

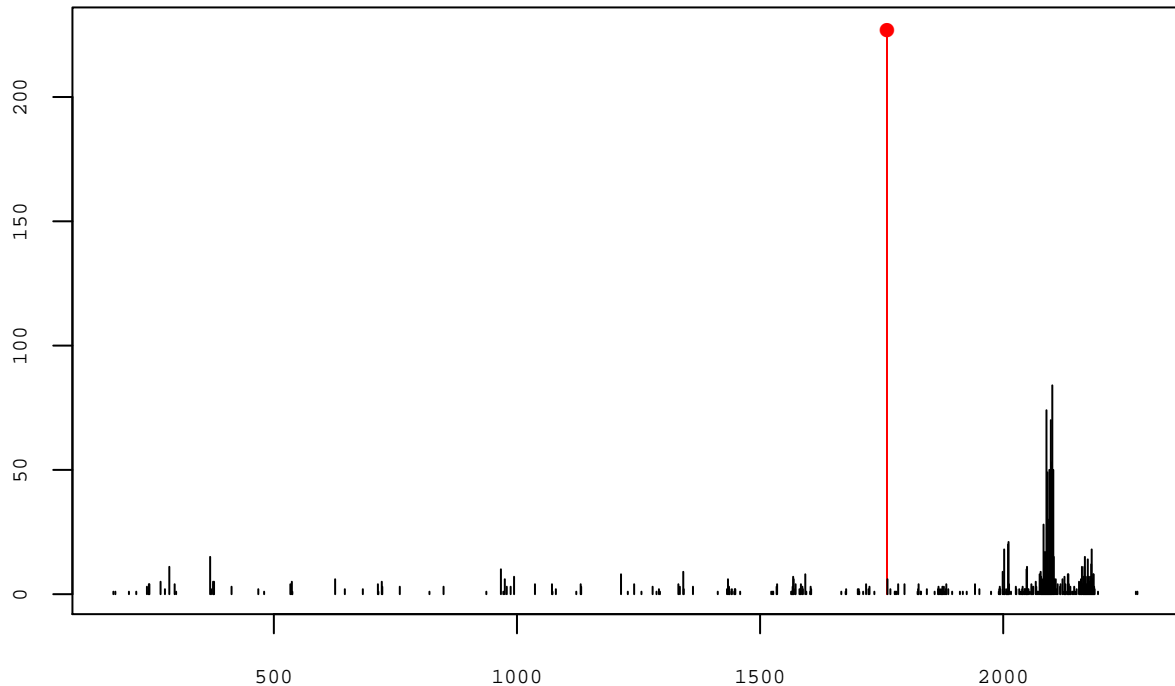
PggD5310_c30_g1_i1

5' TTCCACAGGGGGA-CCCTTCAGTCCAATCCATC '3

|||||o| |||||

3' TCCCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Transcript position

Cleavage site: 1761 Tag abundance: 227 Weighted abundance: 113.5 Category: 0
sRNA abundance: 13228 Alignment score: 2.5 MFE ratio: 0.794 p-value: 0.004