

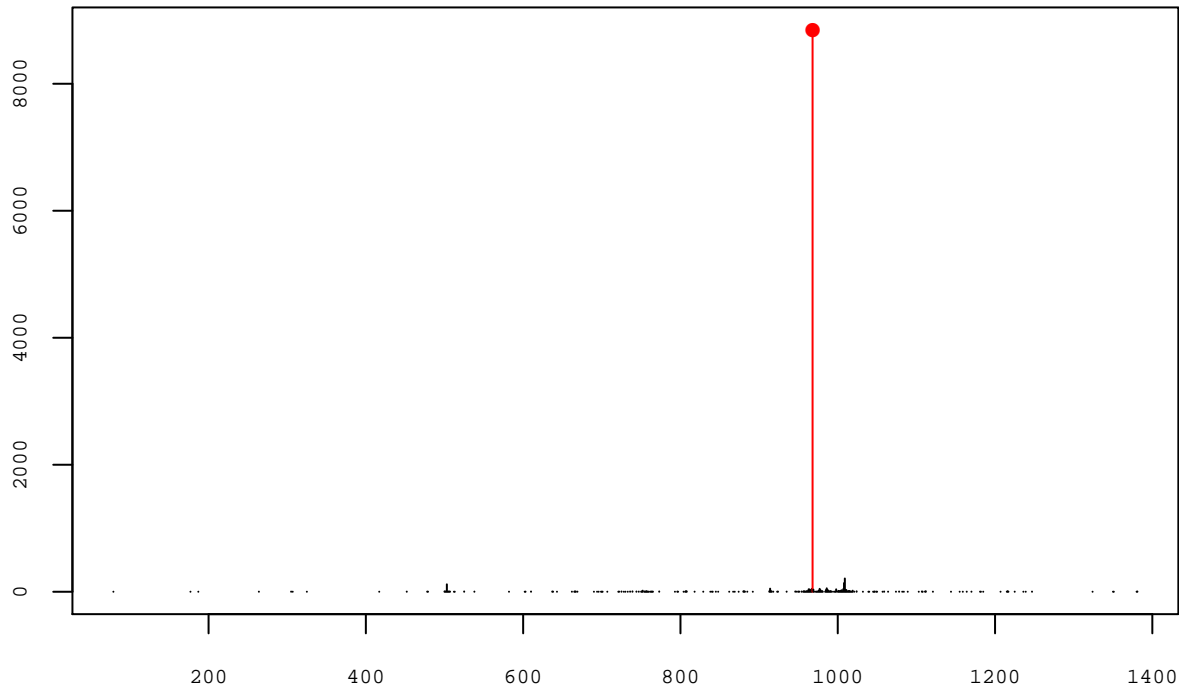
PggD14469_c62_g1_i2

5' TTCCCCTTGCTATCCCTCCTGAGCTGAGCTAG '3

o |||||

3' CGCGATAGGGAGGACTCGAA '5

Fragment Abundance



Transcript position

Cleavage site: 968	Tag abundance: 8848	Weighted abundance: 8848	Category: 0
sRNA abundance: 10	Alignment score: 2.5	MFE ratio: 0.902	p-value: 0.001