

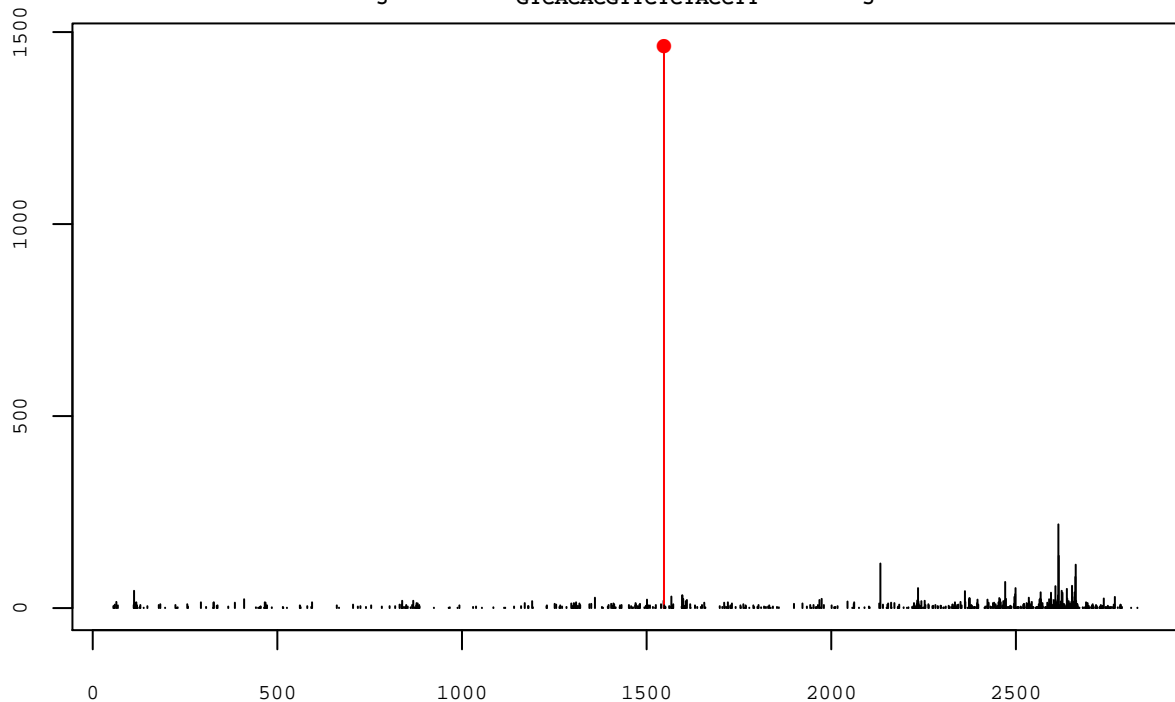
PggD4130_c0_g1_i1

5' GTTCCCTCCAGTA-GCAAGAGATGGACAGATGG '3

|||| |

3' GTCACACGTTCTCTACCTT '5

Fragment Abundance



Transcript position

Cleavage site: 1547 Tag abundance: 1464 Weighted abundance: 732 Category: 0

sRNA abundance: 8 Alignment score: 3 MFE ratio: 0.728 p-value: 0.001