

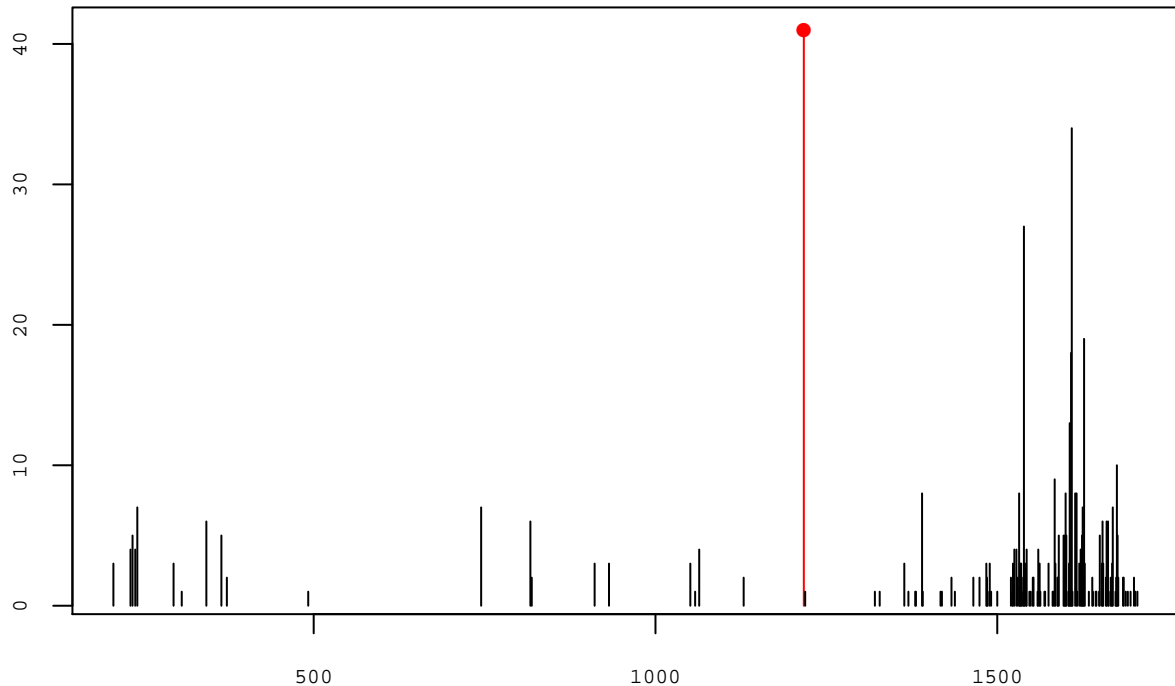
PggD4707\_c6\_g1\_i1

5' TTTAGTAGGGGGA-CCCTTCAGTCCAATTCACC '3

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3' CCCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Transcript position

Cleavage site: 1217 Tag abundance: 41 Weighted abundance: 41 Category: 0  
sRNA abundance: 1130 Alignment score: 2.5 MFE ratio: 0.786 p-value: 0.005