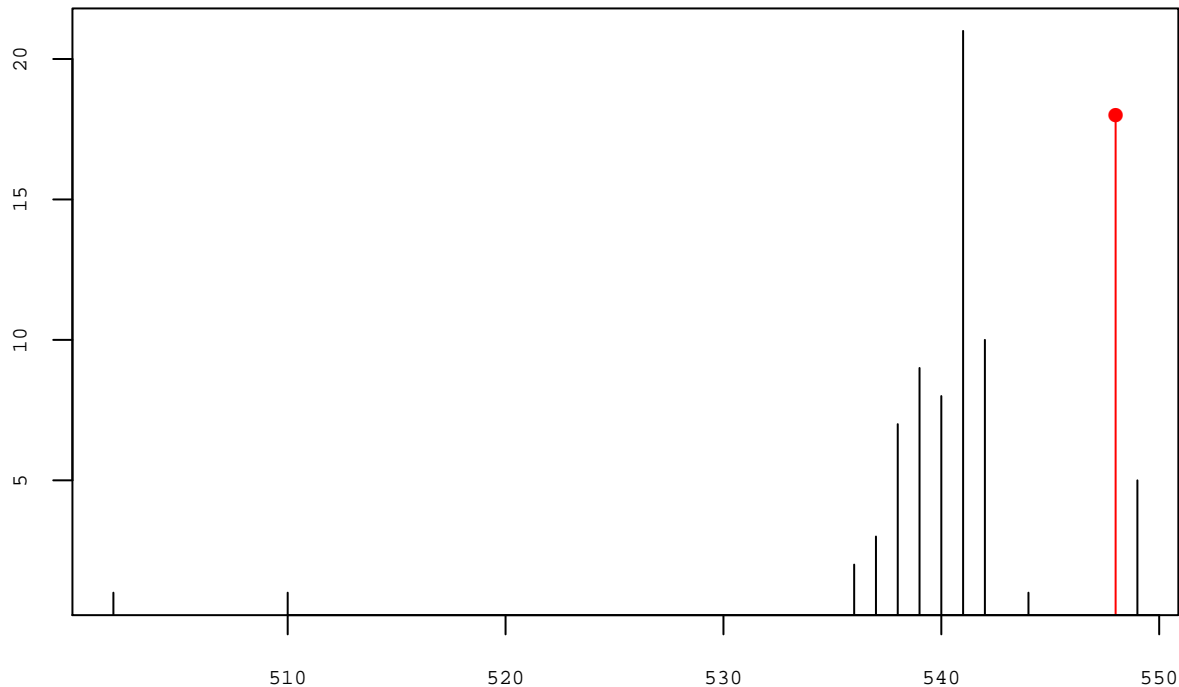


PggD794\_c44\_g1\_i1

5' AATAGCTCGTGTTCGACTCGTTAAGAAGCTCG '3  
o|||||||||||||||||||o  
3' GTCGAGCACAAAGCTGAGCAATTG '5

Fragment Abundance



Transcript position

Cleavage site: 548 Tag abundance: 18 Weighted abundance: 0.947 Category: 2  
sRNA abundance: 6 Alignment score: 2.5 MFE ratio: 0.911 p-value: 0.007