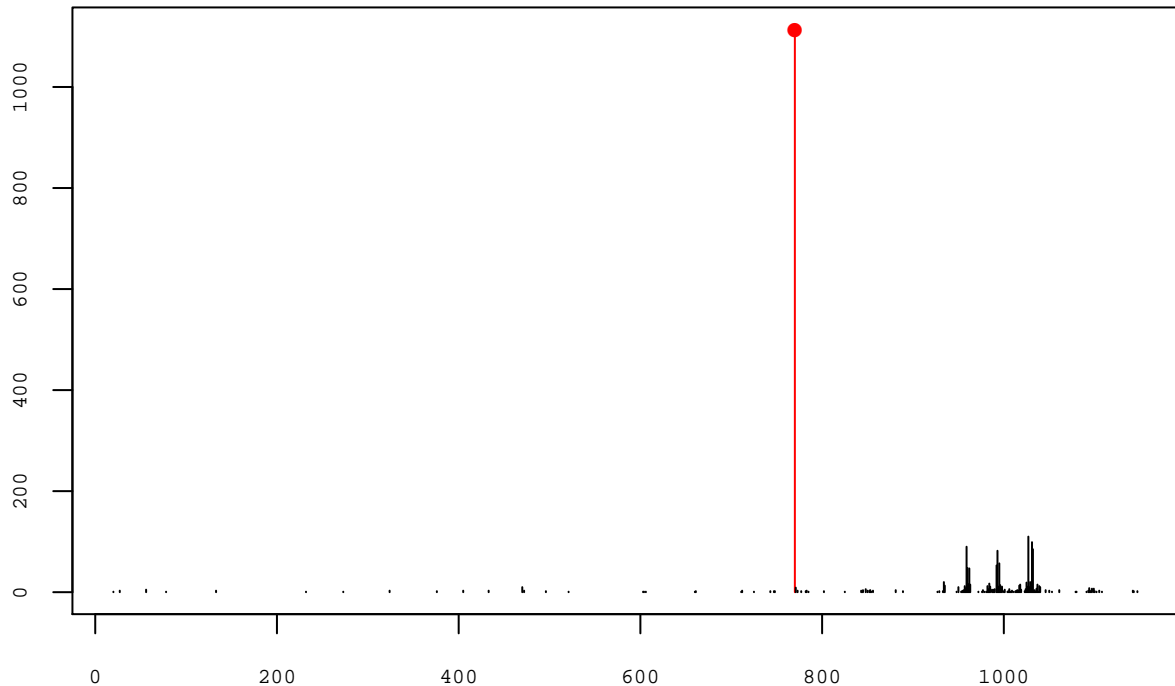


PggD1928\_c0\_g1\_i2

5' AAGGCATTACGTGCCCTGCTTCTCCACAGTGG '3  
o |||||  
3' CGTGCACGGGACGAAGAGGT '5

Fragment Abundance



Transcript position

Cleavage site: 770	Tag abundance: 1113	Weighted abundance: 556.5	Category: 0
sRNA abundance: 74	Alignment score: 1.5	MFE ratio: 0.899	p-value: 0.006