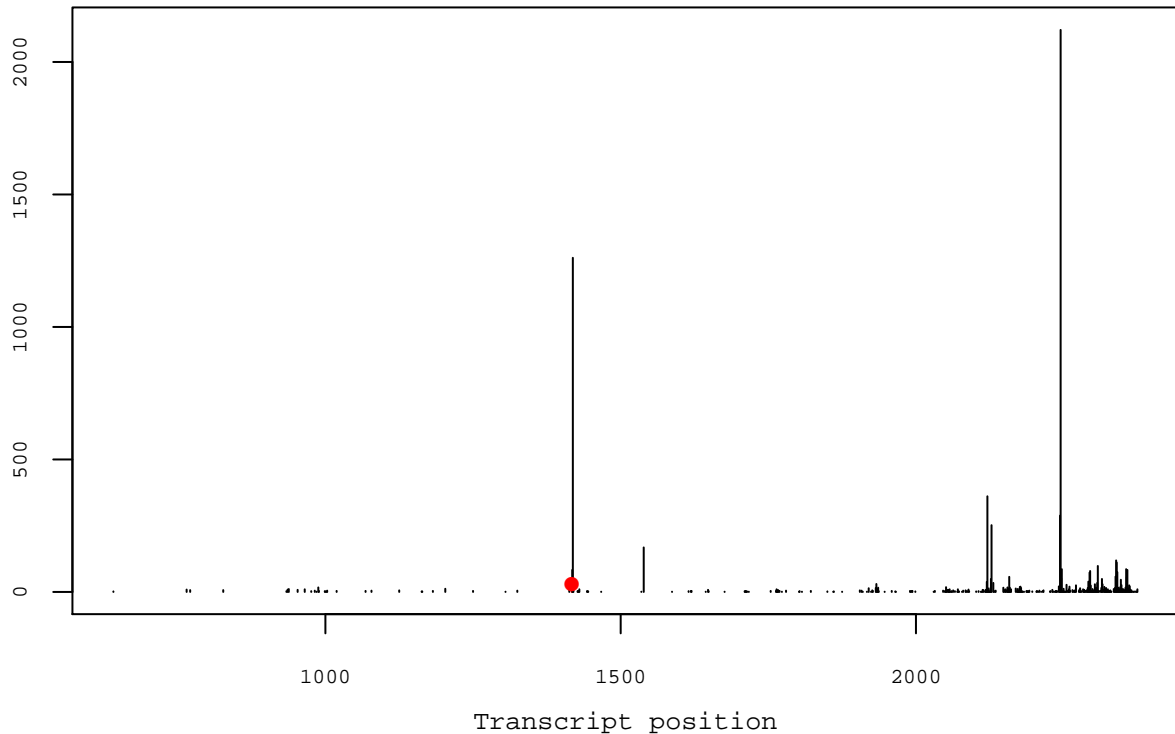


PggD7409_c15_g1_i2

5' GTGAAGATGGAGCTCCCTTCACTCCAATATCC 3'
| | | | | | | | | | | | | | | | | |
3' TCCCTCGAGGGAAGTCAGGT 5'

Fragment Abundance



Cleavage site: 1417 Tag abundance: 30 Weighted abundance: 10 Category: 2
sRNA abundance: 20 Alignment score: 3 MFE ratio: 0.756 p-value: 0.048