

# BSgenome.Celegans.UCSC.ce11

July 17, 2024

---

BSgenome.Celegans.UCSC.ce11

*Full genome sequences for Caenorhabditis elegans (UCSC version ce11)*

---

## Description

Full genome sequences for *Caenorhabditis elegans* (Worm) as provided by UCSC (ce11, Feb. 2013) and stored in Biostrings objects.

## Note

This BSgenome data package was made from the following source data files:

ce11.2bit from <http://hgdownload.soe.ucsc.edu/goldenPath/ce11/bigZips/>

See [?BSgenomeForge](#) and the BSgenomeForge vignette (`vignette("BSgenomeForge")`) in the **BSgenome** software package for how to make a BSgenome data package.

## Author(s)

The Bioconductor Dev Team

## See Also

- [BSgenome](#) objects and the [available.genomes](#) function in the **BSgenome** software package.
- [DNASTring](#) objects in the **Biostrings** package.
- The BSgenomeForge vignette (`vignette("BSgenomeForge")`) in the **BSgenome** software package for how to make a BSgenome data package.

**Examples**

```
BSgenome.Celegans.UCSC.ce11
genome <- BSgenome.Celegans.UCSC.ce11
head(seqlengths(genome))
genome$chrI # same as genome[["chrI"]]

## -----
## Genome-wide motif searching
## -----
## See the GenomeSearching vignette in the BSgenome software
## package for some examples of genome-wide motif searching using
## Biostrings and the BSgenome data packages:
if (interactive())
  vignette("GenomeSearching", package="BSgenome")
```

# Index

\* **data**

BSgenome.Celegans.UCSC.ce11, [1](#)

\* **package**

BSgenome.Celegans.UCSC.ce11, [1](#)

available.genomes, [1](#)

BSgenome, [1](#)

BSgenome.Celegans.UCSC.ce11, [1](#)

BSgenome.Celegans.UCSC.ce11-package  
(BSgenome.Celegans.UCSC.ce11),  
[1](#)

BSgenomeForge, [1](#)

Celegans (BSgenome.Celegans.UCSC.ce11),  
[1](#)

DNAStrng, [1](#)