

# BSgenome.Hsapiens.1000genomes.hs37d5

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BSgenome.Hsapiens.1000genomes.hs37d5

*1000genomes Reference Genome Sequence (hs37d5)*

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## Description

Full 1000genomes Phase2 Reference Genome Sequence (hs37d5), based on NCBI GRCh37.

## Note

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

[ftp://ftp.1000genomes.ebi.ac.uk/vol1/ftp/technical/reference/phase2\\_reference\\_assembly\\_sequence/hs37d5.fa.gz](ftp://ftp.1000genomes.ebi.ac.uk/vol1/ftp/technical/reference/phase2_reference_assembly_sequence/hs37d5.fa.gz)

The genome is composed of:

- Integrated reference sequence from the GRCh37 primary assembly (chromosomal plus unlocalized and unplaced contigs)
- The rCRS mitochondrial sequence (AC:NC\_012920)
- Human herpesvirus 4 type 1 (AC:NC\_007605)
- Concatenated decoy sequences (hs37d5cs.fa.gz)

For details, please see [ftp://ftp.1000genomes.ebi.ac.uk/vol1/ftp/technical/reference/phase2\\_reference\\_assembly\\_sequence/README\\_human\\_reference\\_20110707](ftp://ftp.1000genomes.ebi.ac.uk/vol1/ftp/technical/reference/phase2_reference_assembly_sequence/README_human_reference_20110707).

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## 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.Hsapiens.1000genomes.hs37d5  
genome <- BSgenome.Hsapiens.1000genomes.hs37d5  
seqlengths(genome)
```

# Index

## \* **data**

BSgenome.Hsapiens.1000genomes.hs37d5,

[1](#)

## \* **package**

BSgenome.Hsapiens.1000genomes.hs37d5,

[1](#)

available.genomes, [1](#)

BSgenome, [1](#)

BSgenome.Hsapiens.1000genomes.hs37d5,

[1](#)

BSgenome.Hsapiens.1000genomes.hs37d5-package

(BSgenome.Hsapiens.1000genomes.hs37d5),

[1](#)

DNASTring, [1](#)

hs37d5

(BSgenome.Hsapiens.1000genomes.hs37d5),

[1](#)