Package 'alabaster.vcf'

April 21, 2025

Title Save and Load Variant Data to/from File

Version 1.9.0

Date 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.

License MIT + file LICENSE

Depends 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 devel

git_last_commit 5efb770

git_last_commit_date 2025-04-15

Repository Bioconductor 3.22

Date/Publication 2025-04-21

Author Aaron Lun [aut, cre]

Maintainer Aaron Lun <infinite.monkeys.with.keyboards@gmail.com>

Contents

readVCF		•	•			•	•	•			•		•	•			•			•			2
saveObject,VCF-method	 •	•	•	• •	 •	•	•	•	•	•	•	•	•		•			•		•	•	•	3

4

Index

readVCF

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)</pre>
```

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

Х	Any instance of a VCF class or one of its subclasses.
path	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)</pre>
```

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

Index

altReadObject, 2
loadVCF(readVCF), 2
loadVCFHeader(readVCF), 2
readObjectFile, 2
readVCF, 2, 3

saveObject, 2
saveObject,VCF-method, 3
stageObject,VCF-method
 (saveObject,VCF-method), 3
stageObject,VCFHeader-method
 (saveObject,VCF-method), 3

VCF, 2, 3