

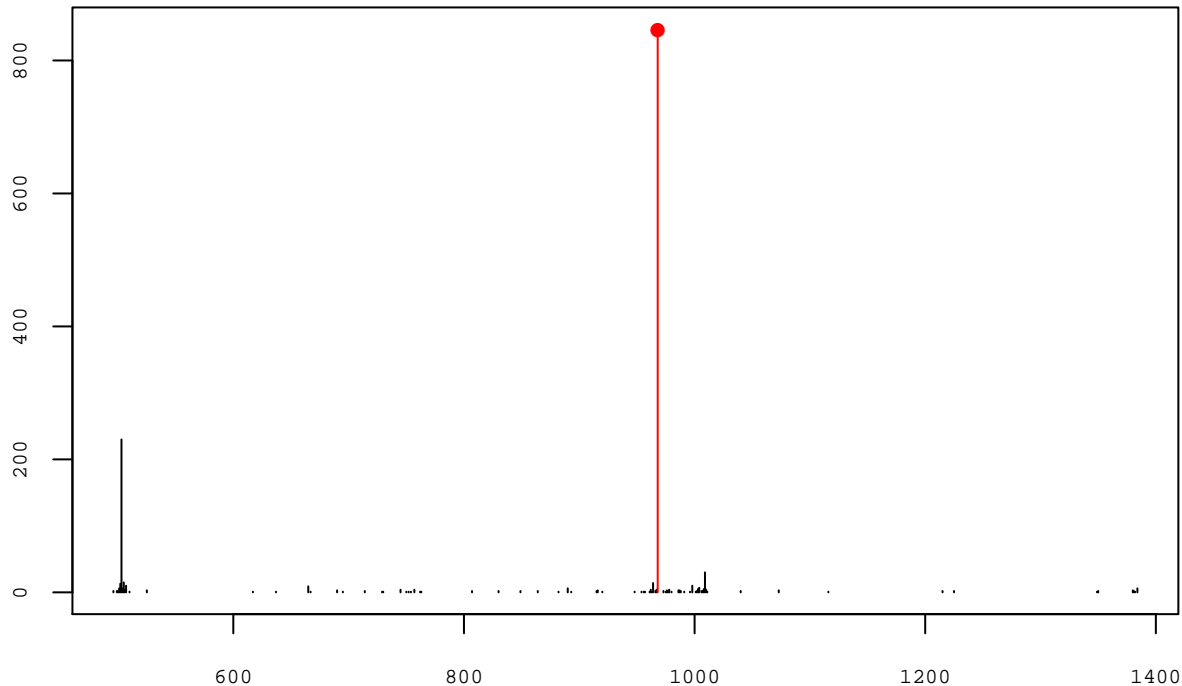
PggD14469\_c62\_g1\_i2

5' TTCCCCTTGCTATCCCTCCTGAGCTGAGCTAG '3

o |||||

3' CCGCGATAGGGAGGACTCGAA '5

Fragment Abundance



Transcript position

Cleavage site: 968 Tag abundance: 846 Weighted abundance: 846 Category: 0  
sRNA abundance: 161 Alignment score: 3.5 MFE ratio: 0.835 p-value: 0.004