

# phastCons30way.UCSC.hg38

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phastCons30way.UCSC.hg38

*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))
```

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citation(gsc)

## End(Not run)

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