

# Package ‘DataPackageR’

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**Type** Package

**Title** Construct Reproducible Analytic Data Sets as R Packages

**Version** 0.15.9

**Description** A framework to help construct R data packages in a reproducible manner. Potentially time consuming processing of raw data sets into analysis ready data sets is done in a reproducible manner and decoupled from the usual 'R CMD build' process so that data sets can be processed into R objects in the data package and the data package can then be shared, built, and installed by others without the need to repeat computationally costly data processing. The package maintains data provenance by turning the data processing scripts into package vignettes, as well as enforcing documentation and version checking of included data objects. Data packages can be version controlled on 'GitHub', and used to share data for manuscripts, collaboration and reproducible research.

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**Depends** R (>= 3.5.0)

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<https://docs.ropensci.org/DataPackageR/>

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**Author** Greg Finak [aut, cph] (Original author and creator of DataPackageR),  
 Paul Obrecht [ctb],  
 Ellis Hughes [ctb] (<<https://orcid.org/0000-0003-0637-4436>>),  
 Jimmy Fulp [ctb],  
 Marie Vendettuoli [ctb] (<<https://orcid.org/0000-0001-9321-1410>>),  
 Dave Slager [ctb, cre] (<<https://orcid.org/0000-0003-2525-2039>>),  
 Jason Taylor [ctb],  
 Kara Woo [rev] (Kara reviewed the package for ropensci, see  
 <<https://github.com/ropensci/onboarding/issues/230>>),  
 William Landau [rev] (William reviewed the package for ropensci, see  
 <<https://github.com/ropensci/onboarding/issues/230>>)

**Maintainer** Dave Slager <dslager@scharp.org>

**Repository** CRAN

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---

assert\_data\_version    *Assert that a data version in a data package matches an expectation.*

---

## Description

Assert that a data version in a data package matches an expectation.

**Usage**

```
assert_data_version(
  data_package_name = NULL,
  version_string = NULL,
  acceptable = "equal",
  ...
)
```

**Arguments**

`data_package_name` character Name of the package.

`version_string` character Version string in "x.y.z" format.

`acceptable` character one of "equal", "equal\_or\_greater", describing what version match is acceptable.

`...` additional arguments passed to `data_version` (such as `lib.loc`)

**Details**

Tests the `DataVersion` string in `data_package_name` against `version_string` testing the major, minor and revision portion.

Tests "data\_package\_name version equal version\_string" or "data\_package\_name version equal\_or\_greater version\_string".

**Value**

invisible logical TRUE if success, otherwise stop on mismatch.

**Examples**

```
if(rmarkdown::pandoc_available()){
  f <- tempdir()
  f <- file.path(f, "foo.Rmd")
  con <- file(f)
  writeLines("```{r}\n vec = 1:10 \n```\n",con = con)
  close(con)
  pname <- basename(tempfile())
  datapackage_skeleton(name = pname,
    path=tempdir(),
    force = TRUE,
    r_object_names = "vec",
    code_files = f)
  package_build(file.path(tempdir(),pname), install = FALSE)

  devtools::load_all(file.path(tempdir(),pname))

  assert_data_version(data_package_name = pname,version_string = "0.1.0",acceptable = "equal")
}
```

---

`construct_yaml_config` *Construct a datapackager.yml configuration*

---

### Description

Constructs a `datapackager.yml` configuration object from a vector of file names and a vector of object names (all quoted). Can be written to disk via `yaml_write`. `render_root` is set to a randomly generated named subdirectory of `tempdir()`.

### Usage

```
construct_yaml_config(code = NULL, data = NULL, render_root = NULL)
```

### Arguments

<code>code</code>	A vector of filenames
<code>data</code>	A vector of quoted object names
<code>render_root</code>	The root directory where the package data processing code will be rendered. Defaults to is set to a randomly generated named subdirectory of <code>tempdir()</code> .

### Value

a `datapackager.yml` configuration represented as an R object

### Examples

```
conf <- construct_yaml_config(code = c('file1.rmd', 'file2.rmd'), data=c('object1', 'object2'))
tmp <- normalizePath(tempdir(), winslash = "/")
yaml_write(conf, path=tmp)
```

---

`DataPackageR` *Process data generation code in 'data-raw'*

---

### Description

Assumes `.R` files in `'data-raw'` generate `rda` files to be stored in `'data'`. Sources `datasets.R` which can source other R files. R files sourced by `datasets.R` must invoke `sys.source('myRfile.R', env=topenv())`. Meant to be called before R CMD build.

### Usage

```
DataPackageR(arg = NULL, deps = TRUE)
```

**Arguments**

arg                    character name of the package to build.  
 deps                  logical should scripts pass data objects to each other (default=TRUE)

**Value**

logical TRUE if successful, FALSE, if not.

---

datapackager\_object\_read

*Read an object created in a previously run processing script.*

---

**Description**

Read an object created in a previously run processing script.

**Usage**

```
datapackager_object_read(name)
```

**Arguments**

name                  character the name of the object. Must be a name available in the configuration objects. Other objects are not saved.

**Details**

This function is only accessible within an R or Rmd file processed by DataPackageR. It searches for an environment named ENVS within the current environment, that holds the object with the given name. Such an environment is constructed and populated with objects specified in the yaml objects property and passed along to subsequent R and Rmd files as DataPackageR processes them in order.

**Value**

An R object.

**Examples**

```
if(rmarkdown::pandoc_available()){
  ENVS <- new.env() # ENVS would be in the environment
                  # where the data processing is run. It is
                  # handled automatically by the package.
  assign("find_me", 100, ENVS) #This is done automatically by DataPackageR

  find_me <- datapackager_object_read("find_me") # This would appear in an Rmd processed by
                                                # DataPackageR to access the object named "find_me" created
                                                # by a previous script. "find_me" would also need to
```

```

    # appear in the objects property of config.yml
  }

```

---

DataPackageR\_options    *Options consulted by DataPackageR*

---

### Description

User-configurable options consulted by DataPackageR, which provide a mechanism for setting default behaviors for various functions.

If the built-in defaults don't suit you, set one or more of these options. Typically, this is done in the `.Rprofile` startup file, which you can open for editing with `usethis::edit_r_profile()` - this will set the specified options for all future R sessions. The following setting is recommended to not be prompted upon each package build for a NEWS update:

```
options(DataPackageR_interact = FALSE)
```

### Options for the DataPackageR package

- `DataPackageR_interact`: Upon package load, this defaults to the value of `interactive()`, unless the option has been previously set (e.g., in `.Rprofile`). `TRUE` prompts user interactively for a NEWS update on `package_build()`. See the example above and the [rOpenSci blog post](<https://ropensci.org/blog/2018/09/>) for more details on how to set this to `FALSE`, which will never prompt user for a NEWS update. `FALSE` is also the setting used for DataPackageR's internal tests.
- `DataPackageR_verbose`: Default upon package load is `TRUE`. `FALSE` suppresses all console output and is currently only used for DataPackageR's automated unit tests.
- `DataPackageR_packagebuilding`: Default upon package load is `FALSE`. This option is used internally for package operations and changing it is not recommended.

---

datapackage\_skeleton    *Create a Data Package skeleton for use with DataPackageR.*

---

### Description

Creates a package skeleton directory structure for use with DataPackageR. Adds the `DataVersion` string to `DESCRIPTION`, creates the `DATADIGEST` file, and the `data-raw` directory. Updates the `Read-and-delete-me` file to reflect the additional necessary steps.

**Usage**

```
datapackage_skeleton(  
  name = NULL,  
  path = ".",  
  force = FALSE,  
  code_files = character(),  
  r_object_names = character(),  
  raw_data_dir = character(),  
  dependencies = character()  
)  
  
datapackage.skeleton(  
  name = NULL,  
  list = character(),  
  environment = .GlobalEnv,  
  path = ".",  
  force = FALSE,  
  code_files = character(),  
  r_object_names = character()  
)
```

**Arguments**

<code>name</code>	character name of the package to create.
<code>path</code>	A character path where the package is located. See <a href="#">package.skeleton</a>
<code>force</code>	logical Force the package skeleton to be recreated even if it exists. see <a href="#">package.skeleton</a>
<code>code_files</code>	Optional character vector of paths to Rmd files that process raw data into R objects.
<code>r_object_names</code>	vector of quoted r object names , tables, etc. created when the files in <code>code_files</code> are run.
<code>raw_data_dir</code>	character pointing to a raw data directory. Will be moved with all its subdirectories to "inst/extdata"
<code>dependencies</code>	vector of character, paths to R files that will be moved to "data-raw" but not included in the yaml config file. e.g., dependency scripts.
<code>list</code>	Not used.
<code>environment</code>	Not used.

**Value**

No return value, called for side effects

**Note**

renamed `datapackage.skeleton()` to `datapackage_skeleton()`.

## Examples

```
if(rmarkdown::pandoc_available()){
  f <- tempdir()
  f <- file.path(f,"foo.Rmd")
  con <- file(f)
  writeLines("```${r}``\n tbl = data.frame(1:10) \n```\n",con=con)
  close(con)
  pname <- basename(tempfile())
  datapackage_skeleton(name = pname,
    path = tempdir(),
    force = TRUE,
    r_object_names = "tbl",
    code_files = f)
}
```

---

data\_version

*Get the DataVersion for a package*

---

## Description

Retrieves the DataVersion of a package if available

## Usage

```
data_version(pkg, lib.loc = NULL)
```

```
dataVersion(pkg, lib.loc = NULL)
```

## Arguments

pkg                    character the package name  
lib.loc                character path to library location.

## Value

Object of class 'package\_version' and 'numeric\_version' specifying the DataVersion of the package

## Note

dataVersion() has been renamed to data\_version()

## See Also

[packageVersion](#)

## Examples

```
if(rmarkdown::pandoc_available()){
  f <- tempdir()
  f <- file.path(f, "foo.Rmd")
  con <- file(f)
  writeLines("```${r}``\n vec = 1:10 \n```\n", con=con)
  close(con)
  pname <- basename(tempfile())
  datapackage_skeleton(name = pname,
    path=tempdir(),
    force = TRUE,
    r_object_names = "vec",
    code_files = f)

  package_build(file.path(tempdir(),pname), install = FALSE)

  devtools::load_all(file.path(tempdir(),pname))
  data_version(pname)
}
```

---

document

*Build documentation for a data package using DataPackageR.*

---

## Description

Build documentation for a data package using DataPackageR.

## Usage

```
document(path = ".", install = TRUE, ...)
```

## Arguments

path	character the path to the data package source root.
install	logical install and reload the package. (default TRUE)
...	additional arguments to install

## Value

Called for side effects. Returns TRUE on successful exit.

## Examples

```
# A simple Rmd file that creates one data object
# named "tbl".
if(rmarkdown::pandoc_available()){
  f <- tempdir()
  f <- file.path(f, "foo.Rmd")
```

```
con <- file(f)
writeLines("```{r}\n tbl = data.frame(1:10) \n```\n",con=con)
close(con)

# construct a data package skeleton named "MyDataPackage" and pass
# in the Rmd file name with full path, and the name of the object(s) it
# creates.

pname <- basename(tempfile())
datapackage_skeleton(name=pname,
  path=tempdir(),
  force = TRUE,
  r_object_names = "tbl",
  code_files = f)

# call package_build to run the "foo.Rmd" processing and
# build a data package.
package_build(file.path(tempdir(), pname), install = FALSE)
document(path = file.path(tempdir(), pname), install=FALSE)

}
```

---

keepDataObjects-defunct

*These functions are no longer available.*

---

## Description

These functions are no longer available.

## Usage

```
keepDataObjects(...)
```

## Arguments

... arguments

## Value

Defunct. No return value.

---

package_build	<i>Pre-process, document and build a data package</i>
---------------	---

---

### Description

Combines the preprocessing, documentation, and build steps into one.

### Usage

```
package_build(  
  packageName = NULL,  
  vignettes = FALSE,  
  log = INFO,  
  deps = TRUE,  
  install = FALSE,  
  ...  
)
```

### Arguments

packageName	character path to package source directory. Defaults to the current path when NULL.
vignettes	logical specify whether to build vignettes. Default FALSE.
log	log level INFO, WARN, DEBUG, FATAL
deps	logical should we pass data objects into subsequent scripts? Default TRUE
install	logical automatically install and load the package after building. Default FALSE
...	additional arguments passed to <code>install.packages</code> when <code>install=TRUE</code> .

### Details

Note that if `package_build` returns an error when rendering an `.Rmd` internally, but that same `.Rmd` can be run successfully manually using `rmarkdown::render`, then the following code facilitates debugging. Set `options(error = function(){ sink(); recover()})` before running `package_build`. This will enable examination of the active function calls at the time of the error, with output printed to the console rather than knitr's default sink. After debugging, evaluate `options(error = NULL)` to revert to default error handling. See section "22.5.3 RMarkdown" at <https://adv-r.hadley.nz/debugging.html> for more details.

### Value

Character vector. File path of the built package.

## Examples

```
if(rmarkdown::pandoc_available()){
  f <- tempdir()
  f <- file.path(f, "foo.Rmd")
  con <- file(f)
  writeLines("```${r}``\n tbl = data.frame(1:10) \n```\n", con=con)
  close(con)
  pname <- basename(tempfile())
  datapackage_skeleton(name=pname,
    path=tempdir(),
    force = TRUE,
    r_object_names = "tbl",
    code_files = f)

  package_build(file.path(tempdir(),pname), install = FALSE)
}
```

---

project_data_path	<i>Get DataPackageR data path</i>
-------------------	-----------------------------------

---

## Description

Get DataPackageR data path

## Usage

```
project_data_path(file = NULL)
```

## Arguments

file                    character or NULL (default).

## Details

Returns the path to the data package data subdirectory, or constructs a path to a file in the data subdirectory from the file argument.

## Value

character

## Examples

```
if(rmarkdown::pandoc_available()){
  project_data_path( file = "data.rda" )
}
```

---

project\_extdata\_path *Get DataPackageR extdata path*

---

**Description**

Get DataPackageR extdata path

**Usage**

```
project_extdata_path(file = NULL)
```

**Arguments**

file                    character or NULL (default).

**Details**

Returns the path to the data package extdata subdirectory, or constructs a path to a file in the extdata subdirectory from the file argument.

**Value**

character

**Examples**

```
if(rmarkdown::pandoc_available()){  
  project_extdata_path(file = "mydata.csv")  
}
```

---

project\_path            *Get DataPackageR Project Root Path*

---

**Description**

Get DataPackageR Project Root Path

**Usage**

```
project_path(file = NULL)
```

**Arguments**

file                    character or NULL (default).

**Details**

Returns the path to the data package project root, or constructs a path to a file in the project root from the file argument.

**Value**

character

**Examples**

```
if(rmarkdown::pandoc_available()){
  project_path( file = "DESCRIPTION" )
}
```

---

use_data_object	<i>Add a data object to a data package.</i>
-----------------	---

---

**Description**

The data object will be added to the yml configuration file.

**Usage**

```
use_data_object(object_name = NULL)
```

**Arguments**

`object_name` Name of the data object. Should be created by a processing script in data-raw. character vector of length 1.

**Value**

invisibly returns TRUE for success.

**Examples**

```
if(rmarkdown::pandoc_available()){
  myfile <- tempfile()
  file <- system.file("extdata", "tests", "extra.rmd",
                     package = "DataPackageR")
  datapackage_skeleton(
    name = "datatest",
    path = tempdir(),
    code_files = file,
    force = TRUE,
    r_object_names = "data")
  use_data_object(object_name = "newobject")
}
```

---

use_ignore	<i>Ignore specific files by git and R build.</i>
------------	--

---

**Description**

Ignore specific files by git and R build.

**Usage**

```
use_ignore(file = NULL, path = NULL)
```

**Arguments**

file	character File to ignore.
path	character Path to the file.

**Value**

invisibly returns 0.

**Examples**

```
datapackage_skeleton(name="test",path = tempdir())  
use_ignore("foo", ".")
```

---

use_processing_script	<i>Add a processing script to a data package.</i>
-----------------------	---

---

**Description**

The Rmd or R file or directory specified by file will be moved into the data-raw directory. It will also be added to the yml configuration file. Any existing file by that name will be overwritten when overwrite is set to TRUE

**Usage**

```
use_processing_script(  
  file = NULL,  
  title = NULL,  
  author = NULL,  
  overwrite = FALSE  
)
```

**Arguments**

file	character path to an existing file or name of a new R or Rmd file to create.
title	character title of the processing script for the yaml header. Used only if file is being created.
author	character author name for the yaml header. Used only if the file is being created.
overwrite	logical default FALSE. Overwrite existing file of the same name.

**Value**

invisibly returns TRUE for success. Stops on failure.

**Examples**

```
if(rmarkdown::pandoc_available()){
  myfile <- tempfile()
  file <- system.file("extdata", "tests", "extra.rmd",
                     package = "DataPackageR")
  datapackage_skeleton(
    name = "datatest",
    path = tempdir(),
    code_files = file,
    force = TRUE,
    r_object_names = "data")
  use_processing_script(file = "newScript.Rmd",
                      title = "Processing a new dataset",
                      author = "Y.N. Here.")
}
```

---

use_raw_dataset	<i>Add a raw data set to inst/extdata</i>
-----------------	---

---

**Description**

The file or directory specified by path will be moved into the inst/extdata directory.

**Usage**

```
use_raw_dataset(path = NULL, ignore = FALSE)
```

**Arguments**

path	character path to file or directory.
ignore	logical whether to ignore the path or file in git and R build.

**Value**

invisibly returns TRUE for success. Stops on failure.

**Examples**

```
if(rmarkdown::pandoc_available()){
  myfile <- tempfile()
  file <- system.file("extdata", "tests", "extra.rmd",
                     package = "DataPackageR")
  raw_data <- system.file("extdata", "tests", "raw_data",
                        package = "DataPackageR")
  datapackage_skeleton(
    name = "datatest",
    path = tempdir(),
    code_files = file,
    force = TRUE,
    r_object_names = "data")
  use_raw_dataset(raw_data)
}
```

---

yml\_find

*Edit DataPackageR yml configuration*

---

**Description**

Edit a yml configuration file via an API.

**Usage**

```
yml_find(path)

yml_add_files(config, filenames)

yml_disable_compile(config, filenames)

yml_enable_compile(config, filenames)

yml_add_objects(config, objects)

yml_list_objects(config)

yml_list_files(config)

yml_remove_objects(config, objects)

yml_remove_files(config, filenames)

yml_write(config, path = NULL)
```

**Arguments**

path	Path to the data package source or path to write config file (for yml_write)
config	an R representation of the datapackager.yml config, returned by yml_find, or a path to the package root.
filenames	A vector of filenames.
objects	A vector of R object names.

**Details**

Add, remove files and objects, enable or disable parsing of specific files, list objects or files in a yaml config, or write a config back to a package.

**Value**

A yaml configuration structured as an R nested list.

**Examples**

```

if(rmarkdown::pandoc_available()){
  f <- tempdir()
  f <- file.path(f,"foo.Rmd")
  con <- file(f)
  writeLines("```{r}\n vec = 1:10\n```\n",con=con)
  close(con)
  pname <- basename(tempfile())
  datapackage_skeleton(name=pname,
    path = tempdir(),
    force = TRUE,
    r_object_names = "vec",
    code_files = f)
  yml <- yml_find(file.path(tempdir(),pname))
  yml <- yml_add_files(yml,"foo.Rmd")
  yml_list_files(yml)
  yml <- yml_disable_compile(yml,"foo.Rmd")
  yml <- yml_enable_compile(yml,"foo.Rmd")
  yml <- yml_add_objects(yml,"data1")
  yml_list_objects(yml)
  yml <- yml_remove_objects(yml,"data1")
  yml <- yml_remove_files(yml,"foo.Rmd")
}

```

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