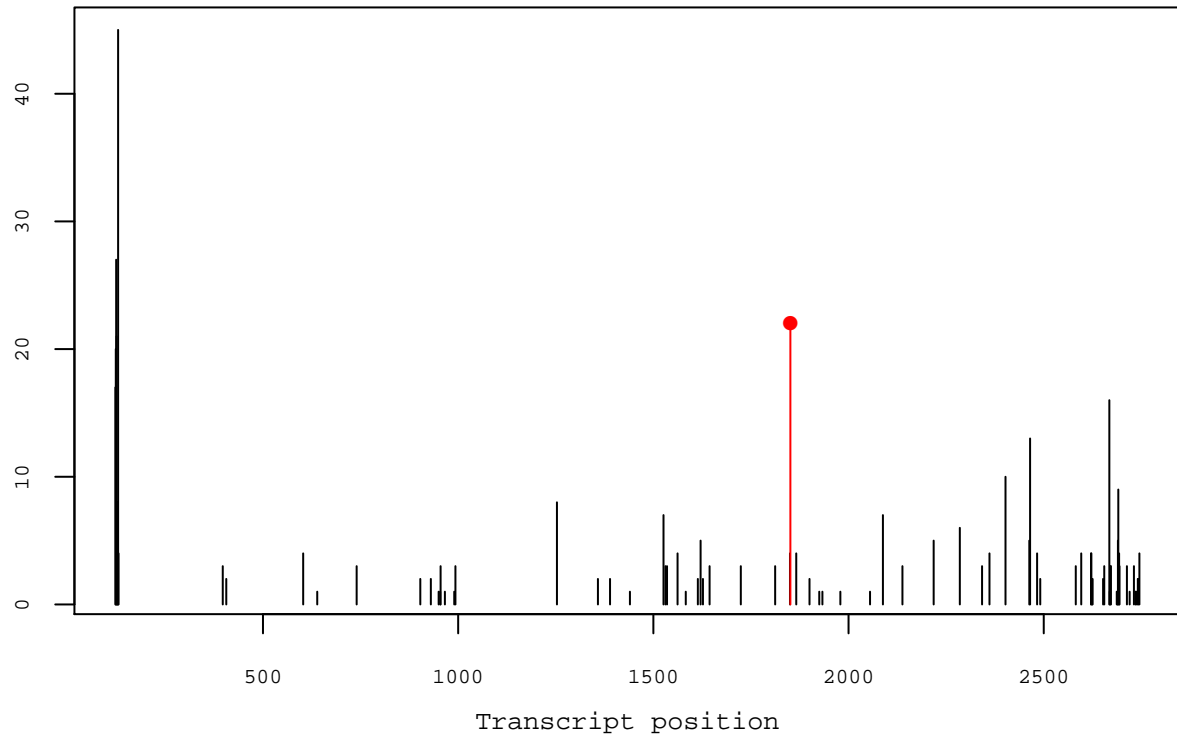


PggD14795_c78_g1_i2

5' AAATACCGAGCTCACTTCAATCCAAGAGAGCC 3'
 ||||| ||||| |||||
3' CTCGAGGGAAGTTAGGTTC 5'

Fragment Abundance



Cleavage site: 1851 Tag abundance: 22 Weighted abundance: 5.5 Category: 2
sRNA abundance: 18 Alignment score: 2 MFE ratio: 0.798 p-value: 0.02