

```
> library(Biostrings)
> # Load packages
> library(Biostrings)
>
> # Check the alphabet of the zikaVirus
> alphabet(zikaVirus)
[1] "A" "C" "G" "T" "M" "R" "W" "S" "Y" "K" "V" "H" "D" "B" "N" "-" "+" "."
>
> # Check the alphabetFrequency of the zikaVirus
> alphabetFrequency(zikaVirus)
  A  C  G  T M R W S Y K V H D B N - + .
[1,] 2991 2359 3139 2305 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
>
> # Check alphabet of the zikaVirus using baseOnly = TRUE
> alphabet(zikaVirus, baseOnly = TRUE)
[1] "A" "C" "G" "T"
```