

BSgenome.CneoformansVarGrubiiKN99.NCBI.ASM221672v1

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`BSgenome.CneoformansVarGrubiiKN99.NCBI.ASM221672v1`

*Full Genome Sequence for Cryptococcus neoformans var. grubii
KN99 (ASM221672v1)*

Description

Full genome sequences for *Cryptococcus neoformans* var. *grubii* KN99 (assembly ASM221672v1 assembly accession GCA_002216725.1).

Note

This BSgenome data package was made from the following source data files:

-- information not available --

See [?BSgenomeForge](#) and the `BSgenomeForge` vignette (`vignette("BSgenomeForge")`) in the **BSgenome** software package for how to make a BSgenome data package.

Author(s)

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See Also

- `BSgenome` objects and the `available.genomes` function in the **BSgenome** software package.
- `DNAString` objects in the **Biostrings** package.
- The `BSgenomeForge` vignette (`vignette("BSgenomeForge")`) in the **BSgenome** software package for how to make a BSgenome data package.

Examples

```
BSgenome.CneoformansVarGrubiiKN99.NCBI.ASM221672v1
genome <- BSgenome.CneoformansVarGrubiiKN99.NCBI.ASM221672v1
head(seqlengths(genome))
genome$CP022321.1 # same as genome[["CP022321.1"]]

## -----
## Genome-wide motif searching
## -----
## See the GenomeSearching vignette in the BSgenome software
## package for some examples of genome-wide motif searching using
## Biostrings and the BSgenome data packages:
if (interactive())
  vignette("GenomeSearching", package="BSgenome")
```

Index

```
* data
  BSgenome.CneoformansVarGrubiiKN99.NCBI.ASM221672v1,
    1
* package
  BSgenome.CneoformansVarGrubiiKN99.NCBI.ASM221672v1,
    1
available.genomes, I
BSgenome, I
BSgenome.CneoformansVarGrubiiKN99.NCBI.ASM221672v1,
  1
BSgenome.CneoformansVarGrubiiKN99.NCBI.ASM221672v1-package
  (BSgenome.CneoformansVarGrubiiKN99.NCBI.ASM221672v1),
    1
BSgenomeForge, I
CneoformansVarGrubiiKN99
  (BSgenome.CneoformansVarGrubiiKN99.NCBI.ASM221672v1),
    1
DNAString, I
```