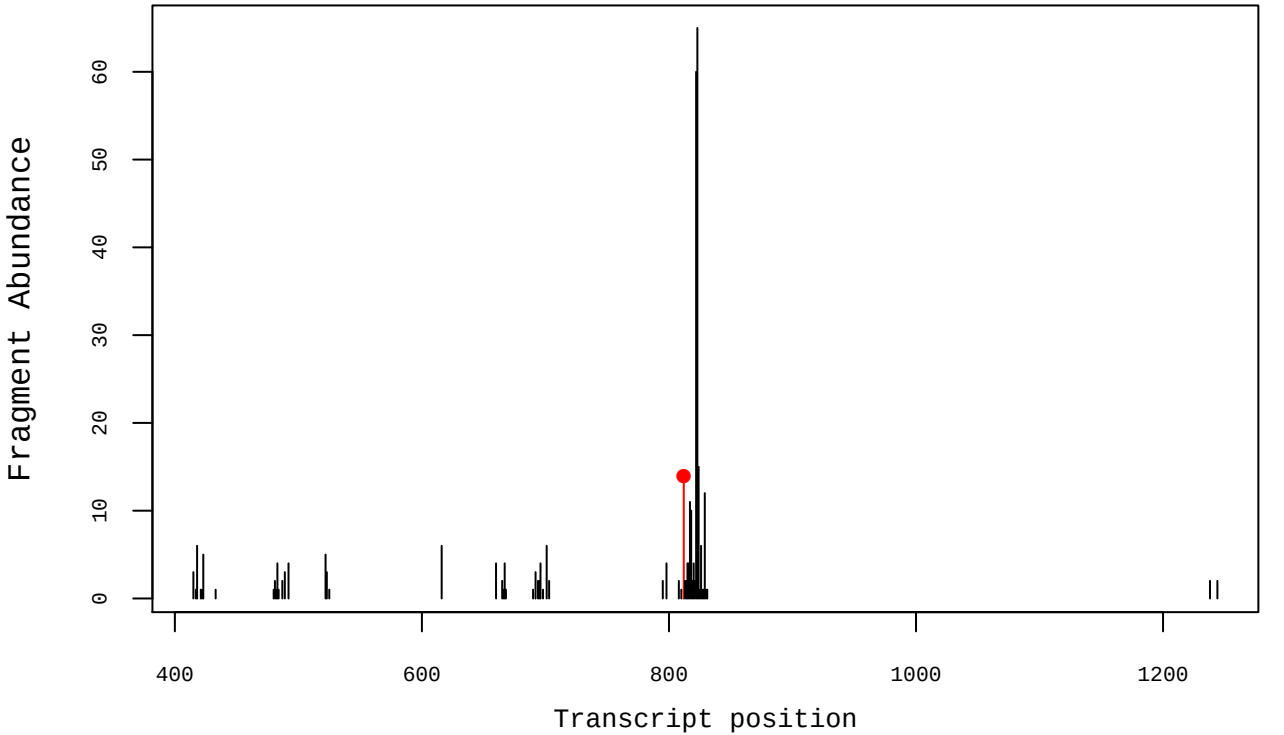
PggD5748_c2_g1_i2

```
5' TCATTGGCTACGCACGTTACTCTACGTGCGTA '3
     ||||| |||||||||| |||||||
3'   TAACCCATGCGTGCAAGGAGATGC       '5
```



Cleavage site: 812 Tag abundance: 14 Weighted abundance: 0.737 Category: 2
sRNA abundance: 10 Alignment score: 3 MFE ratio: 0.715 p-value: 0.036