

phastCons30way.UCSC.hg38

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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.

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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))
```

2

phastCons30way.UCSC.hg38

`citation(gsc)`

`## End(Not run)`

Index

* **data**

phastCons30way.UCSC.hg38, [1](#)

* **package**

phastCons30way.UCSC.hg38, [1](#)

GenomicScores, [1](#)

phastCons30way.UCSC.hg38, [1](#)

phastCons30way.UCSC.hg38-package
(phastCons30way.UCSC.hg38), [1](#)