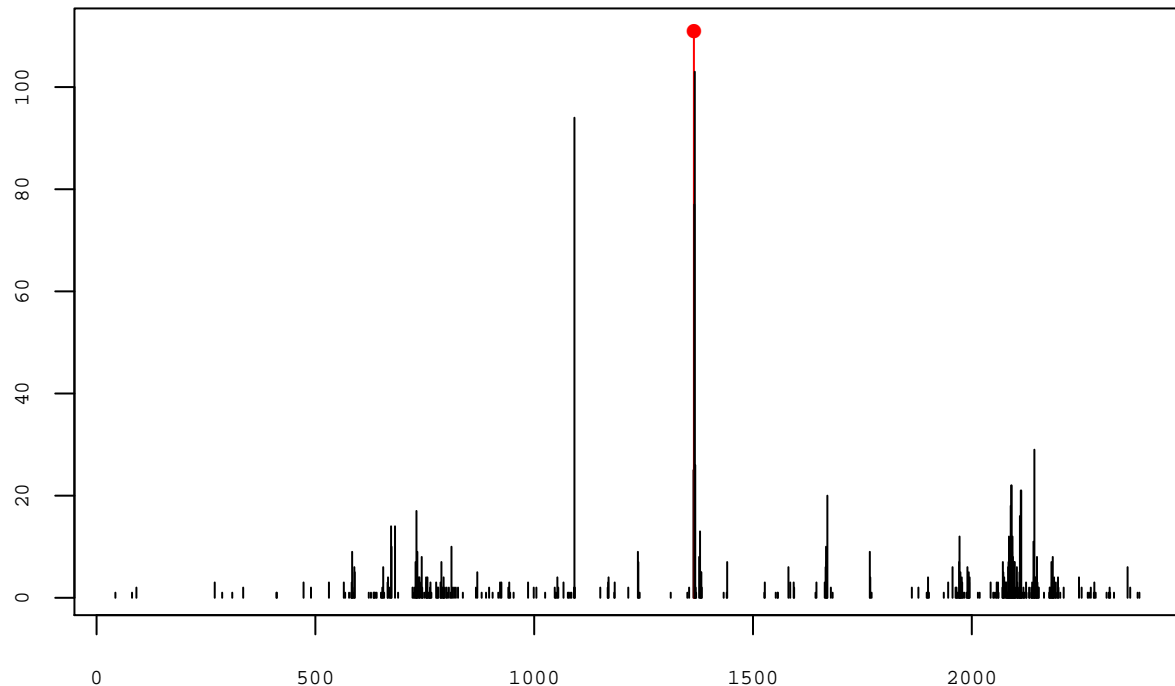


PggD6724\_c18\_g1\_i7

5' CATTGTTTCGCGAACAGCTCGTGTT-CGGCTCGT '3  
||||| ||||| ||||| |||||  
3' ACAAGCACTTGTCTGAGCACAAAC '5

Fragment Abundance



Transcript position

Cleavage site: 1365 Tag abundance: 111 Weighted abundance: 33.033 Category: 0  
sRNA abundance: 13 Alignment score: 3 MFE ratio: 0.833 p-value: 0.027