

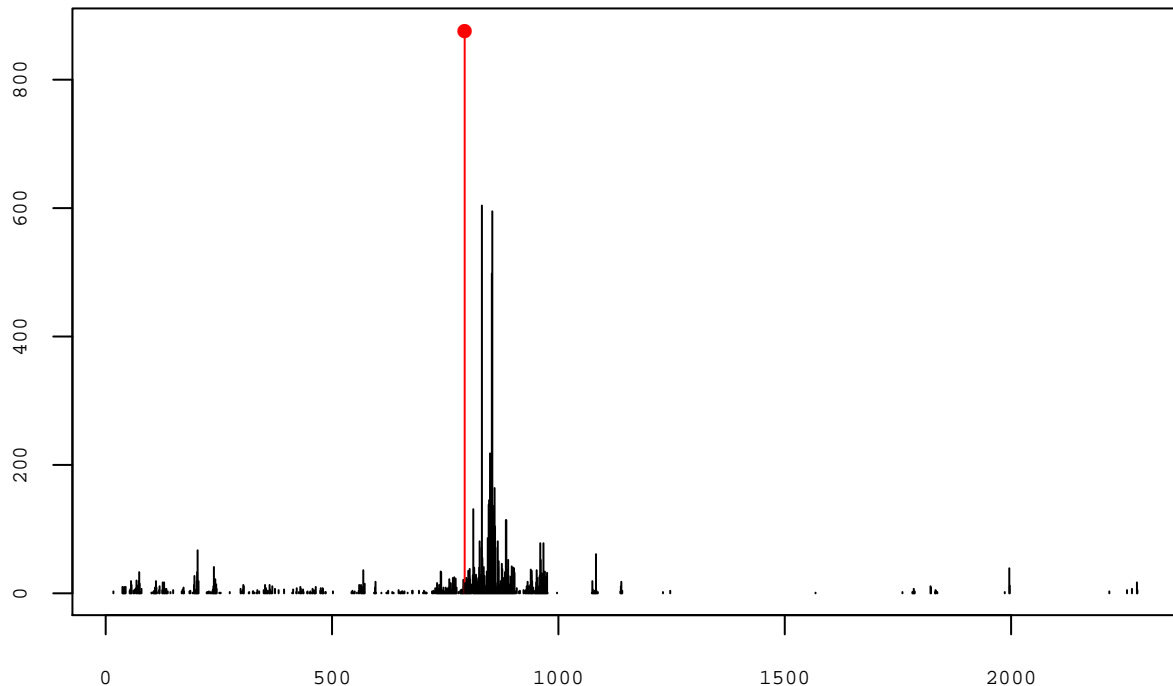
PggD6617\_c1\_g2\_i1

5' TCAAGCATGCTTGCTCTCTTCTGTCAGTAGAT '3

|||||oo|||||||

3' CACGAGTGAGAGAAGACAGT '5

Fragment Abundance



Transcript position

Cleavage site: 793 Tag abundance: 876 Weighted abundance: 876 Category: 0  
sRNA abundance: 26 Alignment score: 2 MFE ratio: 0.858 p-value: 0.008