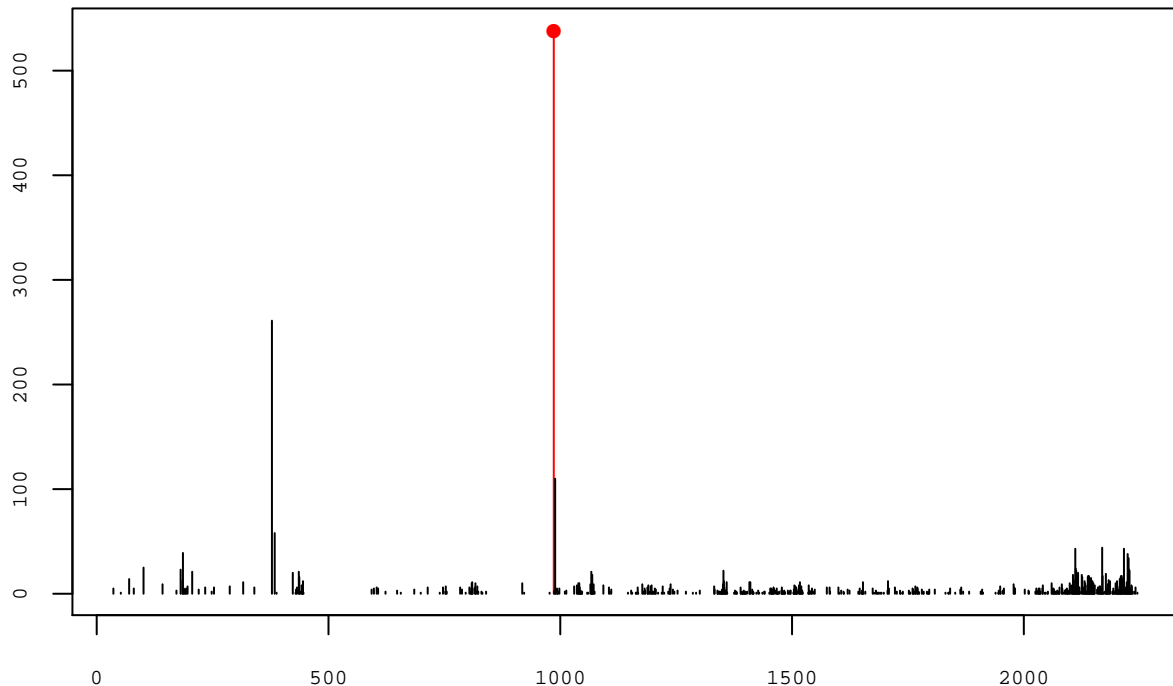


PggD5898\_c0\_g1\_i3

5' GCCCATAGGATATTGGCGCGGCTCAATCATCA '3

|||||||o|||||||  
3' CGCTATAACCGTGCCGAGTT '5

Fragment Abundance



Transcript position

Cleavage site: 986 Tag abundance: 538 Weighted abundance: 269 Category: 0  
sRNA abundance: 83 Alignment score: 3 MFE ratio: 0.878 p-value: 0.002