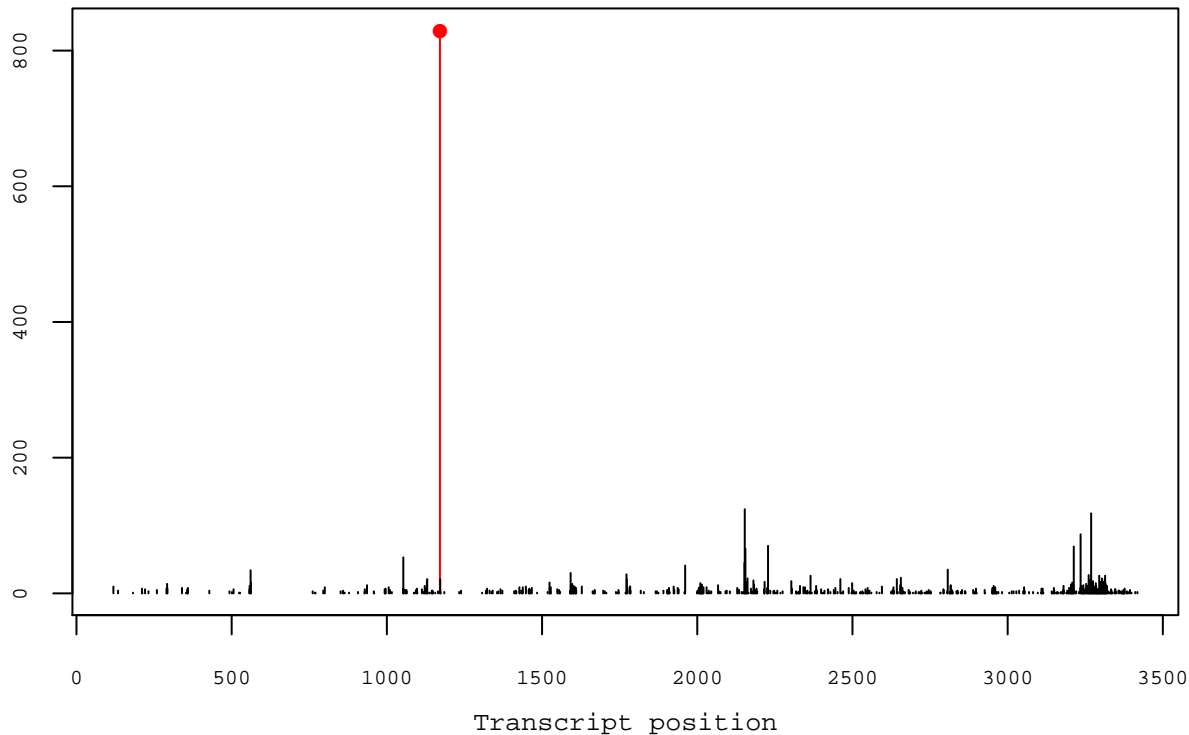


PggD9924_c6_g1_i1

5' GCTTGTTGGAATGAAGCCTGGTCCGGATTCCA '3
|||||
3' CCCTTACTTCGGACCAGGCT '5

Fragment Abundance



| | | | |
|---------------------|----------------------|-------------------------|-------------|
| Cleavage site: 1171 | Tag abundance: 829 | Weighted abundance: 829 | Category: 0 |
| sRNA abundance: 417 | Alignment score: 1.5 | MFE ratio: 0.929 | p-value: 0 |