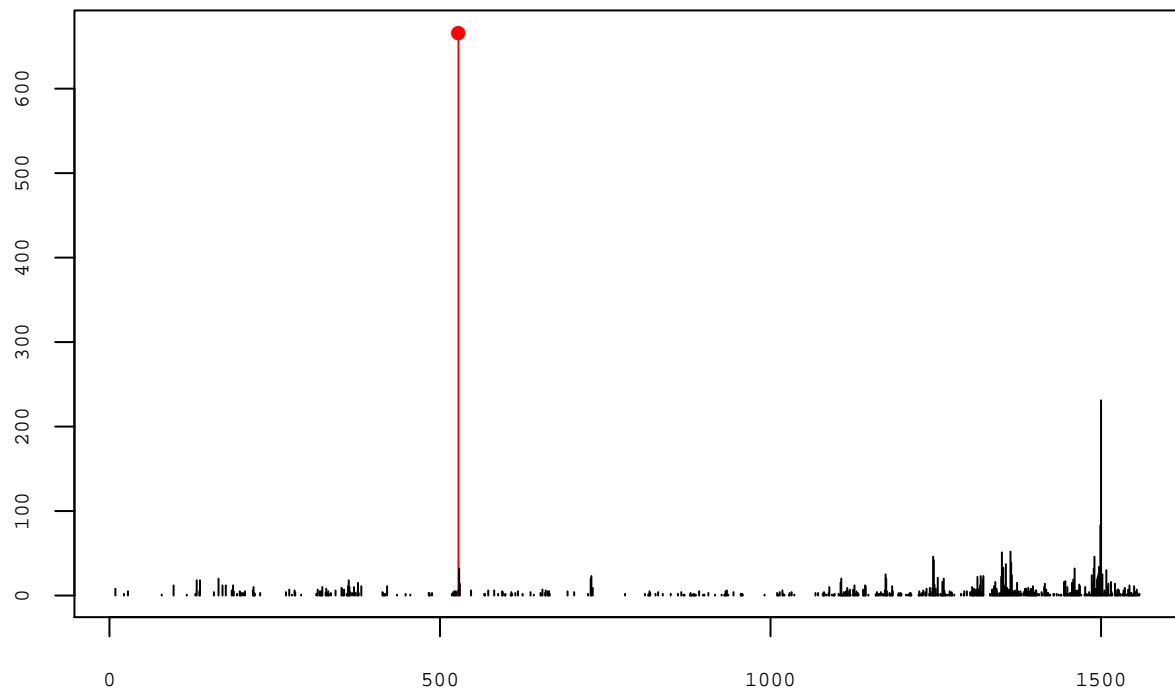


PggD12227_c16_g1_i1

5' GGAGAAAAAGCTCCATTCAATCCAAAGCGCGC '3
 ||||| |||||
3' CTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Transcript position

Cleavage site: 528	Tag abundance: 666	Weighted abundance: 666	Category: 0
sRNA abundance: 738	Alignment score: 3	MFE ratio: 0.738	p-value: 0.015