## Package 'alabaster.spatial'

October 15, 2023

Title Save and Load Spatial 'Omics Data to/from File Description Save SpatialExperiment objects and their images into file artifacts, and load them back into memory. 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. Version 1.0.0 Date 2023-02-28 License MIT + file LICENSE Depends SpatialExperiment, alabaster.base Imports methods, utils, grDevices, S4Vectors, SummarizedExperiment, jsonlite, alabaster.sce Suggests testthat, knitr, rmarkdown, BiocStyle, magick, png, digest VignetteBuilder knitr RoxygenNote 7.2.1 biocViews DataImport, DataRepresentation git\_url https://git.bioconductor.org/packages/alabaster.spatial git\_branch RELEASE\_3\_17 git\_last\_commit 0a9683f git\_last\_commit\_date 2023-04-25 Date/Publication 2023-10-15 Author Aaron Lun [aut, cre]

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

### **R** topics documented:

	loadSpatialExperiment	2
	loadSpatialImage	3
	stageObject,SpatialExperiment-method	3
	stageSpatialImage	4
		_
Index		1

loadSpatialExperiment Load a spatial experiment

#### Description

Load a SpatialExperiment object from its constituent files in DataSetDB.

#### Usage

loadSpatialExperiment(exp.info, project)

#### Arguments

exp.info	Named list of metadata for a spatial 'omics experiment.
project	Any argument accepted by the acquisition functions, see ?acquireFile. By default, this should be a string containing the path to a staging directory.

#### Value

A SpatialExperiment object.

#### Author(s)

Aaron Lun

#### Examples

```
library(SpatialExperiment)
example(read10xVisium, echo=FALSE)
colnames(spe) <- make.unique(colnames(spe)) # forcing unique column names.</pre>
```

```
tmp <- tempfile()
dir.create(tmp)
meta <- stageObject(spe, tmp, "experiment-1")</pre>
```

```
meta$path <- "experiment-1/experiment.json"
loadSpatialExperiment(meta, tmp)</pre>
```

loadSpatialImage Load a spatial image

#### Description

Load an image as a SpatialImage or subclass thereof.

#### Usage

loadSpatialImage(img.info, project)

#### Arguments

img.info	Named list containing the metadata for this assay.
project	Any argument accepted by the acquisition functions, see ?acquireFile. By
	default, this should be a string containing the path to a staging directory.

#### Value

A SpatialImage containing the image data (or a reference to it).

#### Author(s)

Aaron Lun

#### Examples

```
example(read10xVisium, echo=FALSE)
img <- imgData(spe)$data[[1]]
tmp <- tempfile()
dir.create(tmp)
meta <- stageObject(img, tmp, "whee")
out <- loadSpatialImage(meta, tmp)</pre>
```

#### Description

Stage a SpatialExperiment object.

#### Usage

```
## S4 method for signature 'SpatialExperiment'
stageObject(x, dir, path, child = FALSE, ...)
```

#### Arguments

х	A SpatialExperiment object.
dir	String containing the path to the staging directory.
path	String containing a prefix of the relative path inside dir where x is to be saved. The actual path used to save x may include additional components, see Details.
child	Logical scalar indicating whether x is a child of a larger object.
	Further arguments to pass to specific methods.

#### Value

A named list of the same form as that returned by the stageObject method for a SingleCellExperiment, but containing additional fields for the spatial data. A directory is created at path inside dir and is populated with the contents of x.

#### Author(s)

Aaron Lun

#### Examples

```
library(SpatialExperiment)
example(read10xVisium, echo=FALSE)
colnames(spe) <- make.unique(colnames(spe)) # forcing unique column names.
tmp <- tempfile()
dir.create(tmp)</pre>
```

```
stageObject(spe, tmp, "experiment-1")
list.files(tmp, recursive=TRUE)
```

stageSpatialImage Stage images for upload to DataSetDB

#### Description

Stage images from a variety of sources in preparation for upload to DataSetDB.

#### stageSpatialImage

#### Usage

```
## S4 method for signature 'VirtualSpatialImage'
stageObject(x, dir, path, child = FALSE, ...)
## S4 method for signature 'StoredSpatialImage'
stageObject(x, dir, path, child = FALSE, ...)
## S4 method for signature 'RemoteSpatialImage'
```

stageObject(x, dir, path, child = FALSE, ...)

#### Arguments

х	A SpatialImage object.
dir	String containing the path to the staging directory.
path	String containing a prefix of the relative path inside dir where x is to be saved. The actual path used to save x may include additional components, see Details.
child	Logical scalar indicating whether x is a child of a larger object.
	Further arguments to pass to specific methods.

#### Details

Each of the different methods will take advantage of any existing files to avoid an actual save. For example, the RemoteSpatialImage method will download the file directly to path, while the StoredSpatialImage method will create a link or copy the file. The SpatialImage method will fall back to saving the raster directly as a PNG.

#### Value

An image file is created at file.path(dir, path), possibly after appending an appropriate file extension.

The return value should be a named list containing at least:

- \$schema, a string specifying the schema to use to validate the metadata. This may have a package attribute to specify the package where the schema lives (in its inst/schemas directory).
- path, a string containing the path to the file containing the assay contents. This should start with the input path but can be followed by any necessary file extensions.
- child, whether this is a child resource of a larger object.

Other fields can be provided and will be included in the metadata, provided that they are recognized by the specified schema.

#### Author(s)

Aaron Lun

#### stageSpatialImage

#### Examples

```
example(read10xVisium, echo=FALSE)
(img <- imgData(spe)$data[[1]])
# Doing a local run:
tmp <- tempfile()</pre>
```

```
dir.create(tmp)
stageObject(img, tmp, "whee")
```

```
# Forcing a re-save:
Y <- as(img, "LoadedSpatialImage")
stageObject(Y, tmp, "foo")
```

6

# Index

```
acquireFile, 2, 3
loadSpatialExperiment, 2
loadSpatialImage, 3
RemoteSpatialImage, 5
SpatialExperiment, 2-4
SpatialImage, 3, 5
stageObject,4
stageObject,RemoteSpatialImage-method
        (stageSpatialImage), 4
stageObject,SpatialExperiment-method,
        3
stageObject,StoredSpatialImage-method
        (stageSpatialImage), 4
stageObject,VirtualSpatialImage-method
        (stageSpatialImage), 4
stageSpatialImage, 4
StoredSpatialImage, 5
```