

Package: AshkenazimSonChr21 (via r-universe)

October 10, 2026

Type Package

Title Annotated variants on the chromosome 21, human genome 19,
Ashkenazim Trio son sample

Version 1.42.0

Date 2015-04-20

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Maintainer Who to complain to <tomasz.stokowy@k2.uib.no>

Description SonVariantsChr21 is a dataset of annotated genomic
variants coming from Complete Genomics whole genome sequencing.
Data comes from GIAB project, Ashkenazim Trio, sample HG002 run
1. Both vcf and annotated data frame are provided.

https //sites.stanford.edu/abms/content/giab-reference-materials-and-data

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LazyData TRUE

Suggests knitr, VariantAnnotation

VignetteBuilder knitr

biocViews GenomicVariation, Sequencing, WholeGenome

NeedsCompilation no

Repository Bioconductor 3.23

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RemoteUrl https://github.com/bioc/AshkenazimSonChr21

RemoteRef RELEASE_3_23

RemoteSha 439e4bb398098dd6c4c3941aaff5b5013d345f88

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SonVariantsChr21	<i>Annotated variants on the chromosome 21, human genome 19, son genome</i>
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Description

SonVariantsChr21 is a data frame consisting of annotated genomic variants coming from Complete Genomics whole genome sequencing. Data comes from GIAB project, Ashkenazim Trio, sample HG002 run 1 <https://sites.stanford.edu/abms/content/giab-reference-materials-and-data>

Usage

```
data(SonVariantsChr21)
```

Format

data frame

Value

test data

Source

<https://sites.stanford.edu/abms/content/giab-reference-materials-and-data>

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* **datasets**

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