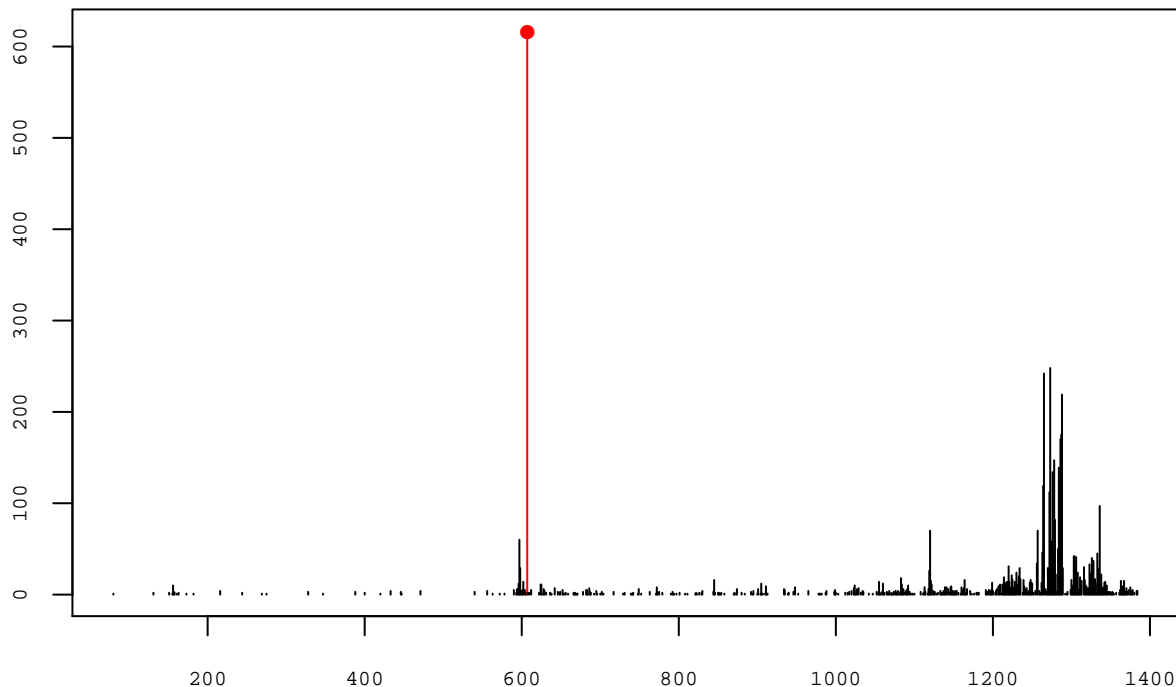


PggD4125_c99_g1_i1

5' AAGAACCGTTCAAGAAAGCCTGTGGAAATCTC '3
|||||
3' CAAGTTCTTTCG-ACACCTT '5

Fragment Abundance



Transcript position

Cleavage site: 607 Tag abundance: 616 Weighted abundance: 154 Category: 0
sRNA abundance: 56 Alignment score: 2 MFE ratio: 0.893 p-value: 0.011