

Package ‘phastCons30way.UCSC.hg38’

December 10, 2025

Type Package

Title phastCons30way.UCSC.hg38 AnnotationHub Resource Metadata

Version 3.13.1

Encoding UTF-8

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Description Store phastCons30way.UCSC.hg38 AnnotationHub Resource Metadata.

VignetteBuilder knitr

License Artistic-2.0

Depends GenomicScores, AnnotationHub

Suggests knitr, rmarkdown, BiocStyle

biocViews AnnotationHub, Genetics, GenomicScores, Homo_sapiens

organism Homo sapiens

provider UCSC

provider_version 03Nov2017

git_url <https://git.bioconductor.org/packages/phastCons30way.UCSC.hg38>

git_branch devel

git_last_commit e50210a

git_last_commit_date 2025-06-09

Repository Bioconductor 3.23

Date/Publication 2025-12-10

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phastCons30way.UCSC.hg38

phastCons30way genomic scores for Homo sapiens (hg38)

Description

This annotation package stores genomic scores for Homo sapiens, genome version hg38.

Note

This data package was made from resources at UCSC based on version 03Nov2017.

Author(s)

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See Also

[GenomicScores](#)

Examples

```
## Not run:
library(GenomicScores)

gsco <- getGScores("phastCons30way.UCSC.hg38")
name(gsco)
type(gsco)
provider(gsco)
providerVersion(gsco)
organism(gsco)
referenceGenome(gsco)
seqlevelsStyle(gsco)
seqinfo(gsco)
head(seqnames(gsco))
head(seqlengths(gsco))
citation(gsco)

## End(Not run)
```

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