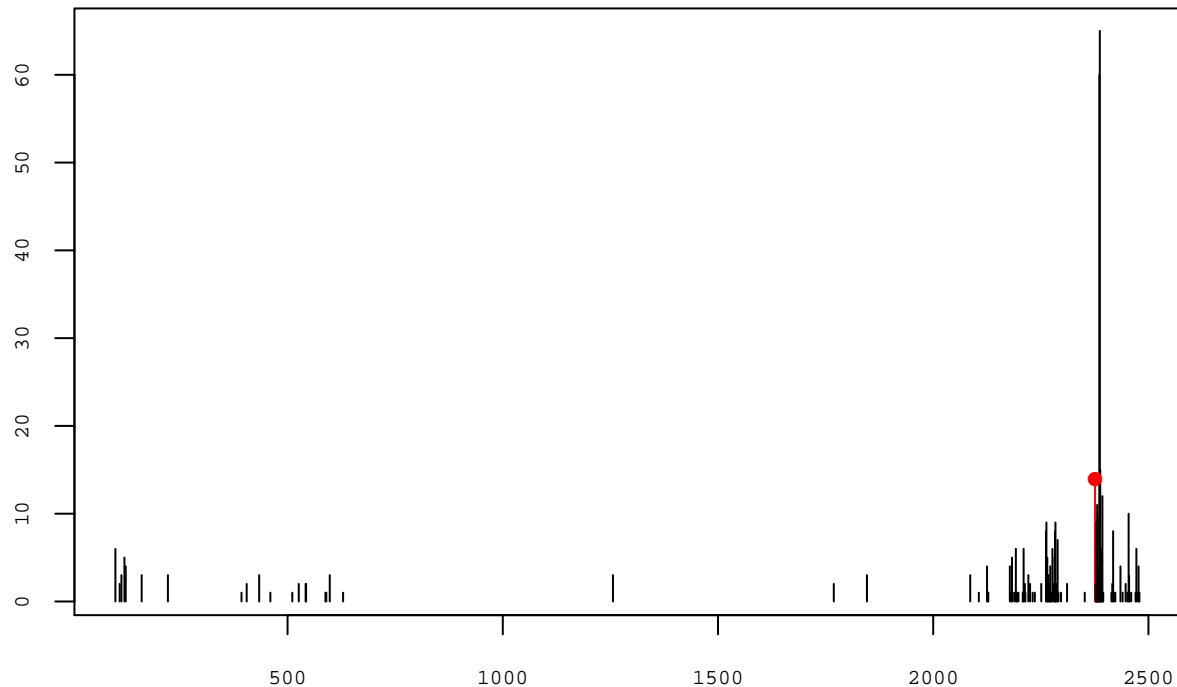


PggD14550\_c21\_g1\_i2

5' TTATTGGGTACGCACGTTCTCTACGTGCGTA '3  
|||||  
3' TAACCCATGCGTGCAAGGAGATGC '5

Fragment Abundance



Transcript position

Cleavage site: 2376 Tag abundance: 14 Weighted abundance: 2.333 Category: 2  
sRNA abundance: 10 Alignment score: 0 MFE ratio: 1 p-value: 0.011