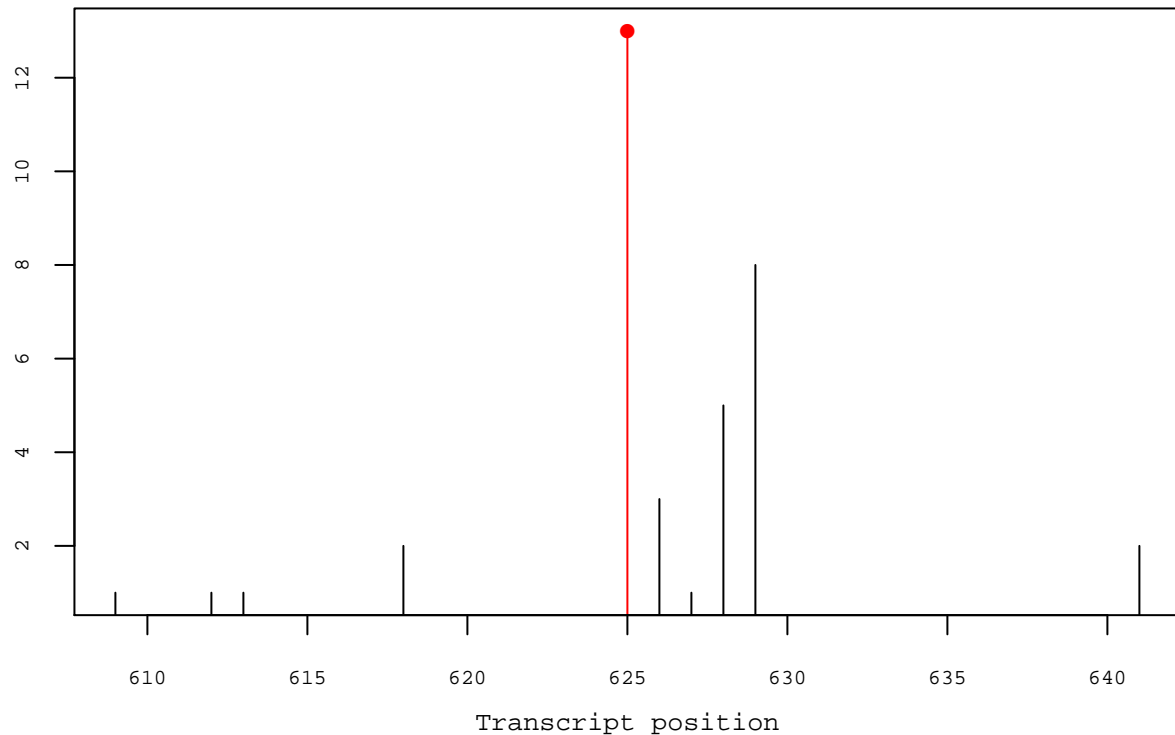


PggD1460_c118_g1_i1

5' CAGACTCCTGACGTAGCTCTACGTGCGGACCA '3
|||||
3' TGAGGACTGCATCGAGATGTACG '5

Fragment Abundance



Cleavage site: 625 Tag abundance: 13 Weighted abundance: 6.5 Category: 0
sRNA abundance: 8 Alignment score: 1 MFE ratio: 0.991 p-value: 0.003