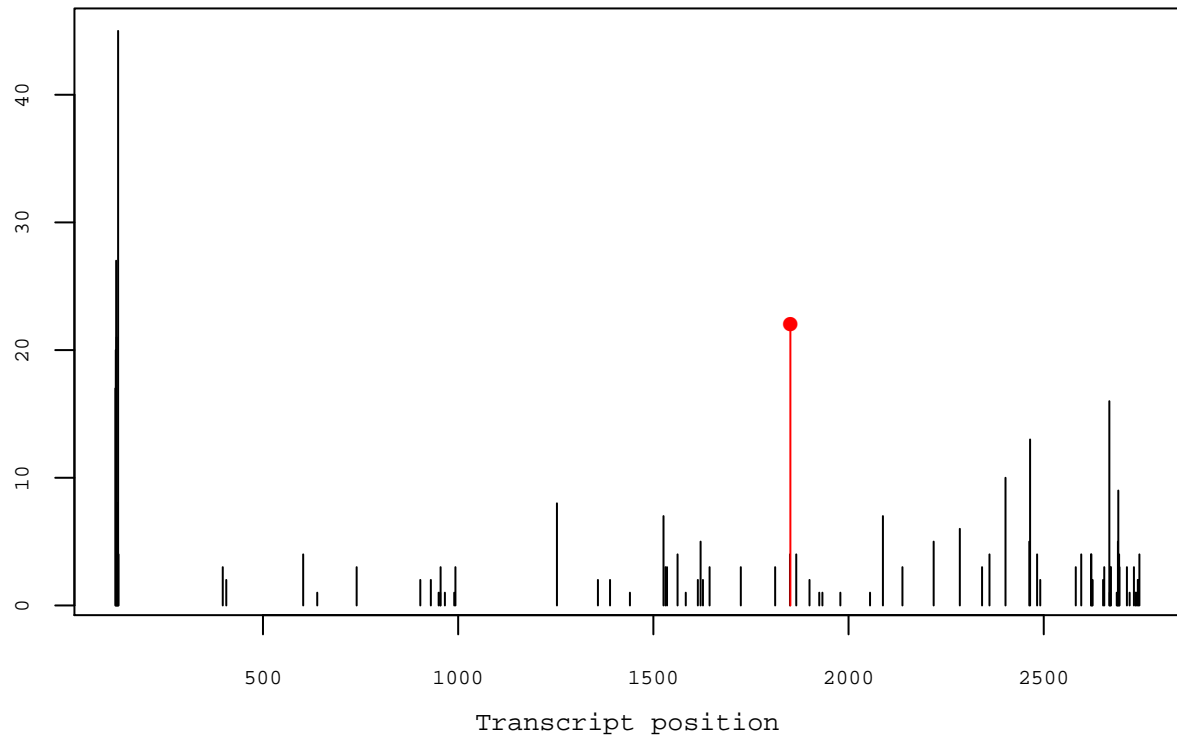


PggD14795_c78_g1_i2

5' AAATACCGAGCTCACTTCAATCCAAGAGAGCC '3
| | | | | | | | | | | | | | | | | |
3' GTCTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Cleavage site: 1851 Tag abundance: 22 Weighted abundance: 5.5 Category: 2
sRNA abundance: 8957 Alignment score: 3.5 MFE ratio: 0.738 p-value: 0.05