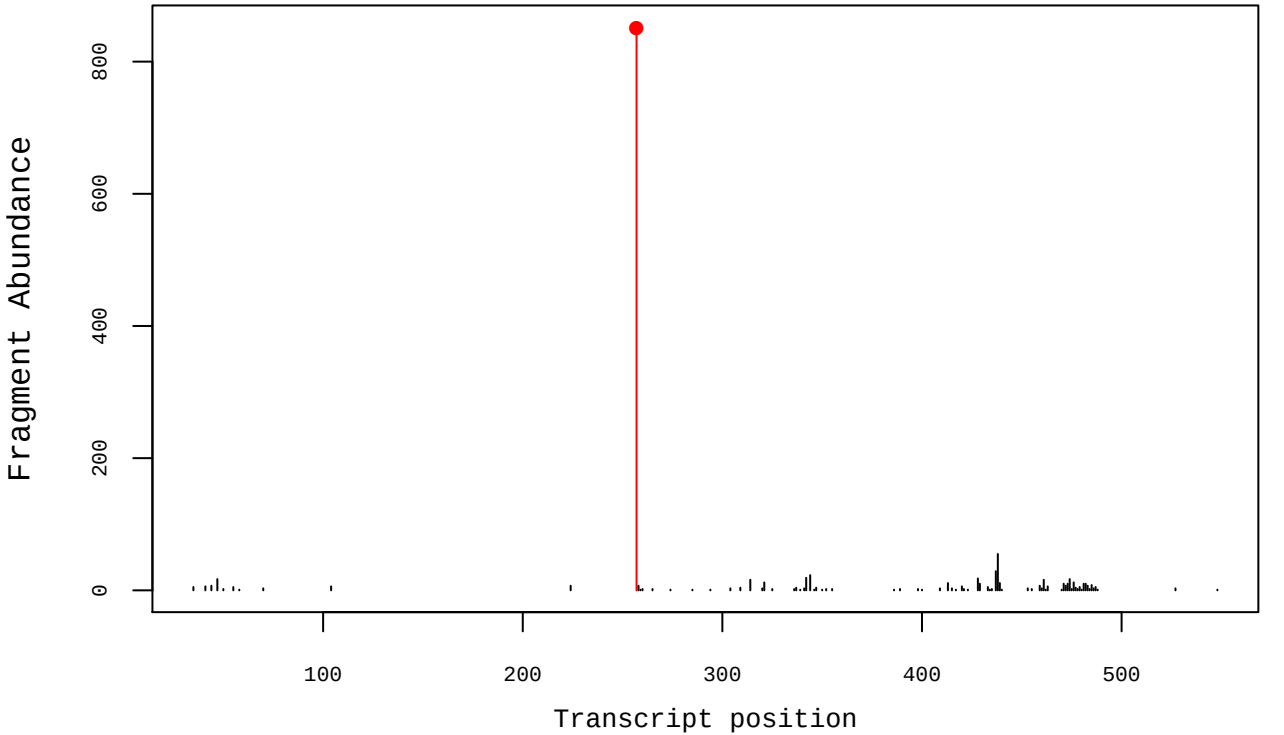
PggD10737_c261_g1_i1

```
5' GAGAGATCACGTGCCCTGCTTCTCCACATCCG '3
          ||||||||||||||||||
3'        CGTGCACGGGACGAAGAGGT       '5
```



Cleavage site: 257 Tag abundance: 851 Weighted abundance: 425.5 Category: 0
sRNA abundance: 45 Alignment score: 1 MFE ratio: 0.932 p-value: 0.009