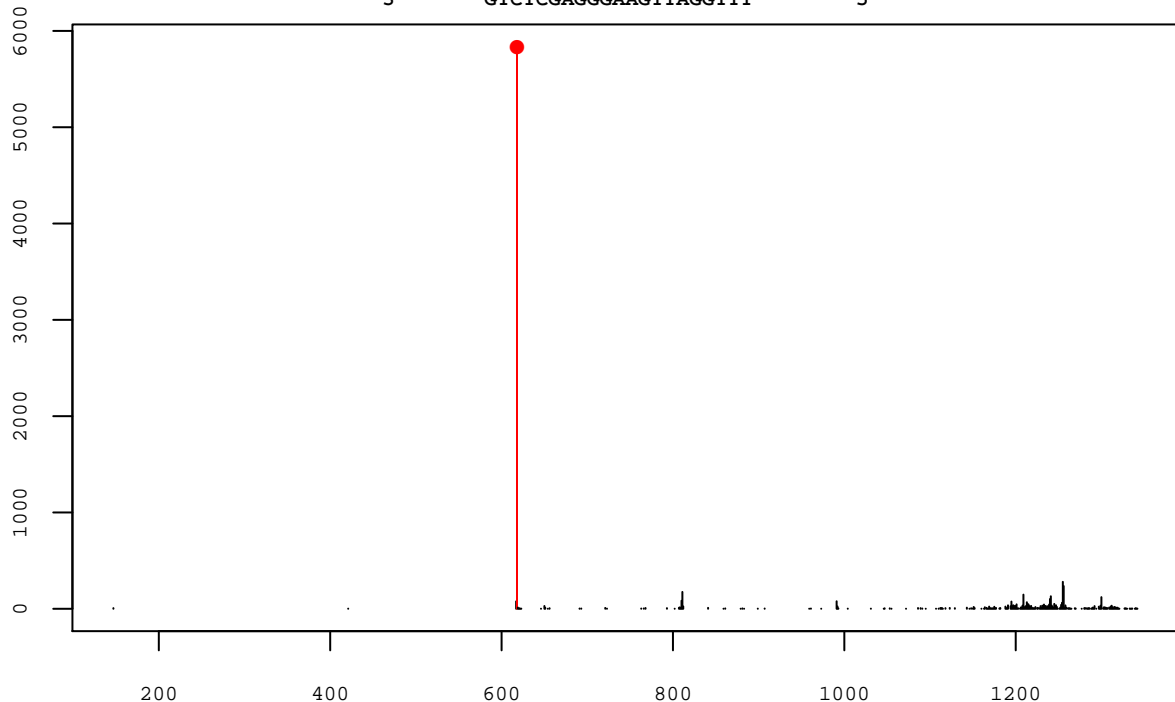


PggD6412\_c14\_g2\_i1

5' GGAAATGGAGCTCCCTTCAAACCAAAGCCATC '3  
oo|||||||||||||  
3' GTCTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Transcript position

Cleavage site: 618    Tag abundance: 5834    Weighted abundance: 5834    Category: 0  
sRNA abundance: 5795    Alignment score: 3    MFE ratio: 0.828    p-value: 0.001