cadd.v1.6.hg19

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`cadd.v1.6.hg19` *CADD Pathogenicity Scores AnnotationHub Resource Metadata for Homo sapiens (hg19)*

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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