

# cadd.v1.6.hg19

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*CADD Pathogenicity Scores AnnotationHub Resource Metadata for Homo sapiens (hg19)*

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## Description

This annotation package stores CADD v1.6 pathogenicity scores AnnotationHub resource metadata for Homo sapiens, genome version hg19.

## Note

This data package was made from resources at University of Washington based on version v1.6.

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## See Also

[GenomicScores](#)

## Examples

```
library(GenomicScores)

gsco <- getGScores("cadd.v1.6.hg19")
name(gsco)
type(gsco)
provider(gsco)
providerVersion(gsco)
organism(gsco)
referenceGenome(gsco)
seqlevelsStyle(gsco)
seqinfo(gsco)
head(seqnames(gsco))
head(seqlengths(gsco))
citation(gsco)
```

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