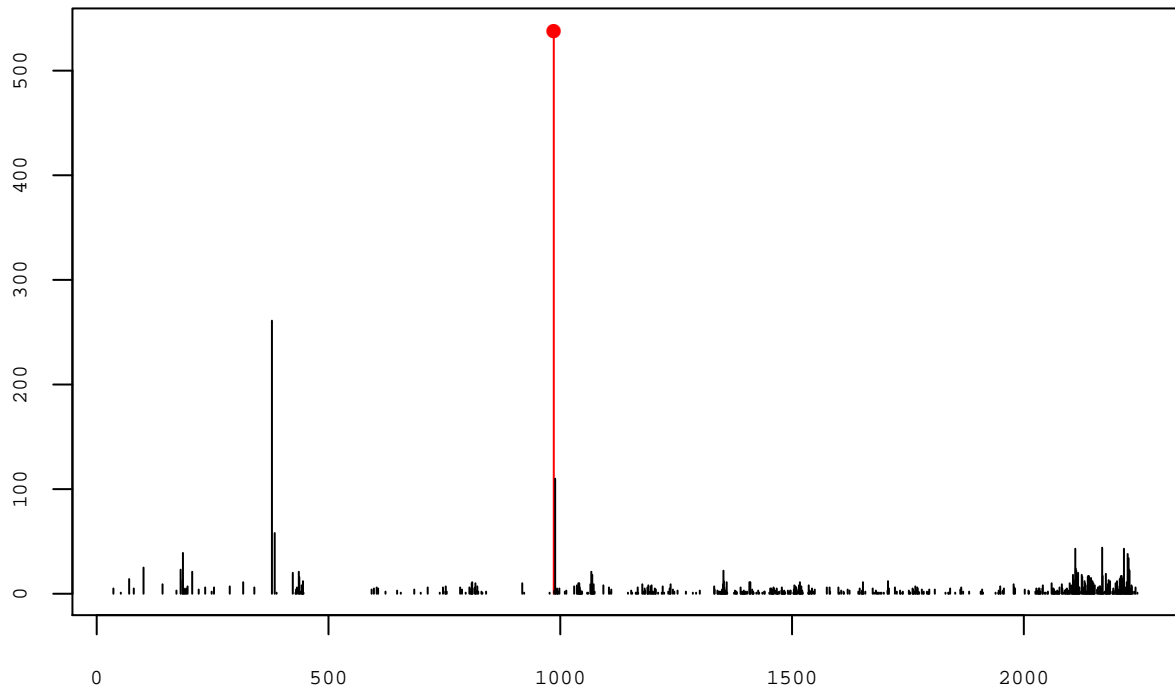


PggD5898\_c0\_g1\_i3

5' GCCCATA-GGATATTGGCGCGGCTCAATCATCA '3  
| | | | | | | | | | | | | | | | | | | | | |  
3' TCACTATAACCGCGCCGAGTT '5

Fragment Abundance



Cleavage site: 986 Tag abundance: 538 Weighted abundance: 269 Category: 0  
sRNA abundance: 20 Alignment score: 2 MFE ratio: 0.886 p-value: 0.002