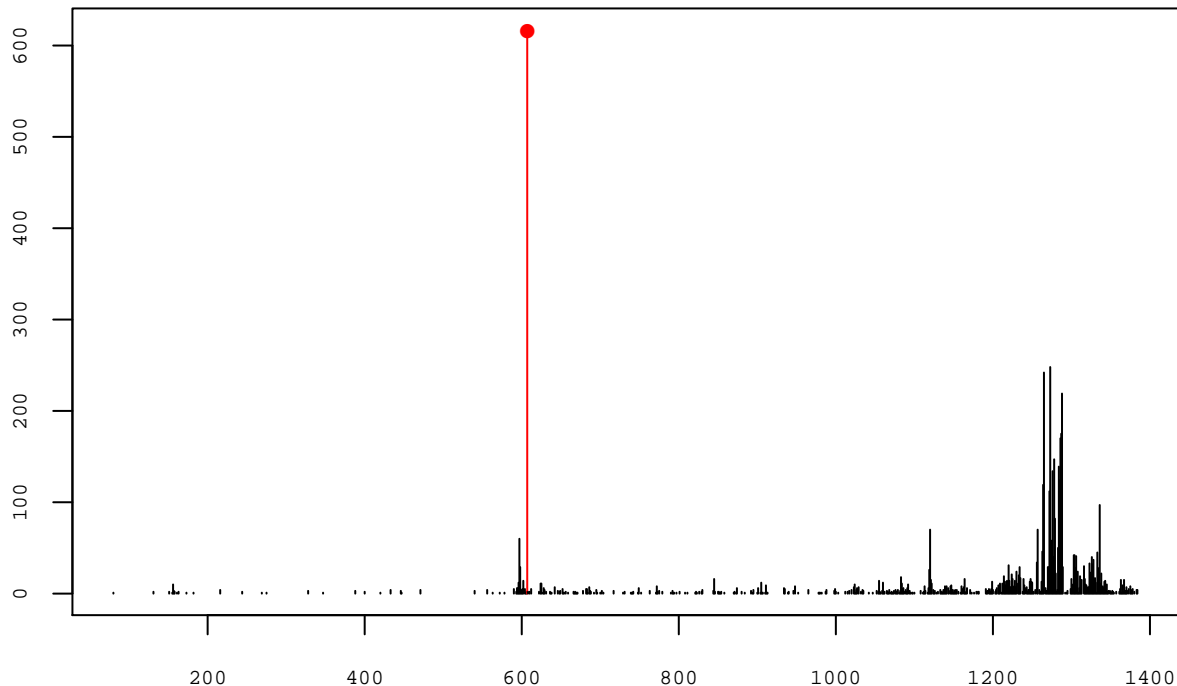


PggD4125_c99_g1_i1

5' AAGAACCGTTCAAGAAAGCCTGTGGAAATCTC '3
|||||
3' TCAAGTTCTTTTCG-ACACCTT '5

Fragment Abundance



Transcript position

Cleavage site: 607	Tag abundance: 616	Weighted abundance: 154	Category: 0
sRNA abundance: 117	Alignment score: 3	MFE ratio: 0.892	p-value: 0.013