

# Package ‘feather’

September 15, 2019

**Title** R Bindings to the Feather 'API'  
**Version** 0.3.5  
**Description** Read and write feather files, a lightweight binary columnar data store designed for maximum speed.  
**Encoding** UTF-8  
**License** Apache License 2.0  
**LazyData** true  
**URL** <https://github.com/wesm/feather>  
**BugReports** <https://github.com/wesm/feather/issues>  
**LinkingTo** Rcpp  
**Imports** Rcpp, tibble (>= 2.0.0), hms  
**Suggests** testthat  
**SystemRequirements** little-endian platform  
**RoxygenNote** 6.1.1  
**NeedsCompilation** yes  
**Author** Hadley Wickham [aut, cre],  
RStudio [cph],  
Feather developers [ctb] (Bundled feather library),  
Google [ctb] (Bundled flatbuffers code),  
LevelDB Authors [ctb]  
**Maintainer** Hadley Wickham <hadley@rstudio.com>  
**Repository** CRAN  
**Date/Publication** 2019-09-15 00:10:22 UTC

## R topics documented:

feather . . . . .	2
feather_metadata . . . . .	2
read_feather . . . . .	3
<b>Index</b>	<b>4</b>

---

feather	<i>Access a feather store like a data frame</i>
---------	---

---

**Description**

These functions permit using a feather dataset much like a regular (read-only) data frame.

**Usage**

```
feather(path)
```

**Arguments**

path	Path to feather file
------	----------------------

**Value**

An object of class feather

---

feather_metadata	<i>Retrieve metadata about a feather file</i>
------------------	---

---

**Description**

Returns the dimensions, field names, and types; and optional dataset description.

**Usage**

```
feather_metadata(path)
```

**Arguments**

path	Path to feather file
------	----------------------

**Value**

A list with class "feather\_metadata".

---

read_feather	<i>Read and write feather files.</i>
--------------	--------------------------------------

---

**Description**

Read and write feather files.

**Usage**

```
read_feather(path, columns = NULL)
```

```
write_feather(x, path)
```

**Arguments**

path	Path to feather file
columns	Columns to read (names or indexes). Default: Read all columns.
x	A data frame to write to disk

**Value**

Both functions return a tibble/data frame. `write_feather` invisibly returns `x` (so you can use this function in a pipeline).

**Examples**

```
mtcars2 <- read_feather(feather_example("mtcars.feather"))  
mtcars2
```

# Index

`feather`, [2](#)

`feather_metadata`, [2](#)

`read_feather`, [3](#)

`write_feather` (`read_feather`), [3](#)