

# cadd.v1.6.hg19

October 16, 2024

---

cadd.v1.6.hg19	<i>CADD Pathogenicity Scores AnnotationHub Resource Metadata for Homo sapiens (hg19)</i>
----------------	--

---

## 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.

## Author(s)

Robert Castelo <robert.castelo@upf.edu> [cre]

## 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(gsc0)`

# Index

\* **data**

cadd.v1.6.hg19, [1](#)

\* **package**

cadd.v1.6.hg19, [1](#)

cadd.v1.6.hg19, [1](#)

cadd.v1.6.hg19-package

(cadd.v1.6.hg19), [1](#)

GenomicScores, [1](#)