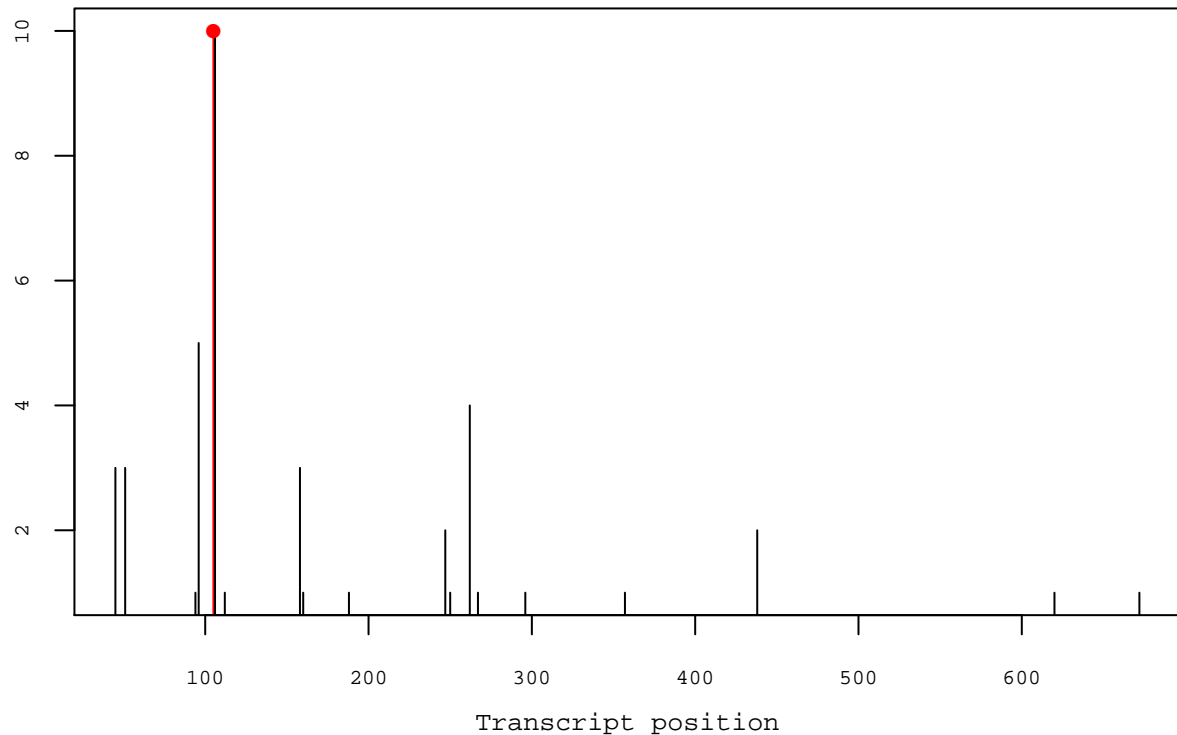


PggD1687_c147_g1_i6

5' AAACACTCCTAACAGCTCTTAAATAGCAATGG '3
|||||
3' TGAGGATTGTCGAGAATTTATC '5

Fragment Abundance



Cleavage site: 105 Tag abundance: 10 Weighted abundance: 0.909 Category: 1
sRNA abundance: 8 Alignment score: 0 MFE ratio: 1 p-value: 0.026