# Package 'alabaster.vcf'

### November 14, 2024

Title Save and Load Variant Data to/from File
Version 1.6.0
<b>Date</b> 2024-01-02
Description Save variant calling SummarizedExperiment to file and load them back as VCF objects.  This is a more portable alternative to serialization of such objects into RDS files.  Each artifact is associated with metadata for further interpretation; downstream applications can enrich this metadata with context-specific properties.
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<b>Depends</b> alabaster.base, VariantAnnotation
Imports methods, S4Vectors, alabaster.se, alabaster.string, Rsamtools
Suggests knitr, rmarkdown, BiocStyle, testthat
RoxygenNote 7.2.3
VignetteBuilder knitr
biocViews DataImport, DataRepresentation
git_url https://git.bioconductor.org/packages/alabaster.vcf
git_branch RELEASE_3_20
git_last_commit 6151a95
git_last_commit_date 2024-10-29
Repository Bioconductor 3.20
Date/Publication 2024-11-14
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readVCF

Read a VCF object from disk

#### Description

Read a VCF object from its on-disk representation.

#### Usage

```
readVCF(path, metadata, ...)
```

#### **Arguments**

path String containing a path to a directory, usually generated by the saveObject

method for VCF object.s

metadata Named list of metadata for this object, see readObjectFile for details.

... Further arguments passed to internal altReadObject calls.

#### Value

A VCF object.

#### Author(s)

Aaron Lun

#### See Also

saveObject, VCF-method, to save VCF objects to disk.

#### **Examples**

```
fl <- system.file("extdata", "structural.vcf", package="VariantAnnotation")
vcf <- readVcf(fl)

tmp <- tempfile()
saveObject(vcf, tmp)
readObject(tmp)</pre>
```

saveObject,VCF-method Save a VCF object to disk

#### **Description**

Save a VCF object to its on-disk representation, namely a VCF file with the same contents.

#### Usage

```
## S4 method for signature 'VCF'
saveObject(x, path, ...)
```

#### **Arguments**

Any instance of a VCF class or one of its subclasses.
 String containing the path to a directory in which to save x.
 Further arguments to pass to specific methods.

#### Value

x is saved to file inside path, and NULL is returned.

#### Author(s)

Aaron Lun

#### See Also

readVCF, to read a VCF object back to the R session.

#### **Examples**

```
fl <- system.file("extdata", "structural.vcf", package="VariantAnnotation")
vcf <- readVcf(fl)

tmp <- tempfile()
saveObject(vcf, tmp)</pre>
```

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