

```
> # Find palindromes in zikv
```

```
> findPalindromes(zikv)
```

```
Views on a 1000-letter DNString subject
```

```
subject:
```

```
AGTTGTTGATCTGTGTGAGTCAGACTGCGACAGT...GGCATACAGTATCAGGTGCATTGGA  
GTCAGCAAT
```

```
views:
```

```
start end width
```

```
[1]  18 26   9 [AGTCAGACT]  
[2] 137 144  8 [ATCCGGAT]  
[3] 143 151  9 [ATTGTCAAT]  
[4] 363 370  8 [AAGATCTT]  
[5] 464 471  8 [GCCATGGC]  
[6] 503 510  8 [TACATGTA]  
[7] 599 606  8 [CACATGTG]  
[8] 835 844 10 [CTTGATCAAG]  
[9] 866 876 11 [AACCCCGGGTT]  
[10] 915 925 11 [GCTCGACGAGC]  
[11] 970 978  9 [ATACAGTAT]
```

*11 palindromic regions in the first 1000 bases of the Zika virus.*