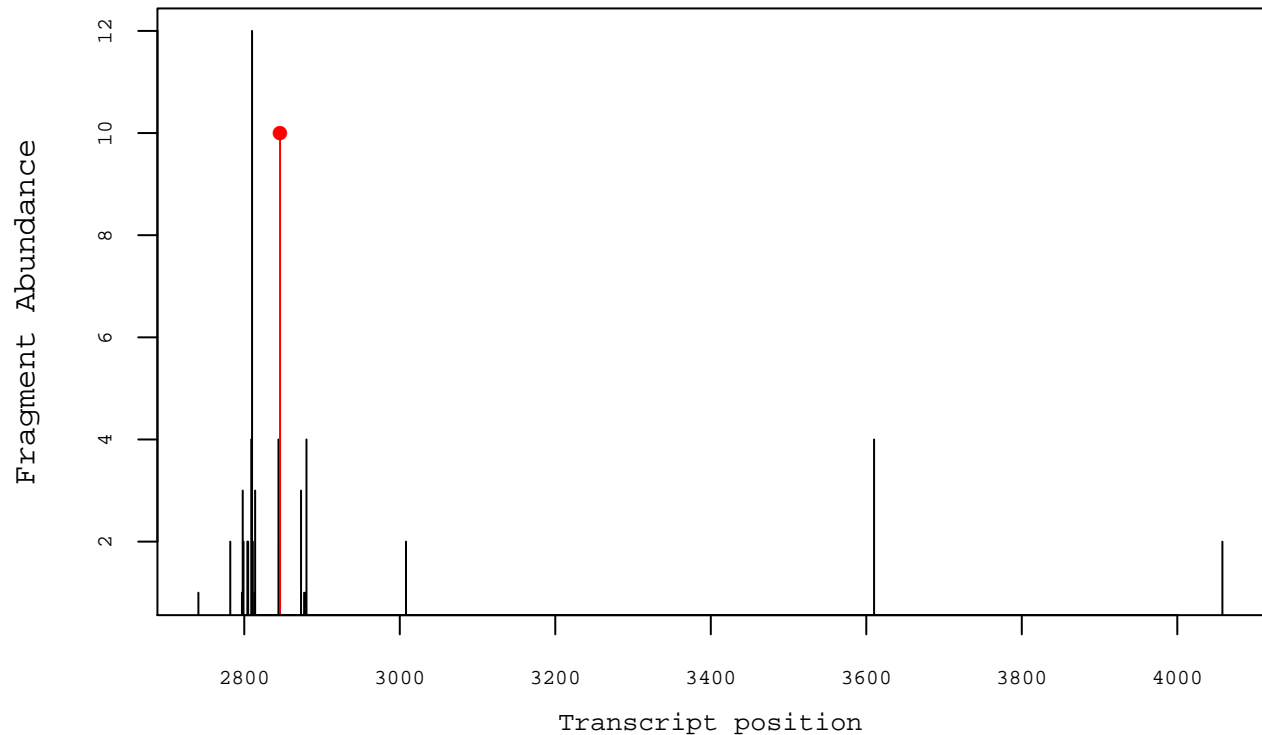


PggD7801\_c34\_g1\_i2

5' ACTGTGATCTACGGCCAGGATCGGACACCGCA '3

○ ||||| ○ |||||

3' GCCTAGATGTCAGTCCTAGCCTG 5'



Cleavage site: 2846	Tag abundance: 10	Weighted abundance: 2.5	Category: 2
sRNA abundance: 1891	Alignment score: 3	MFE ratio: 0.856	p-value: 0.005