

# Package ‘git2rdata’

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**Title** Store and Retrieve Data.frames in a Git Repository

**Version** 0.5.2

**Description** The git2rdata package is an R package for writing and reading dataframes as plain text files. A metadata file stores important information. 1) Storing metadata allows to maintain the classes of variables. By default, git2rdata optimizes the data for file storage. The optimization is most effective on data containing factors. The optimization makes the data less human readable. The user can turn this off when they prefer a human readable format over smaller files. Details on the implementation are available in `vignette("plain_text", package = "git2rdata")`. 2) Storing metadata also allows smaller row based diffs between two consecutive commits. This is a useful feature when storing data as plain text files under version control. Details on this part of the implementation are available in `vignette("version_control", package = "git2rdata")`. Although we envisioned git2rdata with a git workflow in mind, you can use it in combination with other version control systems like subversion or mercurial. 3) git2rdata is a useful tool in a reproducible and traceable workflow. `vignette("workflow", package = "git2rdata")` gives a toy example. 4) `vignette("efficiency", package = "git2rdata")` provides some insight into the efficiency of file storage, git repository size and speed for writing and reading.

**License** GPL-3

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<code>commit</code>	<i>Re-exported Function From git2r</i>
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### Description

See `commit` in `git2r`.

### See Also

Other version\_control: `pull()`, `push()`, `recent_commit()`, `repository()`, `status()`

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<code>data_package</code>	<i>Create a Data Package for a directory of CSV files</i>
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### Description

Create a `datapackage.json` file for a directory of CSV files. The function will look for all `.csv` files in the directory and its subdirectories. It will then create a `datapackage.json` file with the metadata of each CSV file.

### Usage

```
data_package(path = ".")
```

### Arguments

`path` the directory in which to create the `datapackage.json` file.

### See Also

Other storage: `display_metadata()`, `list_data()`, `prune_meta()`, `read_vc()`, `relabel()`, `rename_variable()`, `rm_data()`, `update_metadata()`, `verify_vc()`, `write_vc()`

---

display_metadata	<i>Display metadata for a git2rdata object</i>
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---

### Description

Display metadata for a git2rdata object

### Usage

```
display_metadata(x, minimal = FALSE)
```

### Arguments

x	a git2rdata object
minimal	logical, if TRUE only a message is displayed

### See Also

Other storage: [data\\_package\(\)](#), [list\\_data\(\)](#), [prune\\_meta\(\)](#), [read\\_vc\(\)](#), [relabel\(\)](#), [rename\\_variable\(\)](#), [rm\\_data\(\)](#), [update\\_metadata\(\)](#), [verify\\_vc\(\)](#), [write\\_vc\(\)](#)

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list_data	<i>List Available Git2rdata Files Containing Data</i>
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---

### Description

The function returns the names of all valid git2rdata objects. This implies .tsv files with a matching **valid** metadata file (.yaml). **Invalid** metadata files result in a warning. The function ignores **valid** metadata files without matching raw data (.tsv).

### Usage

```
list_data(root = ".", path = ".", recursive = TRUE)
```

### Arguments

root	the root of the repository. Either a path or a git-repository
path	relative path from the root. Defaults to the root
recursive	logical. Should the listing recurse into directories?

### Value

A character vector of git2rdata object names, including their relative path.

**See Also**

Other storage: [data\\_package\(\)](#), [display\\_metadata\(\)](#), [prune\\_meta\(\)](#), [read\\_vc\(\)](#), [relabel\(\)](#), [rename\\_variable\(\)](#), [rm\\_data\(\)](#), [update\\_metadata\(\)](#), [verify\\_vc\(\)](#), [write\\_vc\(\)](#)

**Examples**

```
## on file system

# create a directory
root <- tempfile("git2rdata-")
dir.create(root)

# store a dataframe as git2rdata object. Capture the result to minimise
# screen output
junk <- write_vc(
  iris[1:6, ], "iris", root, sorting = "Sepal.Length", digits = 6
)
# write a standard tab separate file (non git2rdata object)
write.table(iris, file = file.path(root, "standard.tsv"), sep = "\t")
# write a YAML file
yml <- list(
  authors = list(
    "Research Institute for Nature and Forest" = list(
      href = "https://www.inbo.be/en"))
)
yaml::write_yaml(yml, file = file.path(root, "_pkgdown.yml"))

# list the git2rdata objects
list_data(root)
# list the files
list.files(root, recursive = TRUE)

# remove all .tsv files from valid git2rdata objects
rm_data(root, path = ".")
# check the removal of the .tsv file
list.files(root, recursive = TRUE)
list_data(root)

# remove dangling git2rdata metadata files
prune_meta(root, path = ".")
# check the removal of the metadata
list.files(root, recursive = TRUE)
list_data(root)

## on git repo

# initialise a git repo using git2r
repo_path <- tempfile("git2rdata-repo-")
dir.create(repo_path)
repo <- git2r::init(repo_path)
git2r::config(repo, user.name = "Alice", user.email = "alice@example.org")
```

```

# store a dataframe
write_vc(
  iris[1:6, ], "iris", repo, sorting = "Sepal.Length", stage = TRUE,
  digits = 6
)
# check that the dataframe is stored
status(repo)
list_data(repo)

# commit the current version and check the git repo
commit(repo, "add iris data", session = TRUE)
status(repo)

# remove the data files from the repo
rm_data(repo, path = ".")
# check the removal
list_data(repo)
status(repo)

# remove dangling metadata
prune_meta(repo, path = ".")
# check the removal
list_data(repo)
status(repo)

```

---

prune\_meta

*Prune Metadata Files*


---

## Description

Removes all **valid** metadata (.yml files) from the path when they don't have accompanying data (.tsv file). **Invalid** metadata triggers a warning without removing the metadata file.

Use this function with caution since it will remove all valid metadata files without asking for confirmation. We strongly recommend to use this function on files under version control. See vignette("workflow", package = "git2rdata") for some examples on how to use this.

## Usage

```

prune_meta(root = ".", path = NULL, recursive = TRUE, ...)

## S3 method for class 'git_repository'
prune_meta(root, path = NULL, recursive = TRUE, ..., stage = FALSE)

```

## Arguments

root	The root of a project. Can be a file path or a git-repository. Defaults to the current working directory (".").
path	the directory in which to clean all the data files. The directory is relative to root.

recursive	remove files in subdirectories too.
...	parameters used in some methods
stage	stage the changes after removing the files. Defaults to FALSE.

**Value**

returns invisibly a vector of removed files names. The paths are relative to root.

**See Also**

Other storage: [data\\_package\(\)](#), [display\\_metadata\(\)](#), [list\\_data\(\)](#), [read\\_vc\(\)](#), [relabel\(\)](#), [rename\\_variable\(\)](#), [rm\\_data\(\)](#), [update\\_metadata\(\)](#), [verify\\_vc\(\)](#), [write\\_vc\(\)](#)

**Examples**

```
## on file system

# create a directory
root <- tempfile("git2rdata-")
dir.create(root)

# store a dataframe as git2rdata object. Capture the result to minimise
# screen output
junk <- write_vc(
  iris[1:6, ], "iris", root, sorting = "Sepal.Length", digits = 6
)
# write a standard tab separate file (non git2rdata object)
write.table(iris, file = file.path(root, "standard.tsv"), sep = "\t")
# write a YAML file
yml <- list(
  authors = list(
    "Research Institute for Nature and Forest" = list(
      href = "https://www.inbo.be/en"))
)
yaml::write_yaml(yml, file = file.path(root, "_pkgdown.yml"))

# list the git2rdata objects
list_data(root)
# list the files
list.files(root, recursive = TRUE)

# remove all .tsv files from valid git2rdata objects
rm_data(root, path = ".")
# check the removal of the .tsv file
list.files(root, recursive = TRUE)
list_data(root)

# remove dangling git2rdata metadata files
prune_meta(root, path = ".")
# check the removal of the metadata
list.files(root, recursive = TRUE)
list_data(root)
```

```
## on git repo

# initialise a git repo using git2r
repo_path <- tempfile("git2rdata-repo-")
dir.create(repo_path)
repo <- git2r::init(repo_path)
git2r::config(repo, user.name = "Alice", user.email = "alice@example.org")

# store a dataframe
write_vc(
  iris[1:6, ], "iris", repo, sorting = "Sepal.Length", stage = TRUE,
  digits = 6
)
# check that the dataframe is stored
status(repo)
list_data(repo)

# commit the current version and check the git repo
commit(repo, "add iris data", session = TRUE)
status(repo)

# remove the data files from the repo
rm_data(repo, path = ".")
# check the removal
list_data(repo)
status(repo)

# remove dangling metadata
prune_meta(repo, path = ".")
# check the removal
list_data(repo)
status(repo)
```

---

pull

*Re-exported Function From git2r*

---

### Description

See [pull](#) in `git2r`.

### See Also

Other version\_control: [commit\(\)](#), [push\(\)](#), [recent\\_commit\(\)](#), [repository\(\)](#), [status\(\)](#)

---

push	<i>Re-exported Function From git2r</i>
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---

**Description**

See [push](#) in `git2r`.

**See Also**

Other version\_control: [commit\(\)](#), [pull\(\)](#), [recent\\_commit\(\)](#), [repository\(\)](#), [status\(\)](#)

---

read_vc	<i>Read a Git2rdata Object from Disk</i>
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---

**Description**

`read_vc()` handles `git2rdata` objects stored by `write_vc()`. It reads and verifies the metadata file (`.yaml`). Then it reads and verifies the raw data. The last step is back-transforming any transformation done by `meta()` to return the `data.frame` as stored by `write_vc()`.

`read_vc()` is an S3 generic on `root` which currently handles "character" (a path) and "git-repository" (from `git2r`). S3 methods for other version control system could be added.

**Usage**

```
read_vc(file, root = ".")
```

**Arguments**

<code>file</code>	the name of the <code>git2rdata</code> object. <code>Git2rdata</code> objects cannot have dots in their name. The name may include a relative path. <code>file</code> is a path relative to the root. Note that <code>file</code> must point to a location within root.
<code>root</code>	The root of a project. Can be a file path or a <code>git-repository</code> . Defaults to the current working directory ( <code>"."</code> ).

**Value**

The `data.frame` with the file names and hashes as attributes. It has the additional class "`git2rdata`" to support extra methods to display the descriptions.

**See Also**

Other storage: [data\\_package\(\)](#), [display\\_metadata\(\)](#), [list\\_data\(\)](#), [prune\\_meta\(\)](#), [relabel\(\)](#), [rename\\_variable\(\)](#), [rm\\_data\(\)](#), [update\\_metadata\(\)](#), [verify\\_vc\(\)](#), [write\\_vc\(\)](#)

**Examples**

```

## on file system

# create a directory
root <- tempfile("git2rdata-")
dir.create(root)

# write a dataframe to the directory
write_vc(
  iris[1:6, ], file = "iris", root = root, sorting = "Sepal.Length",
  digits = 6
)
# check that a data file (.tsv) and a metadata file (.yaml) exist.
list.files(root, recursive = TRUE)
# read the git2rdata object from the directory
read_vc("iris", root)

# store a new version with different observations but the same metadata
write_vc(iris[1:5, ], "iris", root)
list.files(root, recursive = TRUE)
# Removing a column requires version requires new metadata.
# Add strict = FALSE to override the existing metadata.
write_vc(
  iris[1:6, -2], "iris", root, sorting = "Sepal.Length", strict = FALSE
)
list.files(root, recursive = TRUE)
# storing the original version again requires another update of the metadata
write_vc(iris[1:6, ], "iris", root, sorting = "Sepal.Width", strict = FALSE)
list.files(root, recursive = TRUE)
# optimize = FALSE stores the data more verbose. This requires larger files.
write_vc(
  iris[1:6, ], "iris2", root, sorting = "Sepal.Width", optimize = FALSE,
  digits = 6
)
list.files(root, recursive = TRUE)

## on git repo using a git2r::git-repository

# initialise a git repo using the git2r package
repo_path <- tempfile("git2rdata-repo-")
dir.create(repo_path)
repo <- git2r::init(repo_path)
git2r::config(repo, user.name = "Alice", user.email = "alice@example.org")

# store a dataframe in git repo.
write_vc(
  iris[1:6, ], file = "iris", root = repo, sorting = "Sepal.Length",
  digits = 6
)
# This git2rdata object is not staged by default.

```

```

status(repo)
# read a dataframe from a git repo
read_vc("iris", repo)

# store a new version in the git repo and stage it in one go
write_vc(iris[1:5, ], "iris", repo, stage = TRUE)
status(repo)

# store a verbose version in a different git2rdata object
write_vc(
  iris[1:6, ], "iris2", repo, sorting = "Sepal.Width", optimize = FALSE,
  digits = 6
)
status(repo)

```

---

recent\_commit

*Retrieve the Most Recent File Change*


---

## Description

Retrieve the most recent commit that added or updated a file or git2rdata object. This does not imply that file still exists at the current HEAD as it ignores the deletion of files.

Use this information to document the current version of file or git2rdata object in an analysis. Since it refers to the most recent change of this file, it remains unchanged by committing changes to other files. You can also use it to track if data got updated, requiring an analysis to be rerun. See `vignette("workflow", package = "git2rdata")`.

## Usage

```
recent_commit(file, root, data = FALSE)
```

## Arguments

<code>file</code>	the name of the git2rdata object. Git2rdata objects cannot have dots in their name. The name may include a relative path. <code>file</code> is a path relative to the root. Note that <code>file</code> must point to a location within <code>root</code> .
<code>root</code>	The root of a project. Can be a file path or a git-repository.
<code>data</code>	does <code>file</code> refers to a data object (TRUE) or to a file (FALSE)? Defaults to FALSE.

## Value

a data.frame with `commit`, `author` and `when` for the most recent commit that adds or updates the file.

## See Also

Other version\_control: [commit\(\)](#), [pull\(\)](#), [push\(\)](#), [repository\(\)](#), [status\(\)](#)

## Examples

```
# initialise a git repo using git2r
repo_path <- tempfile("git2rdata-repo")
dir.create(repo_path)
repo <- git2r::init(repo_path)
git2r::config(repo, user.name = "Alice", user.email = "alice@example.org")

# write and commit a first dataframe
# store the output of write_vc() minimize screen output
junk <- write_vc(
  iris[1:6, ], "iris", repo, sorting = "Sepal.Length", stage = TRUE,
  digits = 6
)
commit(repo, "important analysis", session = TRUE)
list.files(repo_path)

# write and commit a second dataframe
junk <- write_vc(
  iris[7:12, ], "iris2", repo, sorting = "Sepal.Length", stage = TRUE,
  digits = 6
)
commit(repo, "important analysis", session = TRUE)
list.files(repo_path)

# write and commit a new version of the first dataframe
junk <- write_vc(iris[7:12, ], "iris", repo, stage = TRUE)
list.files(repo_path)
commit(repo, "important analysis", session = TRUE)

# find out in which commit a file was last changed

# "iris.tsv" was last updated in the third commit
recent_commit("iris.tsv", repo)
# "iris.yml" was last updated in the first commit
recent_commit("iris.yml", repo)
# "iris2.yml" was last updated in the second commit
recent_commit("iris2.yml", repo)
# the git2rdata object "iris" was last updated in the third commit
recent_commit("iris", repo, data = TRUE)
```

---

relabel

*Relabel Factor Levels by Updating the Metadata*


---

## Description

Imagine the situation where we have a dataframe with a factor variable and we have stored it with `write_vc(optimize = TRUE)`. The raw data file contains the factor indices and the metadata contains the link between the factor index and the corresponding label. See `vignette("version_control", package = "git2rdata")`. In such a case, relabelling a factor can be fast and lightweight by updating the metadata.

**Usage**

```
relabel(file, root = ".", change)
```

**Arguments**

file	the name of the <code>git2rdata</code> object. <code>Git2rdata</code> objects cannot have dots in their name. The name may include a relative path. <code>file</code> is a path relative to the root. Note that <code>file</code> must point to a location within <code>root</code> .
root	The root of a project. Can be a file path or a <code>git</code> -repository. Defaults to the current working directory ( <code>"."</code> ).
change	either a <code>list</code> or a <code>data.frame</code> . In case of a <code>list</code> is a named <code>list</code> with named vectors. The names of <code>list</code> elements must match the names of the variables. The names of the vector elements must match the existing factor labels. The values represent the new factor labels. In case of a <code>data.frame</code> it needs to have the variables <code>factor</code> (name of the factor), <code>old</code> (the old) factor label and <code>new</code> (the new factor label). <code>relabel()</code> ignores all other columns.

**Value**

invisible `NULL`.

**See Also**

Other storage: [data\\_package\(\)](#), [display\\_metadata\(\)](#), [list\\_data\(\)](#), [prune\\_meta\(\)](#), [read\\_vc\(\)](#), [rename\\_variable\(\)](#), [rm\\_data\(\)](#), [update\\_metadata\(\)](#), [verify\\_vc\(\)](#), [write\\_vc\(\)](#)

**Examples**

```
# initialise a git repo using git2r
repo_path <- tempfile("git2rdata-repo-")
dir.create(repo_path)
repo <- git2r::init(repo_path)
git2r::config(repo, user.name = "Alice", user.email = "alice@example.org")

# Create a dataframe and store it as an optimized git2rdata object.
# Note that write_vc() uses optimization by default.
# Stage and commit the git2rdata object.
ds <- data.frame(
  a = c("a1", "a2"),
  b = c("b2", "b1"),
  stringsAsFactors = TRUE
)
junk <- write_vc(ds, "relabel", repo, sorting = "b", stage = TRUE)
cm <- commit(repo, "initial commit")
# check that the workspace is clean
status(repo)

# Define new labels as a list and apply them to the git2rdata object.
new_labels <- list(
  a = list(a2 = "a3")
```

```

)
relabel("relabel", repo, new_labels)
# check the changes
read_vc("relabel", repo)
# relabel() changed the metadata, not the raw data
status(repo)
git2r::add(repo, "relabel.*")
cm <- commit(repo, "relabel using a list")

# Define new labels as a dataframe and apply them to the git2rdata object
change <- data.frame(
  factor = c("a", "a", "b"),
  old = c("a3", "a1", "b2"),
  new = c("c2", "c1", "b3"),
  stringsAsFactors = TRUE
)
relabel("relabel", repo, change)
# check the changes
read_vc("relabel", repo)
# relabel() changed the metadata, not the raw data
status(repo)

```

---

rename\_variable

*Rename a Variable*

---

## Description

The raw data file contains a header with the variable names. The metadata list the variable names and their type. Changing a variable name and overwriting the `git2rdata` object with result in an error. Because it will look like removing an existing variable and adding a new one. Overwriting the object with `strict = FALSE` potentially changes the order of the variables, leading to a large diff.

## Usage

```

rename_variable(file, change, root = ".", ...)

## S3 method for class 'character'
rename_variable(file, change, root = ".", ...)

## Default S3 method:
rename_variable(file, change, root, ...)

## S3 method for class 'git_repository'
rename_variable(file, change, root, ..., stage = FALSE, force = FALSE)

```

**Arguments**

file	the name of the git2rdata object. Git2rdata objects cannot have dots in their name. The name may include a relative path. file is a path relative to the root. Note that file must point to a location within root.
change	A named vector with the old names as values and the new names as names.
root	The root of a project. Can be a file path or a git-repository. Defaults to the current working directory (".").
...	parameters used in some methods
stage	Logical value indicating whether to stage the changes after writing the data. Defaults to FALSE.
force	Add ignored files. Default is FALSE.

**Details**

This function solves this by only updating the raw data header and the metadata.

**Value**

a named vector with the file paths relative to root. The names contain the hashes of the files.

**See Also**

Other storage: [data\\_package\(\)](#), [display\\_metadata\(\)](#), [list\\_data\(\)](#), [prune\\_meta\(\)](#), [read\\_vc\(\)](#), [relabel\(\)](#), [rm\\_data\(\)](#), [update\\_metadata\(\)](#), [verify\\_vc\(\)](#), [write\\_vc\(\)](#)

**Examples**

```
# initialise a git repo using git2r
repo_path <- tempfile("git2rdata-repo-")
dir.create(repo_path)
repo <- git2r::init(repo_path)
git2r::config(repo, user.name = "Alice", user.email = "alice@example.org")

# Create a dataframe and store it as an optimized git2rdata object.
# Note that write_vc() uses optimization by default.
# Stage and commit the git2rdata object.
ds <- data.frame(
  a = c("a1", "a2"),
  b = c("b2", "b1"),
  stringsAsFactors = TRUE
)
junk <- write_vc(ds, "rename", repo, sorting = "b", stage = TRUE)
cm <- commit(repo, "initial commit")
# check that the workspace is clean
status(repo)

# Define change.
change <- c(new_name = "a")
rename_variable(file = "rename", change = change, root = repo)
```

```
# check the changes
read_vc("rename", repo)
status(repo)
```

---

 repository

*Re-exported Function From git2r*


---

### Description

See [repository](#) in git2r.

### See Also

Other version\_control: [commit\(\)](#), [pull\(\)](#), [push\(\)](#), [recent\\_commit\(\)](#), [status\(\)](#)

---

 rm\_data

*Remove Data Files From Git2rdata Objects*


---

### Description

Remove the data (.tsv) file from all valid git2rdata objects at the path. The metadata remains untouched. A warning lists any git2rdata object with **invalid** metadata. The function keeps any .tsv file with invalid metadata or from non-git2rdata objects.

Use this function with caution since it will remove all valid data files without asking for confirmation. We strongly recommend to use this function on files under version control. See `vignette("workflow", package = "git2rdata")` for some examples on how to use this.

### Usage

```
rm_data(root = ".", path = NULL, recursive = TRUE, ...)
```

```
## S3 method for class 'git_repository'
rm_data(
  root,
  path = NULL,
  recursive = TRUE,
  ...,
  stage = FALSE,
  type = c("unmodified", "modified", "ignored", "all")
)
```

**Arguments**

root	The root of a project. Can be a file path or a git-repository. Defaults to the current working directory (".").
path	the directory in which to clean all the data files. The directory is relative to root.
recursive	remove files in subdirectories too.
...	parameters used in some methods
stage	stage the changes after removing the files. Defaults to FALSE.
type	Defines the classes of files to remove. <code>unmodified</code> are files in the git history and unchanged since the last commit. <code>modified</code> are files in the git history and changed since the last commit. <code>ignored</code> refers to file listed in a <code>.gitignore</code> file. Selecting <code>modified</code> will remove both unmodified and modified data files. Selecting <code>ignored</code> will remove unmodified, modified and ignored data files. <code>all</code> refers to all visible data files, including untracked files.

**Value**

returns invisibly a vector of removed files names. The paths are relative to root.

**See Also**

Other storage: [data\\_package\(\)](#), [display\\_metadata\(\)](#), [list\\_data\(\)](#), [prune\\_meta\(\)](#), [read\\_vc\(\)](#), [relabel\(\)](#), [rename\\_variable\(\)](#), [update\\_metadata\(\)](#), [verify\\_vc\(\)](#), [write\\_vc\(\)](#)

**Examples**

```
## on file system

# create a directory
root <- tempfile("git2rdata-")
dir.create(root)

# store a dataframe as git2rdata object. Capture the result to minimise
# screen output
junk <- write_vc(
  iris[1:6, ], "iris", root, sorting = "Sepal.Length", digits = 6
)
# write a standard tab separate file (non git2rdata object)
write.table(iris, file = file.path(root, "standard.tsv"), sep = "\t")
# write a YAML file
yaml <- list(
  authors = list(
    "Research Institute for Nature and Forest" = list(
      href = "https://www.inbo.be/en"))
)
yaml::write_yaml(yaml, file = file.path(root, "_pkgdown.yaml"))

# list the git2rdata objects
list_data(root)
# list the files
list.files(root, recursive = TRUE)
```

```
# remove all .tsv files from valid git2rdata objects
rm_data(root, path = ".")
# check the removal of the .tsv file
list.files(root, recursive = TRUE)
list_data(root)

# remove dangling git2rdata metadata files
prune_meta(root, path = ".")
# check the removal of the metadata
list.files(root, recursive = TRUE)
list_data(root)

## on git repo

# initialise a git repo using git2r
repo_path <- tempfile("git2rdata-repo-")
dir.create(repo_path)
repo <- git2r::init(repo_path)
git2r::config(repo, user.name = "Alice", user.email = "alice@example.org")

# store a dataframe
write_vc(
  iris[1:6, ], "iris", repo, sorting = "Sepal.Length", stage = TRUE,
  digits = 6
)
# check that the dataframe is stored
status(repo)
list_data(repo)

# commit the current version and check the git repo
commit(repo, "add iris data", session = TRUE)
status(repo)

# remove the data files from the repo
rm_data(repo, path = ".")
# check the removal
list_data(repo)
status(repo)

# remove dangling metadata
prune_meta(repo, path = ".")
# check the removal
list_data(repo)
status(repo)
```

**Description**

See [status](#) in `git2r`.

**See Also**

Other version\_control: [commit\(\)](#), [pull\(\)](#), [push\(\)](#), [recent\\_commit\(\)](#), [repository\(\)](#)

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update_metadata	<i>Update the description of a git2rdata object</i>
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---

**Description**

Allows to update the description of the fields, the table name, the title, and the description of a `git2rdata` object. All arguments are optional. Setting an argument to `NA` or an empty string will remove the corresponding field from the metadata.

**Usage**

```
update_metadata(  
  file,  
  root = ".",  
  field_description,  
  name,  
  title,  
  description,  
  ...  
)
```

**Arguments**

<code>file</code>	the name of the <code>git2rdata</code> object. <code>Git2rdata</code> objects cannot have dots in their name. The name may include a relative path. <code>file</code> is a path relative to the root. Note that <code>file</code> must point to a location within <code>root</code> .
<code>root</code>	The root of a project. Can be a file path or a <code>git-repository</code> . Defaults to the current working directory (" <code>.</code> ").
<code>field_description</code>	a named character vector with the new descriptions for the fields. The names of the vector must match the variable names.
<code>name</code>	a character string with the new table name of the object.
<code>title</code>	a character string with the new title of the object.
<code>description</code>	a character string with the new description of the object.
<code>...</code>	parameters used in some methods

**See Also**

Other storage: [data\\_package\(\)](#), [display\\_metadata\(\)](#), [list\\_data\(\)](#), [prune\\_meta\(\)](#), [read\\_vc\(\)](#), [relabel\(\)](#), [rename\\_variable\(\)](#), [rm\\_data\(\)](#), [verify\\_vc\(\)](#), [write\\_vc\(\)](#)

---

verify_vc	<i>Read a file and verify the presence of variables</i>
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**Description**

Reads the file with `read_vc()`. Then verifies that every variable listed in `variables` is present in the `data.frame`.

**Usage**

```
verify_vc(file, root, variables)
```

**Arguments**

<code>file</code>	the name of the <code>git2rdata</code> object. <code>Git2rdata</code> objects cannot have dots in their name. The name may include a relative path. <code>file</code> is a path relative to the root. Note that <code>file</code> must point to a location within <code>root</code> .
<code>root</code>	The root of a project. Can be a file path or a <code>git-repository</code> . Defaults to the current working directory ( <code>"."</code> ).
<code>variables</code>	a character vector with variable names.

**See Also**

Other storage: `data_package()`, `display_metadata()`, `list_data()`, `prune_meta()`, `read_vc()`, `relabel()`, `rename_variable()`, `rm_data()`, `update_metadata()`, `write_vc()`

---

write_vc	<i>Store a Data.Frame as a Git2rdata Object on Disk</i>
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---

**Description**

A `git2rdata` object consists of two files. The `".tsv"` file contains the raw data as a plain text tab separated file. The `".yaml"` contains the metadata on the columns in plain text YAML format. See `vignette("plain text", package = "git2rdata")` for more details on the implementation.

**Usage**

```
write_vc(
  x,
  file,
  root = ".",
  sorting,
  strict = TRUE,
  optimize = TRUE,
  na = "NA",
```

```

    ...,
    split_by,
    convert
  )

## S3 method for class 'character'
write_vc(
  x,
  file,
  root = ".",
  sorting,
  strict = TRUE,
  optimize = TRUE,
  na = "NA",
  ...,
  append = FALSE,
  split_by = character(0),
  digits,
  convert = list()
)

## S3 method for class 'git_repository'
write_vc(
  x,
  file,
  root,
  sorting,
  strict = TRUE,
  optimize = TRUE,
  na = "NA",
  ...,
  stage = FALSE,
  force = FALSE
)

```

### Arguments

x	the data.frame.
file	the name of the git2rdata object. Git2rdata objects cannot have dots in their name. The name may include a relative path. file is a path relative to the root. Note that file must point to a location within root.
root	The root of a project. Can be a file path or a git-repository. Defaults to the current working directory (".").
sorting	an optional vector of column names defining which columns to use for sorting x and in what order to use them. The default empty sorting yields a warning. Add sorting to avoid this warning. Strongly recommended in combination with version control. See vignette("efficiency", package = "git2rdata") for an illustration of the importance of sorting.

strict	What to do when the metadata changes. <code>strict = FALSE</code> overwrites the data and the metadata with a warning listing the changes, <code>strict = TRUE</code> returns an error and leaves the data and metadata as is. Defaults to <code>TRUE</code> .
optimize	If <code>TRUE</code> , recode the data to get smaller text files. If <code>FALSE</code> , <code>meta()</code> converts the data to character. Defaults to <code>TRUE</code> .
na	the string to use for missing values in the data.
...	parameters used in some methods
split_by	An optional vector of variables name to split the text files. This creates a separate file for every combination. We prepend these variables to the vector of sorting variables.
convert	An optional named list for column conversions. Names must be present in the column names of <code>x</code> . Each element must be a character vector of length 2 with names <code>write</code> and <code>read</code> , containing function names in the package::function format. The <code>write</code> function is applied before storing, and <code>read</code> function is applied when reading back the data.
append	logical. Only relevant if <code>file</code> is a character string. If <code>TRUE</code> , the output is appended to the file. If <code>FALSE</code> , any existing file of the name is destroyed.
digits	The number of significant digits of the smallest absolute value. The function applies the rounding automatically. Only relevant for numeric variables. Either a single positive integer or a named vector where the names link to the variables in the <code>data.frame</code> . Defaults to 6 with a warning.
stage	Logical value indicating whether to stage the changes after writing the data. Defaults to <code>FALSE</code> .
force	Add ignored files. Default is <code>FALSE</code> .

**Value**

a named vector with the file paths relative to `root`. The names contain the hashes of the files.

**Note**

`..generic` is a reserved name for the metadata and is a forbidden column name in a `data.frame`.

**See Also**

Other storage: [data\\_package\(\)](#), [display\\_metadata\(\)](#), [list\\_data\(\)](#), [prune\\_meta\(\)](#), [read\\_vc\(\)](#), [relabel\(\)](#), [rename\\_variable\(\)](#), [rm\\_data\(\)](#), [update\\_metadata\(\)](#), [verify\\_vc\(\)](#)

**Examples**

```
## on file system

# create a directory
root <- tempfile("git2rdata-")
dir.create(root)

# write a dataframe to the directory
```

```

write_vc(
  iris[1:6, ], file = "iris", root = root, sorting = "Sepal.Length",
  digits = 6
)
# check that a data file (.tsv) and a metadata file (.yaml) exist.
list.files(root, recursive = TRUE)
# read the git2rdata object from the directory
read_vc("iris", root)

# store a new version with different observations but the same metadata
write_vc(iris[1:5, ], "iris", root)
list.files(root, recursive = TRUE)
# Removing a column requires version requires new metadata.
# Add strict = FALSE to override the existing metadata.
write_vc(
  iris[1:6, -2], "iris", root, sorting = "Sepal.Length", strict = FALSE
)
list.files(root, recursive = TRUE)
# storing the original version again requires another update of the metadata
write_vc(iris[1:6, ], "iris", root, sorting = "Sepal.Width", strict = FALSE)
list.files(root, recursive = TRUE)
# optimize = FALSE stores the data more verbose. This requires larger files.
write_vc(
  iris[1:6, ], "iris2", root, sorting = "Sepal.Width", optimize = FALSE,
  digits = 6
)
list.files(root, recursive = TRUE)

## on git repo using a git2r::git-repository

# initialise a git repo using the git2r package
repo_path <- tempfile("git2rdata-repo-")
dir.create(repo_path)
repo <- git2r::init(repo_path)
git2r::config(repo, user.name = "Alice", user.email = "alice@example.org")

# store a dataframe in git repo.
write_vc(
  iris[1:6, ], file = "iris", root = repo, sorting = "Sepal.Length",
  digits = 6
)
# This git2rdata object is not staged by default.
status(repo)
# read a dataframe from a git repo
read_vc("iris", repo)

# store a new version in the git repo and stage it in one go
write_vc(iris[1:5, ], "iris", repo, stage = TRUE)
status(repo)

# store a verbose version in a different git2rdata object

```

```
write_vc(  
  iris[1:6, ], "iris2", repo, sorting = "Sepal.Width", optimize = FALSE,  
  digits = 6  
)  
status(repo)
```

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