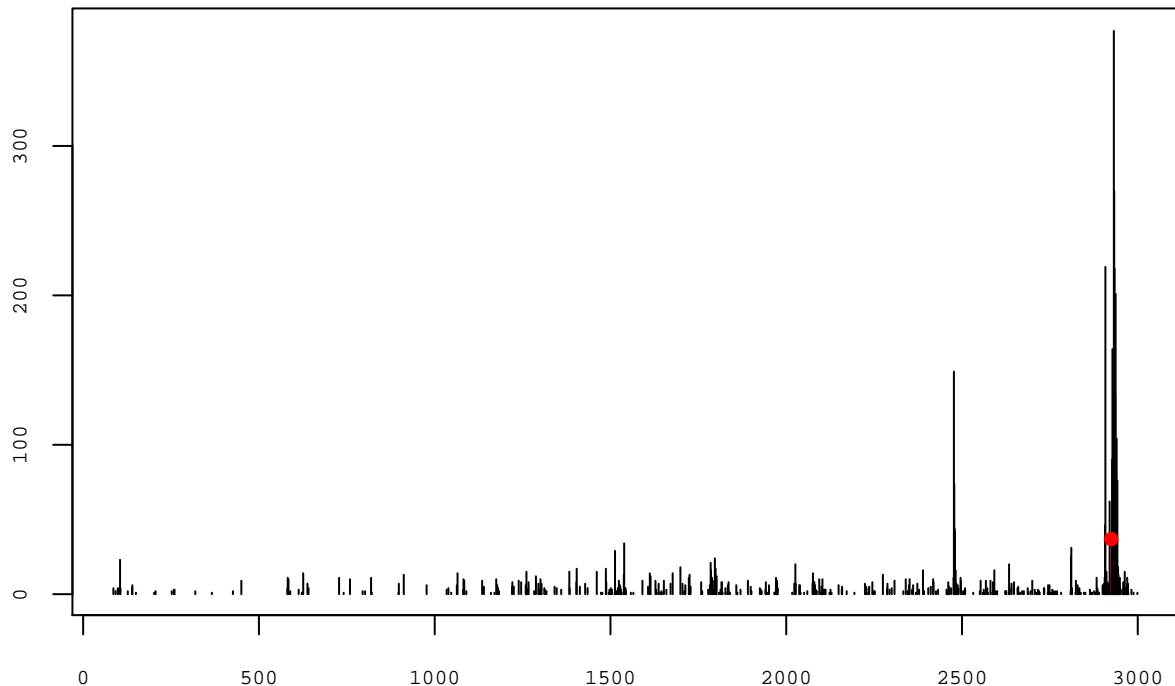


PggD11190\_c0\_g1\_i2

5' TGACATGAT-GTCTGATGATTGAATACTTATAT '3  
          ||||| |||○|||||||  
3'   CTACTACCAGGCTACTAACTTATA       '5

Fragment Abundance



Transcript position

Cleavage site: 2925    Tag abundance: 37    Weighted abundance: 12.333    Category: 2  
sRNA abundance: 5    Alignment score: 3.5    MFE ratio: 0.703    p-value: 0.047