

# AlphaMissense.v2023.hg38

December 4, 2024

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*AlphaMissense Pathogenicity Scores AnnotationHub Resource Metadata for Homo sapiens (hg38)*

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## Description

This annotation package stores AlphaMissense v2023 pathogenicity scores AnnotationHub resource metadata for Homo sapiens, genome version hg38.

## Note

This data package was made from resources at Google DeepMind based on version v2023.

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## See Also

[GenomicScores](#)

## Examples

```
library(GenomicScores)

gsco <- getGScores("AlphaMissense.v2023.hg38")
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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