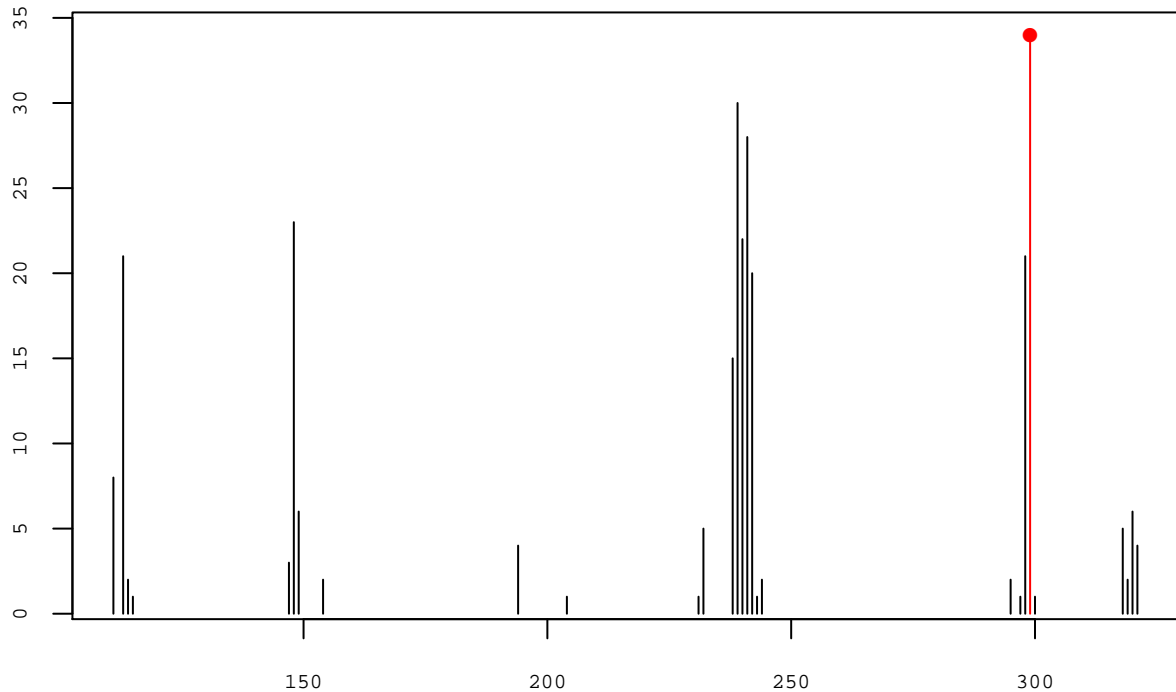


PggD1687\_c409\_g1\_i1

5' CAGTTTCAGG-GTTAAATATGTAAATATATAAG '3

3' AAAGTCCACAATTTATACATTTAT



Transcript position

Cleavage site: 299

sRNA abundance: 5

Tag abundance: 34

Alignment score: 1

Weighted abundance: 0.454

MFE ratio: 0.83

Category: 0

p-value: 0.015