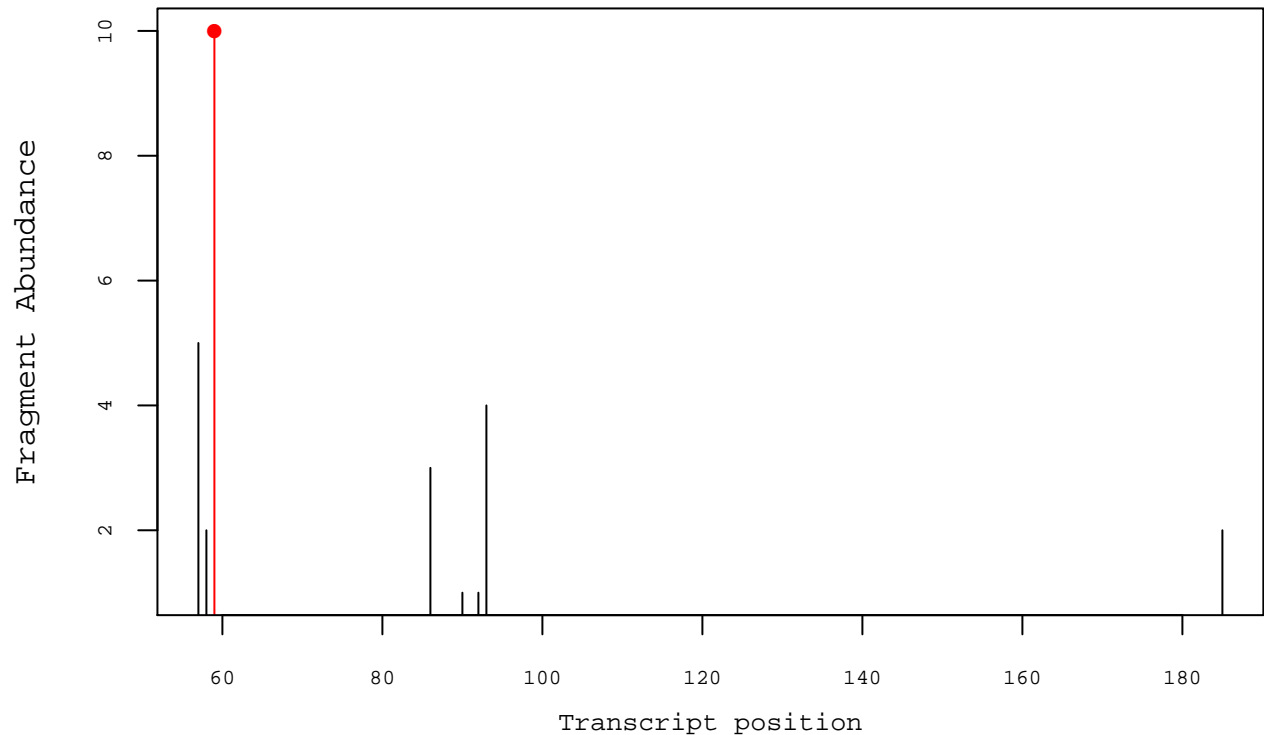


PggD5826_c3_g2_i2

5' TTCGTGATCTATGGTCAGGATCGGACACCGCA '3
 ||||| |○| ||||| |||||
 3' GCACTAGATGCCAGTCC TAGCCTA '5



Cleavage site: 59	Tag abundance: 10	Weighted abundance: 2.5	Category: 0
sRNA abundance: 111	Alignment score: 1.5	MFE ratio: 0.951	p-value: 0.003