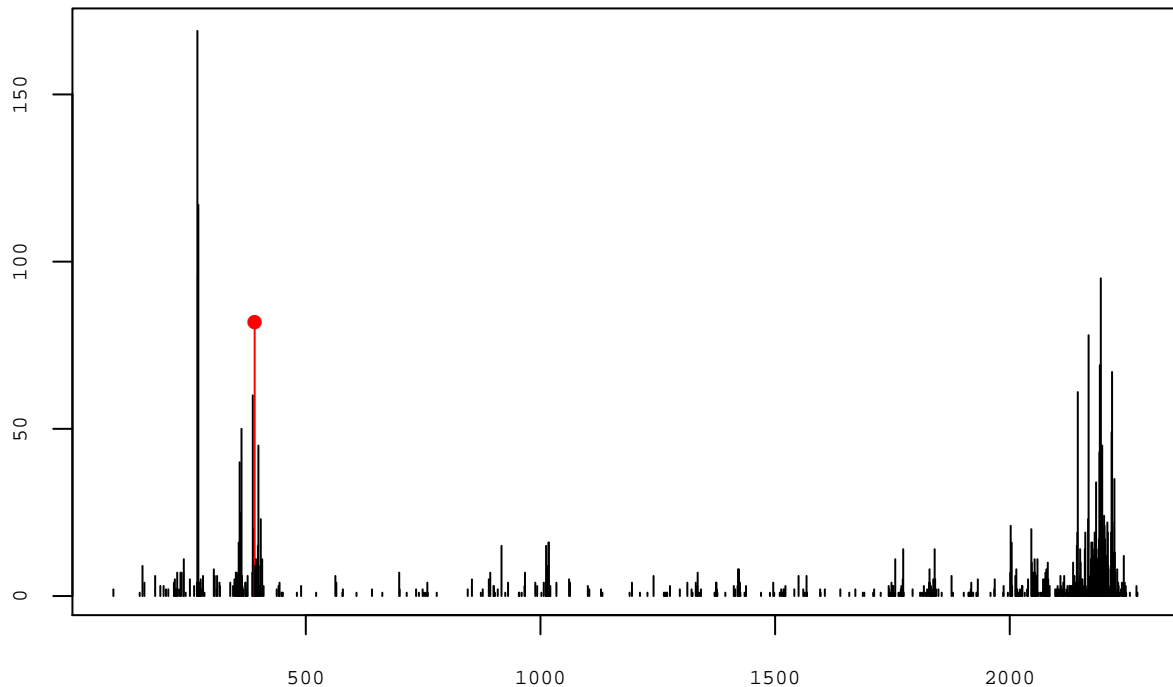


PggD1315_c0_g1_i1

5' TGCCACTCTGAAATAGCTCCGAAATACACATT '3
||||| |||||o| |||||
3' TGAGA-TTTATTGCGGCTTTAT '5

Fragment Abundance



Transcript position

Cleavage site: 391 Tag abundance: 82 Weighted abundance: 82 Category: 2
sRNA abundance: 5 Alignment score: 4 MFE ratio: 0.704 p-value: 0.033