

# BSgenome.Ggallus.UCSC.galGal5

June 3, 2026

---

BSgenome.Ggallus.UCSC.galGal5

*Full genome sequences for Gallus gallus (UCSC version galGal5)*

---

## Description

Full genome sequences for Gallus gallus (Chicken) as provided by UCSC (galGal5, Dec. 2015) and stored in Biostrings objects.

## Note

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

galGal5.2bit from <http://hgdownload.cse.ucsc.edu/goldenPath/galGal5/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.Ggallus.UCSC.galGal5
genome <- BSgenome.Ggallus.UCSC.galGal5
head(seqlengths(genome))
genome$chr1 # same as genome[["chr1"]]
```

```
## -----
## 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.Ggallus.UCSC.galGal5, [1](#)

**\* package**

BSgenome.Ggallus.UCSC.galGal5, [1](#)

available.genomes, [1](#)

BSgenome, [1](#)

BSgenome.Ggallus.UCSC.galGal5, [1](#)

BSgenome.Ggallus.UCSC.galGal5-package  
(BSgenome.Ggallus.UCSC.galGal5),  
[1](#)

BSgenomeForge, [1](#)

DNAStrng, [1](#)

Ggallus

(BSgenome.Ggallus.UCSC.galGal5),  
[1](#)