

# phastCons30way.UCSC.hg38

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*phastCons30way genomic scores for Homo sapiens (hg38)*

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## 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))
```

```
citation(gscs)
```

```
## End(Not run)
```

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