

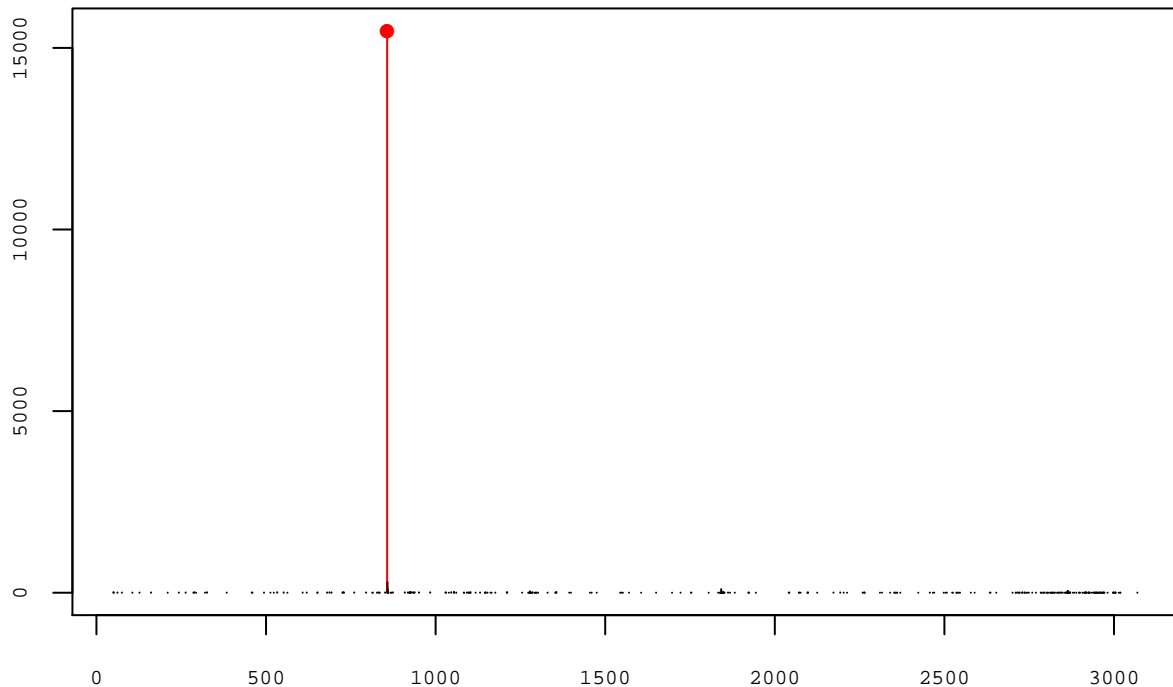
PggD11091_c20_g1_i1

5' GATGGTCGGGATGAAGCCTGGTCCGGATTCTA '3

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3' TCCCTTACTTCGGACCAGGCT '5

Fragment Abundance



Transcript position

| | | | |
|---------------------|----------------------|--------------------------|----------------|
| Cleavage site: 857 | Tag abundance: 15468 | Weighted abundance: 7734 | Category: 0 |
| sRNA abundance: 696 | Alignment score: 3.5 | MFE ratio: 0.825 | p-value: 0.001 |