# Package 'arrow'

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Title Integration to 'Apache' 'Arrow'

Version 1.0.0

**Description** 'Apache' 'Arrow' <a href="https://arrow.apache.org/">https://arrow.apache.org/</a> is a cross-language development platform for in-memory data. It specifies a standardized language-independent columnar memory format for flat and hierarchical data, organized for efficient analytic operations on modern hardware. This package provides an interface to the 'Arrow C++' library.

**Depends** R (>= 3.1)

License Apache License (>= 2.0)

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BugReports https://issues.apache.org/jira/projects/ARROW/issues

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# NeedsCompilation yes

Author Romain François [aut] (<https://orcid.org/0000-0002-2444-4226>), Jeroen Ooms [aut], Neal Richardson [aut, cre], Javier Luraschi [ctb], Jeffrey Wong [ctb], Apache Arrow [aut, cph]

Maintainer Neal Richardson <neal@ursalabs.org>

**Repository** CRAN

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array
```

Arrow Arrays

# Description

An Array is an immutable data array with some logical type and some length. Most logical types are contained in the base Array class; there are also subclasses for DictionaryArray, ListArray, and StructArray.

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#### Factory

The Array\$create() factory method instantiates an Array and takes the following arguments:

- x: an R vector, list, or data.frame
- type: an optional data type for x. If omitted, the type will be inferred from the data.

Array\$create() will return the appropriate subclass of Array, such as DictionaryArray when given an R factor.

To compose a DictionaryArray directly, call DictionaryArray\$create(), which takes two arguments:

- x: an R vector or Array of integers for the dictionary indices
- dict: an R vector or Array of dictionary values (like R factor levels but not limited to strings only)

#### Usage

```
a <- Array$create(x)
length(a)
```

print(a) a == a

#### Methods

- \$IsNull(i): Return true if value at index is null. Does not boundscheck
- \$IsValid(i): Return true if value at index is valid. Does not boundscheck
- \$length(): Size in the number of elements this array contains
- \$offset(): A relative position into another array's data, to enable zero-copy slicing
- \$null\_count(): The number of null entries in the array
- \$type(): logical type of data
- \$type\_id(): type id
- \$Equals(other) : is this array equal to other
- \$ApproxEquals(other) :
- \$data(): return the underlying ArrayData
- \$as\_vector(): convert to an R vector
- \$ToString(): string representation of the array
- \$Slice(offset, length = NULL): Construct a zero-copy slice of the array with the indicated offset and length. If length is NULL, the slice goes until the end of the array.
- \$Take(i): return an Array with values at positions given by integers (R vector or Array Array) i.
- \$Filter(i, keep\_na = TRUE): return an Array with values at positions where logical vector (or Arrow boolean Array) i is TRUE.
- \$RangeEquals(other, start\_idx, end\_idx, other\_start\_idx) :
- \$cast(target\_type, safe = TRUE, options = cast\_options(safe)): Alter the data in the array to change its type.

# ArrayData

- \$View(type): Construct a zero-copy view of this array with the given type.
- \$Validate() : Perform any validation checks to determine obvious inconsistencies within the array's internal data. This can be an expensive check, potentially 0(length)

ArrayData ArrayData class

#### Description

The ArrayData class allows you to get and inspect the data inside an arrow: : Array.

#### Usage

```
data <- Array$create(x)$data()
data$type()
data$length()
data$null_count()
data$offset()
data$buffers()</pre>
```

#### Methods

•••

arrow\_available Is the C++ Arrow library available?

#### Description

You won't generally need to call this function, but it's here in case it helps for development purposes.

#### Usage

```
arrow_available()
```

#### Value

TRUE or FALSE depending on whether the package was installed with the Arrow C++ library. If FALSE, you'll need to install the C++ library and then reinstall the R package. See install\_arrow() for help.

#### Examples

arrow\_available()

buffer

# Description

A Buffer is an object containing a pointer to a piece of contiguous memory with a particular size.

# Usage

buffer(x)

# Arguments

х

R object. Only raw, numeric and integer vectors are currently supported

# Value

an instance of Buffer that borrows memory from x

# Factory

buffer() lets you create an arrow: :Buffer from an R object

# Methods

- \$is\_mutable():
- \$ZeroPadding():
- \$size():
- \$capacity():

cast\_options Cast options

# Description

Cast options

# Usage

```
cast_options(
  safe = TRUE,
  allow_int_overflow = !safe,
  allow_time_truncate = !safe,
  allow_float_truncate = !safe
)
```

#### ChunkedArray

#### Arguments

ChunkedArray ChunkedArray class

#### Description

A ChunkedArray is a data structure managing a list of primitive Arrow Arrays logically as one large array. Chunked arrays may be grouped together in a Table.

# Usage

chunked\_array(..., type = NULL)

# Arguments

• • •	Vectors to coerce
type	currently ignored

# Factory

The ChunkedArray\$create() factory method instantiates the object from various Arrays or R vectors. chunked\_array() is an alias for it.

#### Methods

- \$length(): Size in the number of elements this array contains
- \$chunk(i): Extract an Array chunk by integer position
- \$as\_vector(): convert to an R vector
- \$Slice(offset, length = NULL): Construct a zero-copy slice of the array with the indicated offset and length. If length is NULL, the slice goes until the end of the array.
- \$Take(i): return a ChunkedArray with values at positions given by integers i. If i is an Arrow Array or ChunkedArray, it will be coerced to an R vector before taking.
- \$Filter(i, keep\_na = TRUE): return a ChunkedArray with values at positions where logical vector or Arrow boolean-type (Chunked)Array i is TRUE.
- \$cast(target\_type, safe = TRUE, options = cast\_options(safe)): Alter the data in the array to change its type.
- \$null\_count(): The number of null entries in the array

- \$chunks(): return a list of Arrays
- \$num\_chunks(): integer number of chunks in the ChunkedArray
- \$type(): logical type of data
- \$View(type): Construct a zero-copy view of this ChunkedArray with the given type.
- \$Validate(): Perform any validation checks to determine obvious inconsistencies within the array's internal data. This can be an expensive check, potentially 0(length)

#### See Also

Array

Codec

Compression Codec class

#### Description

Codecs allow you to create compressed input and output streams.

#### Factory

The Codec\$create() factory method takes the following arguments:

- type: string name of the compression method. Possible values are "uncompressed", "snappy", "gzip", "brotli", "zstd", "lz4", "lzo", or "bz2". type may be upper- or lower-cased. Not all methods may be available; support depends on build-time flags for the C++ library. See codec\_is\_available(). Most builds support at least "snappy" and "gzip". All support "uncompressed".
- compression\_level: compression level, the default value (NA) uses the default compression level for the selected compression type.

codec\_is\_available Check whether a compression codec is available

# Description

Support for compression libraries depends on the build-time settings of the Arrow C++ library. This function lets you know which are available for use.

#### Usage

```
codec_is_available(type)
```

#### Arguments

type

A string, one of "uncompressed", "snappy", "gzip", "brotli", "zstd", "lz4", "lzo", or "bz2", case insensitive.

#### compression

# Value

Logical: is type available?

compression

Compressed stream classes

# Description

CompressedInputStream and CompressedOutputStream allow you to apply a compression Codec to an input or output stream.

# Factory

The CompressedInputStream\$create() and CompressedOutputStream\$create() factory methods instantiate the object and take the following arguments:

- stream An InputStream or OutputStream, respectively
- codec A Codec, either a Codec instance or a string
- compression\_level compression level for when the codec argument is given as a string

#### Methods

Methods are inherited from InputStream and OutputStream, respectively

cpu\_count

Manage the global CPU thread pool in libarrow

#### Description

Manage the global CPU thread pool in libarrow

#### Usage

```
cpu_count()
```

```
set_cpu_count(num_threads)
```

#### Arguments

num\_threads integer: New number of threads for thread pool

CsvReadOptions

#### Description

CsvReadOptions, CsvParseOptions, CsvConvertOptions, JsonReadOptions, and JsonParseOptions are containers for various file reading options. See their usage in read\_csv\_arrow() and read\_json\_arrow(), respectively.

#### Factory

The CsvReadOptions\$create() and JsonReadOptions\$create() factory methods take the following arguments:

- use\_threads Whether to use the global CPU thread pool
- block\_size Block size we request from the IO layer; also determines the size of chunks when use\_threads is TRUE. NB: if FALSE, JSON input must end with an empty line.

CsvReadOptions\$create() further accepts these additional arguments:

- skip\_rows Number of lines to skip before reading data (default 0)
- column\_names Character vector to supply column names. If length-0 (the default), the first non-skipped row will be parsed to generate column names, unless autogenerate\_column\_names is TRUE.
- autogenerate\_column\_names Logical: generate column names instead of using the first nonskipped row (the default)? If TRUE, column names will be "f0", "f1", ..., "fN".

CsvParseOptions\$create() takes the following arguments:

- delimiter Field delimiting character (default ", ")
- quoting Logical: are strings quoted? (default TRUE)
- quote\_char Quoting character, if quoting is TRUE
- double\_quote Logical: are quotes inside values double-quoted? (default TRUE)
- escaping Logical: whether escaping is used (default FALSE)
- escape\_char Escaping character, if escaping is TRUE
- newlines\_in\_values Logical: are values allowed to contain CR (0x0d) and LF (0x0a) characters? (default FALSE)
- ignore\_empty\_lines Logical: should empty lines be ignored (default) or generate a row of missing values (if FALSE)?

JsonParseOptions\$create() accepts only the newlines\_in\_values argument.

CsvConvertOptions\$create() takes the following arguments:

- check\_utf8 Logical: check UTF8 validity of string columns? (default TRUE)
- null\_values character vector of recognized spellings for null values. Analogous to the na.strings argument to read.csv() or na in readr::read\_csv().
- strings\_can\_be\_null Logical: can string / binary columns have null values? Similar to the quoted\_na argument to readr::read\_csv(). (default FALSE)

#### CsvTableReader

#### Methods

These classes have no implemented methods. They are containers for the options.

CsvTableReader Arrow CSV and JSON table reader classes

#### Description

CsvTableReader and JsonTableReader wrap the Arrow C++ CSV and JSON table readers. See their usage in read\_csv\_arrow() and read\_json\_arrow(), respectively.

## Factory

The CsvTableReader\$create() and JsonTableReader\$create() factory methods take the following arguments:

- file A character path to a local file, or an Arrow input stream
- convert\_options (CSV only), parse\_options, read\_options: see CsvReadOptions
- ... additional parameters.

#### Methods

• \$Read(): returns an Arrow Table.

data-type Apache Arrow data types

#### Description

These functions create type objects corresponding to Arrow types. Use them when defining a schema() or as inputs to other types, like struct. Most of these functions don't take arguments, but a few do.

# Usage

int8()
int16()
int32()
int64()
uint8()

uint16()

uint32()

uint64()

float16()

halffloat()

float32()

float()

float64()

boolean()

bool()

utf8()

large\_utf8()

binary()

large\_binary()

fixed\_size\_binary(byte\_width)

string()

date32()

date64()

time32(unit = c("ms", "s"))

time64(unit = c("ns", "us"))

null()

timestamp(unit = c("s", "ms", "us", "ns"), timezone = "")

decimal(precision, scale)

list\_of(type)

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# data-type

large\_list\_of(type)

fixed\_size\_list\_of(type, list\_size)

struct(...)

# Arguments

byte_width	byte width for FixedSizeBinary type.
unit	For time/timestamp types, the time unit. time32() can take either "s" or "ms", while time64() can be "us" or "ns". timestamp() can take any of those four values.
timezone	For timestamp(), an optional time zone string.
precision	For decimal(), precision
scale	For decimal(), scale
type	For list_of(), a data type to make a list-of-type
list_size	list size for FixedSizeList type.
	For struct(), a named list of types to define the struct columns

# Details

A few functions have aliases:

- utf8() and string()
- float16() and halffloat()
- float32() and float()
- bool() and boolean()
- Called from schema() or struct(), double() also is supported as a way of creating a float64()

date32() creates a datetime type with a "day" unit, like the R Date class. date64() has a "ms" unit.

#### Value

An Arrow type object inheriting from DataType.

# See Also

dictionary() for creating a dictionary (factor-like) type.

# Examples

```
bool()
struct(a = int32(), b = double())
timestamp("ms", timezone = "CEST")
time64("ns")
```

Dataset

#### Description

Arrow Datasets allow you to query against data that has been split across multiple files. This sharding of data may indicate partitioning, which can accelerate queries that only touch some partitions (files).

A Dataset contains one or more Fragments, such as files, of potentially differing type and partitioning.

For Dataset\$create(), see open\_dataset(), which is an alias for it.

DatasetFactory is used to provide finer control over the creation of Datasets.

#### Factory

DatasetFactory is used to create a Dataset, inspect the Schema of the fragments contained in it, and declare a partitioning. FileSystemDatasetFactory is a subclass of DatasetFactory for discovering files in the local file system, the only currently supported file system.

For the DatasetFactory\$create() factory method, see dataset\_factory(), an alias for it. A DatasetFactory has:

- \$Inspect(unify\_schemas): If unify\_schemas is TRUE, all fragments will be scanned and a unified Schema will be created from them; if FALSE (default), only the first fragment will be inspected for its schema. Use this fast path when you know and trust that all fragments have an identical schema.
- \$Finish(schema, unify\_schemas): Returns a Dataset. If schema is provided, it will be used for the Dataset; if omitted, a Schema will be created from inspecting the fragments (files) in the dataset, following unify\_schemas as described above.

FileSystemDatasetFactory\$create() is a lower-level factory method and takes the following arguments:

- filesystem: A FileSystem
- selector: A FileSelector
- format: A FileFormat
- partitioning: Either Partitioning, PartitioningFactory, or NULL

#### Methods

A Dataset has the following methods:

- \$NewScan(): Returns a ScannerBuilder for building a query
- \$schema: Active binding that returns the Schema of the Dataset; you may also replace the dataset's schema by using ds\$schema <-new\_schema. This method currently supports only adding, removing, or reordering fields in the schema: you cannot alter or cast the field types.</li>

FileSystemDataset has the following methods:

# dataset\_factory

- \$files: Active binding, returns the files of the FileSystemDataset
- \$format: Active binding, returns the FileFormat of the FileSystemDataset

UnionDataset has the following methods:

• \$children: Active binding, returns all child Datasets.

#### See Also

open\_dataset() for a simple interface to creating a Dataset

dataset\_factory Create a DatasetFactory

# Description

A Dataset can constructed using one or more DatasetFactorys. This function helps you construct a DatasetFactory that you can pass to open\_dataset().

# Usage

```
dataset_factory(
    x,
    filesystem = NULL,
    format = c("parquet", "arrow", "ipc", "feather", "csv", "tsv", "text"),
    partitioning = NULL,
    ...
)
```

### Arguments

x	A string file x containing data files, or a list of DatasetFactory objects whose datasets should be grouped. If this argument is specified it will be used to construct a UnionDatasetFactory and other arguments will be ignored.
filesystem	A FileSystem object; if omitted, the FileSystem will be detected from x
format	A FileFormat object, or a string identifier of the format of the files in x. Currently supported values:
	• "parquet"
	• "ipc"/"arrow"/"feather", all aliases for each other; for Feather, note that only version 2 files are supported
	• "csv"/"text", aliases for the same thing (because comma is the default de- limiter for text files
	<ul> <li>"tsv", equivalent to passing format = "text", delimiter = "\t"</li> </ul>
	Default is "parquet", unless a delimiter is also specified, in which case it is assumed to be "text".
partitioning	One of

- A Schema, in which case the file paths relative to sources will be parsed, and path segments will be matched with the schema fields. For example, schema(year = int16(),month = int8()) would create partitions for file paths like "2019/01/file.parquet", "2019/02/file.parquet", etc.
- A character vector that defines the field names corresponding to those path segments (that is, you're providing the names that would correspond to a Schema but the types will be autodetected)
- A HivePartitioning or HivePartitioningFactory, as returned by hive\_partition() which parses explicit or autodetected fields from Hive-style path segments
- NULL for no partitioning

Additional format-specific options, passed to FileFormat\$create(). For CSV options, note that you can specify them either with the Arrow C++ library naming ("delimiter", "quoting", etc.) or the readr-style naming used in read\_csv\_arrow() ("delim", "quote", etc.)

#### Details

. . .

If you would only have a single DatasetFactory (for example, you have a single directory containing Parquet files), you can call open\_dataset() directly. Use dataset\_factory() when you want to combine different directories, file systems, or file formats.

#### Value

A DatasetFactory object. Pass this to open\_dataset(), in a list potentially with other DatasetFactory objects, to create a Dataset.

DataType

class arrow::DataType

#### Description

class arrow::DataType

#### Methods

TODO

# Description

default arrow::MemoryPool

# Usage

default\_memory\_pool()

# Value

the default arrow::MemoryPool

dictionary

Create a dictionary type

# Description

Create a dictionary type

# Usage

```
dictionary(index_type = int32(), value_type = utf8(), ordered = FALSE)
```

# Arguments

index_type	A DataType for the indices (default int32())
value_type	A DataType for the values (default utf8())
ordered	Is this an ordered dictionary (default FALSE)?

# Value

A DictionaryType

# See Also

Other Arrow data types

DictionaryType cla

# class DictionaryType

#### Description

class DictionaryType

# Methods

TODO

Expression

#### Description

Expressions are used to define filter logic for passing to a Dataset Scanner.

Arrow expressions

Expression\$scalar(x) constructs an Expression which always evaluates to the provided scalar (length-1) R value.

Expression\$field\_ref(name) is used to construct an Expression which evaluates to the named column in the Dataset against which it is evaluated.

Expression\$compare(OP,e1,e2) takes two Expression operands, constructing an Expression which will evaluate these operands then compare them with the relation specified by OP (e.g. "==", "!=", ">", etc.) For example, to filter down to rows where the column named "alpha" is less than 5: Expression\$compare("<",Expression\$field\_ref("alpha"),Expression\$scalar(5))

Expression\$and(e1,e2), Expression\$or(e1,e2), and Expression\$not(e1) construct an Expression combining their arguments with Boolean operators.

Expression\$is\_valid(x) is essentially (an inversion of) is.na() for Expressions.

Expression\$in\_(x, set) evaluates x and returns whether or not it is a member of the set.

FeatherReader FeatherReader class

# Description

This class enables you to interact with Feather files. Create one to connect to a file or other Input-Stream, and call Read() on it to make an arrow: :Table. See its usage in read\_feather().

# Field

# Factory

The FeatherReader\$create() factory method instantiates the object and takes the following arguments:

- file an Arrow file connection object inheriting from RandomAccessFile.
- mmap Logical: whether to memory-map the file (default TRUE)
- ... Additional arguments, currently ignored

# Methods

- \$Read(columns): Returns a Table of the selected columns, a vector of integer indices
- \$version: Active binding, returns 1 or 2, according to the Feather file version

Field

Field class

# Description

field() lets you create an arrow::Field that maps a DataType to a column name. Fields are contained in Schemas.

#### Usage

field(name, type, metadata)

# Arguments

name	field name
type	logical type, instance of DataType
metadata	currently ignored

#### Methods

- f\$ToString(): convert to a string
- f\$Equals(other): test for equality. More naturally called as f == other

# Examples

field("x", int32())

FileFormat

#### Description

A FileFormat holds information about how to read and parse the files included in a Dataset. There are subclasses corresponding to the supported file formats (ParquetFileFormat and IpcFileFormat).

#### Factory

FileFormat\$create() takes the following arguments:

- format: A string identifier of the file format. Currently supported values:
  - "parquet"
  - "ipc"/"arrow"/"feather", all aliases for each other; for Feather, note that only version 2 files are supported
  - "csv"/"text", aliases for the same thing (because comma is the default delimiter for text files
  - "tsv", equivalent to passing format = "text", delimiter = "\t"
- ...: Additional format-specific options

'format = "parquet"":

- use\_buffered\_stream: Read files through buffered input streams rather than loading entire row groups at once. This may be enabled to reduce memory overhead. Disabled by default.
- buffer\_size: Size of buffered stream, if enabled. Default is 8KB.
- dict\_columns: Names of columns which should be read as dictionaries.

format = "text": see CsvReadOptions. Note that you can specify them either with the Arrow C++ library naming ("delimiter", "quoting", etc.) or the readr-style naming used in read\_csv\_arrow() ("delim", "quote", etc.)

It returns the appropriate subclass of FileFormat (e.g. ParquetFileFormat)

FileInfo

FileSystem entry info

#### Description

FileSystem entry info

# Methods

- base\_name() : The file base name (component after the last directory separator).
- extension() : The file extension

# FileSelector

#### Active bindings

- \$type: The file type
- \$path: The full file path in the filesystem
- \$size: The size in bytes, if available. Only regular files are guaranteed to have a size.
- \$mtime: The time of last modification, if available.

FileSelector *file selector* 

# Description

file selector

# Factory

The \$create() factory method instantiates a FileSelector given the 3 fields described below.

# Fields

- base\_dir: The directory in which to select files. If the path exists but doesn't point to a directory, this should be an error.
- allow\_not\_found: The behavior if base\_dir doesn't exist in the filesystem. If FALSE, an error is returned. If TRUE, an empty selection is returned
- recursive: Whether to recurse into subdirectories.

FileSystem

FileSystem classes

#### Description

FileSystem is an abstract file system API, LocalFileSystem is an implementation accessing files on the local machine. SubTreeFileSystem is an implementation that delegates to another implementation after prepending a fixed base path

#### Factory

The \$create() factory methods instantiate the FileSystem object and take the following arguments, depending on the subclass:

- no argument is needed for instantiating a LocalFileSystem
- base\_path and base\_fs for instantiating a SubTreeFileSystem

#### Methods

- \$GetFileInfo(x): x may be a FileSelector or a character vector of paths. Returns a list of FileInfo
- \$CreateDir(path, recursive = TRUE): Create a directory and subdirectories.
- \$DeleteDir(path): Delete a directory and its contents, recursively.
- \$DeleteDirContents(path): Delete a directory's contents, recursively. Like \$DeleteDir(), but doesn't delete the directory itself. Passing an empty path ("") will wipe the entire filesystem tree.
- \$DeleteFile(path) : Delete a file.
- \$DeleteFiles(paths) : Delete many files. The default implementation issues individual delete operations in sequence.
- \$Move(src, dest): Move / rename a file or directory. If the destination exists: if it is a nonempty directory, an error is returned otherwise, if it has the same type as the source, it is replaced otherwise, behavior is unspecified (implementation-dependent).
- \$CopyFile(src, dest): Copy a file. If the destination exists and is a directory, an error is returned. Otherwise, it is replaced.
- \$OpenInputStream(path): Open an input stream for sequential reading.
- \$OpenInputFile(path): Open an input file for random access reading.
- \$OpenOutputStream(path): Open an output stream for sequential writing.
- \$OpenAppendStream(path): Open an output stream for appending.

FixedWidthType class arrow::FixedWidthType

# Description

class arrow::FixedWidthType

#### Methods

TODO

hive\_partition Construct Hive partitioning

#### Description

Hive partitioning embeds field names and values in path segments, such as "/year=2019/month=2/data.parquet".

#### Usage

hive\_partition(...)

#### InputStream

# Arguments

named list of data types, passed to schema()

# Details

Because fields are named in the path segments, order of fields passed to hive\_partition() does not matter.

#### Value

A HivePartitioning, or a HivePartitioningFactory if calling hive\_partition() with no arguments.

#### Examples

hive\_partition(year = int16(), month = int8())

InputStream InputStream classes

#### Description

RandomAccessFile inherits from InputStream and is a base class for: ReadableFile for reading from a file; MemoryMappedFile for the same but with memory mapping; and BufferReader for reading from a buffer. Use these with the various table readers.

#### Factory

The \$create() factory methods instantiate the InputStream object and take the following arguments, depending on the subclass:

- path For ReadableFile, a character file name
- x For BufferReader, a Buffer or an object that can be made into a buffer via buffer().

To instantiate a MemoryMappedFile, call mmap\_open().

#### Methods

- \$GetSize():
- \$supports\_zero\_copy(): Logical
- \$seek(position): go to that position in the stream
- \$tell(): return the position in the stream
- \$close(): close the stream
- \$Read(nbytes): read data from the stream, either a specified nbytes or all, if nbytes is not provided
- \$ReadAt(position, nbytes): similar to \$seek(position)\$Read(nbytes)
- \$Resize(size): for a MemoryMappedFile that is writeable

```
install_arrow
```

# Description

Use this function to install the latest release of arrow, to switch to or from a nightly development version, or on Linux to try reinstalling with all necessary C++ dependencies.

# Usage

```
install_arrow(
    nightly = FALSE,
    binary = Sys.getenv("LIBARROW_BINARY", TRUE),
    use_system = Sys.getenv("ARROW_USE_PKG_CONFIG", FALSE),
    minimal = Sys.getenv("LIBARROW_MINIMAL", FALSE),
    repos = getOption("repos"),
    ...
)
```

# Arguments

nightly	logical: Should we install a development version of the package, or should we install from CRAN (the default).
binary	On Linux, value to set for the environment variable LIBARROW_BINARY, which governs how C++ binaries are used, if at all. The default value, TRUE, tells the installation script to detect the Linux distribution and version and find an appropriate C++ library. FALSE would tell the script not to retrieve a binary and instead build Arrow C++ from source. Other valid values are strings corresponding to a Linux distribution-version, to override the value that would be detected. See vignette("install", package = "arrow") for further details.
use_system	logical: Should we use pkg-config to look for Arrow system packages? Default is FALSE. If TRUE, source installation may be faster, but there is a risk of version mismatch.
minimal	logical: If building from source, should we build without optional dependencies (compression libraries, for example)? Default is FALSE.
repos	character vector of base URLs of the repositories to install from (passed to install.packages())
	Additional arguments passed to install.packages()

# See Also

arrow\_available() to see if the package was configured with necessary C++ dependencies. vignette("install", packag = "arrow") for more ways to tune installation on Linux. install\_pyarrow Install pyarrow for use with reticulate

#### Description

pyarrow is the Python package for Apache Arrow. This function helps with installing it for use with reticulate.

#### Usage

install\_pyarrow(envname = NULL, nightly = FALSE, ...)

# Arguments

envname	The name or full path of the Python environment to install into. This can be a vir- tualenv or conda environment created by reticulate. See reticulate::py_install().
nightly	logical: Should we install a development version of the package? Default is to use the official release version.
	additional arguments passed to reticulate::py_install().
map_batches	Apply a function to a stream of RecordBatches

#### Description

As an alternative to calling collect() on a Dataset query, you can use this function to access the stream of RecordBatches in the Dataset. This lets you aggregate on each chunk and pull the intermediate results into a data.frame for further aggregation, even if you couldn't fit the whole Dataset result in memory.

#### Usage

map\_batches(X, FUN, ..., .data.frame = TRUE)

#### Arguments

Х	A Dataset or arrow_dplyr_query object, as returned by the dplyr methods on Dataset.
FUN	A function or purrr-style lambda expression to apply to each batch
•••	Additional arguments passed to FUN
.data.frame	logical: collect the resulting chunks into a single data.frame? Default TRUE

# Details

This is experimental and not recommended for production use.

MemoryPool

# Description

class arrow::MemoryPool

# Methods

TODO

Message

# class arrow::Message

# Description

class arrow::Message

# Methods

TODO

MessageReader class arrow::MessageReader

# Description

class arrow::MessageReader

# Methods

TODO

mmap\_create

# Description

Create a new read/write memory mapped file of a given size

# Usage

```
mmap_create(path, size)
```

# Arguments

path	file path
size	size in bytes

# Value

a arrow::io::MemoryMappedFile

mmap\_open

Open a memory mapped file

# Description

Open a memory mapped file

# Usage

```
mmap_open(path, mode = c("read", "write", "readwrite"))
```

# Arguments

path	file path
mode	file mode (read/write/readwrite)

open\_dataset

# Description

Arrow Datasets allow you to query against data that has been split across multiple files. This sharding of data may indicate partitioning, which can accelerate queries that only touch some partitions (files). Call open\_dataset() to point to a directory of data files and return a Dataset, then use dplyr methods to query it.

# Usage

```
open_dataset(
   sources,
   schema = NULL,
   partitioning = hive_partition(),
   unify_schemas = NULL,
   ...
)
```

```
Arguments
```

sources	Either:
	• a string path to a directory containing data files
	<ul> <li>a list of Dataset objects as created by this function</li> </ul>
	<ul> <li>a list of DatasetFactory objects as created by dataset_factory().</li> </ul>
schema	Schema for the dataset. If NULL (the default), the schema will be inferred from the data sources.
partitioning	When sources is a file path, one of
	<ul> <li>a Schema, in which case the file paths relative to sources will be parsed, and path segments will be matched with the schema fields. For example, schema(year = int16(),month = int8()) would create partitions for file paths like "2019/01/file.parquet", "2019/02/file.parquet", etc.</li> </ul>
	• a character vector that defines the field names corresponding to those path segments (that is, you're providing the names that would correspond to a Schema but the types will be autodetected)
	• a HivePartitioning or HivePartitioningFactory, as returned by hive_partition() which parses explicit or autodetected fields from Hive-style path segments
	NULL for no partitioning
	The default is to autodetect Hive-style partitions.
unify_schemas	logical: should all data fragments (files, Datasets) be scanned in order to create a unified schema from them? If FALSE, only the first fragment will be inspected for its schema. Use this fast path when you know and trust that all fragments have an identical schema. The default is FALSE when creating a dataset from a

# OutputStream

file path (because there may be many files and scanning may be slow) but TRUE when sources is a list of Datasets (because there should be few Datasets in the list and their Schemas are already in memory).
 additional arguments passed to dataset_factory() when sources is a file path, otherwise ignored. These may include "format" to indicate the file format, or other format-specific options.

# Value

A Dataset R6 object. Use dplyr methods on it to query the data, or call \$NewScan() to construct a query directly.

# See Also

vignette("dataset",package = "arrow")

OutputStream

# OutputStream classes

#### Description

FileOutputStream is for writing to a file; BufferOutputStream writes to a buffer; You can create one and pass it to any of the table writers, for example.

# Factory

The \$create() factory methods instantiate the OutputStream object and take the following arguments, depending on the subclass:

- path For FileOutputStream, a character file name
- initial\_capacity For BufferOutputStream, the size in bytes of the buffer.

#### Methods

- \$tell(): return the position in the stream
- \$close(): close the stream
- \$write(x): send x to the stream
- \$capacity(): for BufferOutputStream
- \$finish(): for BufferOutputStream
- \$GetExtentBytesWritten(): for MockOutputStream, report how many bytes were sent.

ParquetFileReader ParquetFileReader class

#### Description

This class enables you to interact with Parquet files.

# Factory

The ParquetFileReader\$create() factory method instantiates the object and takes the following arguments:

- file A character file name, raw vector, or Arrow file connection object (e.g. RandomAccessFile).
- props Optional ParquetReaderProperties
- mmap Logical: whether to memory-map the file (default TRUE)
- ... Additional arguments, currently ignored

# Methods

- \$ReadTable(col\_select): get an arrow::Table from the file, possibly with columns filtered by a character vector of column names or a tidyselect specification.
- \$GetSchema(): get the arrow::Schema of the data in the file

# Examples

```
f <- system.file("v0.7.1.parquet", package="arrow")
pq <- ParquetFileReader$create(f)
pq$GetSchema()
if (codec_is_available("snappy")) {
    # This file has compressed data columns
    tab <- pq$ReadTable(starts_with("c"))
    tab$schema
}</pre>
```

ParquetFileWriter ParquetFileWriter class

#### Description

This class enables you to interact with Parquet files.

# Factory

The ParquetFileWriter\$create() factory method instantiates the object and takes the following arguments:

- schema A Schema
- sink An arrow::io::OutputStream or a string which is interpreted as a file path
- properties An instance of ParquetWriterProperties
- arrow\_properties An instance of ParquetArrowWriterProperties

ParquetReaderProperties

ParquetReaderProperties class

# Description

This class holds settings to control how a Parquet file is read by ParquetFileReader.

#### Factory

The ParquetReaderProperties\$create() factory method instantiates the object and takes the following arguments:

• use\_threads Logical: whether to use multithreading (default TRUE)

#### Methods

- \$read\_dictionary(column\_index)
- \$set\_read\_dictionary(column\_index, read\_dict)
- \$use\_threads(use\_threads)

ParquetWriterProperties

ParquetWriterProperties class

# Description

This class holds settings to control how a Parquet file is read by ParquetFileWriter.

#### Details

The parameters compression, compression\_level, use\_dictionary and write\_statistics' support various patterns:

- The default NULL leaves the parameter unspecified, and the C++ library uses an appropriate default for each column (defaults listed above)
- A single, unnamed, value (e.g. a single string for compression) applies to all columns
- An unnamed vector, of the same size as the number of columns, to specify a value for each column, in positional order
- A named vector, to specify the value for the named columns, the default value for the setting is used when not supplied

Unlike the high-level write\_parquet, ParquetWriterProperties arguments use the C++ defaults. Currently this means "uncompressed" rather than "snappy" for the compression argument.

# Factory

The ParquetWriterProperties\$create() factory method instantiates the object and takes the following arguments:

- table: table to write (required)
- version: Parquet version, "1.0" or "2.0". Default "1.0"
- compression: Compression type, algorithm "uncompressed"
- compression\_level: Compression level; meaning depends on compression algorithm
- use\_dictionary: Specify if we should use dictionary encoding. Default TRUE
- write\_statistics: Specify if we should write statistics. Default TRUE
- data\_page\_size: Set a target threshold for the approximate encoded size of data pages within a column chunk (in bytes). Default 1 MiB.

# See Also

write\_parquet

Partitioning

Define Partitioning for a Dataset

#### Description

Pass a Partitioning object to a FileSystemDatasetFactory's \$create() method to indicate how the file's paths should be interpreted to define partitioning.

DirectoryPartitioning describes how to interpret raw path segments, in order. For example, schema(year = int16(),month = int8()) would define partitions for file paths like "2019/01/file.parquet", "2019/02/file.parquet", etc.

#### read\_arrow

HivePartitioning is for Hive-style partitioning, which embeds field names and values in path segments, such as "/year=2019/month=2/data.parquet". Because fields are named in the path segments, order does not matter.

PartitioningFactory subclasses instruct the DatasetFactory to detect partition features from the file paths.

#### Factory

Both DirectoryPartitioning\$create() and HivePartitioning\$create() methods take a Schema as a single input argument. The helper function hive\_partition(...) is shorthand for HivePartitioning\$create(schema

With DirectoryPartitioningFactory\$create(), you can provide just the names of the path segments (in our example, c("year", "month")), and the DatasetFactory will infer the data types for those partition variables. HivePartitioningFactory\$create() takes no arguments: both variable names and their types can be inferred from the file paths. hive\_partition() with no arguments returns a HivePartitioningFactory.

read\_arrow

Read Arrow IPC stream format

#### Description

Apache Arrow defines two formats for serializing data for interprocess communication (IPC): a "stream" format and a "file" format, known as Feather. read\_ipc\_stream() and read\_feather() read those formats, respectively.

#### Usage

```
read_arrow(file, ...)
read_ipc_stream(file, as_data_frame = TRUE, ...)
```

#### Arguments

file	A character file name, raw vector, or an Arrow input stream. If a file name, a
	memory-mapped Arrow InputStream will be opened and closed when finished.
	If an input stream is provided, it will be left open.
	extra parameters passed to read_feather().
as_data_frame	Should the function return a data.frame (default) or an Arrow Table?

#### Details

read\_arrow(), a wrapper around read\_ipc\_stream() and read\_feather(), is deprecated. You should explicitly choose the function that will read the desired IPC format (stream or file) since a file or InputStream may contain either.

# Value

A data.frame if as\_data\_frame is TRUE (the default), or an Arrow Table otherwise

#### See Also

read\_feather() for writing IPC files. RecordBatchReader for a lower-level interface.

read\_delim\_arrow Read a CSV or other delimited file with Arrow

#### Description

These functions uses the Arrow C++ CSV reader to read into a data.frame. Arrow C++ options have been mapped to argument names that follow those of readr::read\_delim(), and col\_select was inspired by vroom::vroom().

#### Usage

```
read_delim_arrow(
  file,
  delim = ","
  quote = "\""
  escape_double = TRUE,
  escape_backslash = FALSE,
  col_names = TRUE,
  col_select = NULL,
  na = c("", "NA"),
  quoted_na = TRUE,
  skip_empty_rows = TRUE,
  skip = 0L,
  parse_options = NULL,
  convert_options = NULL,
  read_options = NULL,
  as_data_frame = TRUE
)
read_csv_arrow(
  file,
  quote = " \setminus "",
  escape_double = TRUE,
  escape_backslash = FALSE,
  col_names = TRUE,
  col_select = NULL,
  na = c("", "NA"),
  quoted_na = TRUE,
  skip_empty_rows = TRUE,
  skip = 0L,
```

```
parse_options = NULL,
 convert_options = NULL,
 read_options = NULL,
 as_data_frame = TRUE
)
read_tsv_arrow(
  file,
 quote = " \"",
 escape_double = TRUE,
 escape_backslash = FALSE,
  col_names = TRUE,
  col_select = NULL,
 na = c("", "NA"),
 quoted_na = TRUE,
  skip_empty_rows = TRUE,
  skip = 0L,
 parse_options = NULL,
 convert_options = NULL,
 read_options = NULL,
 as_data_frame = TRUE
)
```

# Arguments

file	A character file name, raw vector, or an Arrow input stream. If a file name, a memory-mapped Arrow InputStream will be opened and closed when finished; compression will be detected from the file extension and handled automatically. If an input stream is provided, it will be left open.
delim	Single character used to separate fields within a record.
quote	Single character used to quote strings.
escape_double	Does the file escape quotes by doubling them? i.e. If this option is TRUE, the value """" represents a single quote, $\$ .
escape_backslas	h
	Does the file use backslashes to escape special characters? This is more general than escape_double as backslashes can be used to escape the delimiter character, the quote character, or to add special characters like \\n.
col_names	If TRUE, the first row of the input will be used as the column names and will not be included in the data frame. If FALSE, column names will be generated by Arrow, starting with "f0", "f1",, "fN". Alternatively, you can specify a character vector of column names.
col_select	A character vector of column names to keep, as in the "select" argument to data.table::fread(), or a tidy selection specification of columns, as used in dplyr::select().
na	A character vector of strings to interpret as missing values.

quoted_na	Should missing values inside quotes be treated as missing values (the default) or strings. (Note that this is different from the the Arrow C++ default for the corresponding convert option, $strings_can_be_null$ .)	
skip_empty_rows	3	
	Should blank rows be ignored altogether? If TRUE, blank rows will not be represented at all. If FALSE, they will be filled with missings.	
skip	Number of lines to skip before reading data.	
parse_options	see file reader options. If given, this overrides any parsing options provided in other arguments (e.g. delim, quote, etc.).	
convert_options		
	see file reader options	
read_options	see file reader options	
as_data_frame	Should the function return a data.frame (default) or an Arrow Table?	

#### Details

read\_csv\_arrow() and read\_tsv\_arrow() are wrappers around read\_delim\_arrow() that specify a delimiter.

Note that not all readr options are currently implemented here. Please file an issue if you encounter one that arrow should support.

If you need to control Arrow-specific reader parameters that don't have an equivalent in readr::read\_csv(), you can either provide them in the parse\_options, convert\_options, or read\_options arguments, or you can use CsvTableReader directly for lower-level access.

# Value

A data.frame, or a Table if as\_data\_frame = FALSE.

# Examples

```
tf <- tempfile()
on.exit(unlink(tf))
write.csv(mtcars, file = tf)
df <- read_csv_arrow(tf)
dim(df)
# Can select columns
df <- read_csv_arrow(tf, col_select = starts_with("d"))</pre>
```

 $read_feather$ 

# Description

Feather provides binary columnar serialization for data frames. It is designed to make reading and writing data frames efficient, and to make sharing data across data analysis languages easy. This function reads both the original, limited specification of the format and the version 2 specification, which is the Apache Arrow IPC file format.

#### Usage

```
read_feather(file, col_select = NULL, as_data_frame = TRUE, ...)
```

# Arguments

file	A character file name, raw vector, or an Arrow input stream. If a file name, a memory-mapped Arrow InputStream will be opened and closed when finished. If an input stream is provided, it will be left open.
col_select	A character vector of column names to keep, as in the "select" argument to data.table::fread(), or a tidy selection specification of columns, as used in dplyr::select().
as_data_frame	Should the function return a data.frame (default) or an Arrow Table?
	additional parameters, passed to FeatherReader\$create()

# Value

A data.frame if as\_data\_frame is TRUE (the default), or an Arrow Table otherwise

# See Also

FeatherReader and RecordBatchReader for lower-level access to reading Arrow IPC data.

```
tf <- tempfile()
on.exit(unlink(tf))
write_feather(mtcars, tf)
df <- read_feather(tf)
dim(df)
# Can select columns
df <- read_feather(tf, col_select = starts_with("d"))</pre>
```

read\_json\_arrow Read a JSON file

# Description

Using JsonTableReader

# Usage

```
read_json_arrow(file, col_select = NULL, as_data_frame = TRUE, ...)
```

# Arguments

file	A character file name, raw vector, or an Arrow input stream. If a file name, a memory-mapped Arrow InputStream will be opened and closed when finished; compression will be detected from the file extension and handled automatically. If an input stream is provided, it will be left open.
col_select	A character vector of column names to keep, as in the "select" argument to data.table::fread(), or a tidy selection specification of columns, as used in dplyr::select().
as_data_frame	Should the function return a data.frame (default) or an Arrow Table?
	Additional options, passed to json_table_reader()

# Value

A data.frame, or an Table if as\_data\_frame = FALSE.

```
tf <- tempfile()
on.exit(unlink(tf))
writeLines('
    { "hello": 3.5, "world": false, "yo": "thing" }
    { "hello": 3.25, "world": null }
    { "hello": 0.0, "world": true, "yo": null }
    ', tf, useBytes=TRUE)
df <- read_json_arrow(tf)</pre>
```

read\_message

Read a Parquet file

# Description

Read a Message from a stream

# Usage

read\_message(stream)

# Arguments

stream an InputStream

read\_parquet

# Description

'Parquet' is a columnar storage file format. This function enables you to read Parquet files into R.

# Usage

```
read_parquet(
   file,
   col_select = NULL,
   as_data_frame = TRUE,
   props = ParquetReaderProperties$create(),
   ...
)
```

# Arguments

file	A character file name, raw vector, or an Arrow input stream. If a file name, a memory-mapped Arrow InputStream will be opened and closed when finished. If an input stream is provided, it will be left open.
col_select	A character vector of column names to keep, as in the "select" argument to data.table::fread(), or a tidy selection specification of columns, as used in dplyr::select().
as_data_frame	Should the function return a data.frame (default) or an Arrow Table?
props	ParquetReaderProperties
	Additional arguments passed to ParquetFileReader\$create()

# Value

A arrow::Table, or a data.frame if as\_data\_frame is TRUE (the default).

# Examples

```
tf <- tempfile()
on.exit(unlink(tf))
write_parquet(mtcars, tf)
df <- read_parquet(tf, col_select = starts_with("d"))
head(df)</pre>
```

read\_schema read a Schema from a stream

# Description

read a Schema from a stream

## Usage

read\_schema(stream, ...)

# Arguments

stream	a Message, InputStream, or Buffer
	currently ignored

# Value

A Schema

RecordBatch RecordBatch class

# Description

A record batch is a collection of equal-length arrays matching a particular Schema. It is a table-like data structure that is semantically a sequence of fields, each a contiguous Arrow Array.

# Usage

record\_batch(..., schema = NULL)

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# RecordBatch

#### Arguments

	A data.frame or a named set of Arrays or vectors. If given a mixture of data.frames and vectors, the inputs will be autospliced together (see examples). Alternatively, you can provide a single Arrow IPC InputStream, Message, Buffer, or R raw object containing a Buffer.
schema	a Schema, or NULL (the default) to infer the schema from the data in When providing an Arrow IPC buffer, schema is required.

## S3 Methods and Usage

Record batches are data-frame-like, and many methods you expect to work on a data.frame are implemented for RecordBatch. This includes [, [[, \$, names, dim, nrow, ncol, head, and tail. You can also pull the data from an Arrow record batch into R with as.data.frame(). See the examples.

A caveat about the \$ method: because RecordBatch is an R6 object, \$ is also used to access the object's methods (see below). Methods take precedence over the table's columns. So, batch\$Slice would return the "Slice" method function even if there were a column in the table called "Slice".

# **R6 Methods**

In addition to the more R-friendly S3 methods, a RecordBatch object has the following R6 methods that map onto the underlying C++ methods:

- \$Equals(other): Returns TRUE if the other record batch is equal
- \$column(i): Extract an Array by integer position from the batch
- \$column\_name(i): Get a column's name by integer position
- \$names(): Get all column names (called by names(batch))
- \$GetColumnByName(name): Extract an Array by string name
- \$RemoveColumn(i): Drops a column from the batch by integer position
- \$select(spec): Return a new record batch with a selection of columns. This supports the usual character, numeric, and logical selection methods as well as "tidy select" expressions.
- \$Slice(offset, length = NULL): Create a zero-copy view starting at the indicated integer offset and going for the given length, or to the end of the table if NULL, the default.
- \$Take(i): return an RecordBatch with rows at positions given by integers (R vector or Array Array) i.
- \$Filter(i, keep\_na = TRUE): return an RecordBatch with rows at positions where logical vector (or Arrow boolean Array) i is TRUE.
- \$serialize(): Returns a raw vector suitable for interprocess communication
- \$cast(target\_schema, safe = TRUE, options = cast\_options(safe)): Alter the schema of the record batch.

There are also some active bindings

- \$num\_columns
- \$num\_rows
- \$schema

- \$metadata: Returns the key-value metadata of the Schema as a named list. Modify or replace by assigning in (batch\$metadata <-new\_metadata). All list elements are coerced to string.
- \$columns: Returns a list of Arrays

# Examples

```
batch <- record_batch(name = rownames(mtcars), mtcars)
dim(batch)
dim(head(batch))
names(batch)
batch$mpg
batch[["cyl"]]
as.data.frame(batch[4:8, c("gear", "hp", "wt")])</pre>
```

RecordBatchReader RecordBatchReader classes

#### Description

Apache Arrow defines two formats for serializing data for interprocess communication (IPC): a "stream" format and a "file" format, known as Feather. RecordBatchStreamReader and RecordBatchFileReader are interfaces for accessing record batches from input sources those formats, respectively.

For guidance on how to use these classes, see the examples section.

## Factory

The RecordBatchFileReader\$create() and RecordBatchStreamReader\$create() factory methods instantiate the object and take a single argument, named according to the class:

- file A character file name, raw vector, or Arrow file connection object (e.g. RandomAccess-File).
- stream A raw vector, Buffer, or InputStream.

## Methods

- \$read\_next\_batch(): Returns a RecordBatch, iterating through the Reader. If there are no further batches in the Reader, it returns NULL.
- \$schema: Returns a Schema (active binding)
- \$batches(): Returns a list of RecordBatches
- \$read\_table(): Collects the reader's RecordBatches into a Table
- \$get\_batch(i): For RecordBatchFileReader, return a particular batch by an integer index.
- \$num\_record\_batches(): For RecordBatchFileReader, see how many batches are in the file.

#### See Also

read\_ipc\_stream() and read\_feather() provide a much simpler interface for reading data from these formats and are sufficient for many use cases.

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## RecordBatchWriter

## Examples

```
tf <- tempfile()</pre>
on.exit(unlink(tf))
batch <- record_batch(chickwts)</pre>
# This opens a connection to the file in Arrow
file_obj <- FileOutputStream$create(tf)</pre>
# Pass that to a RecordBatchWriter to write data conforming to a schema
writer <- RecordBatchFileWriter$create(file_obj, batch$schema)</pre>
writer$write(batch)
# You may write additional batches to the stream, provided that they have
# the same schema.
# Call "close" on the writer to indicate end-of-file/stream
writer$close()
# Then, close the connection--closing the IPC message does not close the file
file_obj$close()
# Now, we have a file we can read from. Same pattern: open file connection,
# then pass it to a RecordBatchReader
read_file_obj <- ReadableFile$create(tf)</pre>
reader <- RecordBatchFileReader$create(read_file_obj)</pre>
# RecordBatchFileReader knows how many batches it has (StreamReader does not)
reader$num_record_batches
# We could consume the Reader by calling $read_next_batch() until all are,
# consumed, or we can call $read_table() to pull them all into a Table
tab <- reader$read_table()</pre>
# Call as.data.frame to turn that Table into an R data.frame
df <- as.data.frame(tab)</pre>
# This should be the same data we sent
all.equal(df, chickwts, check.attributes = FALSE)
# Unlike the Writers, we don't have to close RecordBatchReaders,
# but we do still need to close the file connection
read_file_obj$close()
```

RecordBatchWriter RecordBatchWriter classes

## Description

Apache Arrow defines two formats for serializing data for interprocess communication (IPC): a "stream" format and a "file" format, known as Feather. RecordBatchStreamWriter and RecordBatchFileWriter are interfaces for writing record batches to those formats, respectively.

For guidance on how to use these classes, see the examples section.

#### Factory

The RecordBatchFileWriter\$create() and RecordBatchStreamWriter\$create() factory methods instantiate the object and take the following arguments:

- sink An OutputStream
- schema A Schema for the data to be written
- use\_legacy\_format logical: write data formatted so that Arrow libraries versions 0.14 and lower can read it? Default is FALSE. You can also enable this by setting the environment variable ARROW\_PRE\_0\_15\_IPC\_FORMAT=1.
- metadata\_version: A string like "V5" or the equivalent integer indicating the Arrow IPC MetadataVersion. Default (NULL) will use the latest version, unless the environment variable ARROW\_PRE\_1\_0\_METADATA\_VERSION=1, in which case it will be V4.

# Methods

- \$write(x): Write a RecordBatch, Table, or data.frame, dispatching to the methods below appropriately
- \$write\_batch(batch): Write a RecordBatch to stream
- \$write\_table(table): Write a Table to stream
- \$close(): close stream. Note that this indicates end-of-file or end-of-stream-it does not close the connection to the sink. That needs to be closed separately.

## See Also

write\_ipc\_stream() and write\_feather() provide a much simpler interface for writing data to these formats and are sufficient for many use cases. write\_to\_raw() is a version that serializes data to a buffer.

## Examples

```
tf <- tempfile()
on.exit(unlink(tf))
batch <- record_batch(chickwts)
# This opens a connection to the file in Arrow
file_obj <- FileOutputStream$create(tf)
# Pass that to a RecordBatchWriter to write data conforming to a schema
writer <- RecordBatchFileWriter$create(file_obj, batch$schema)
writer$write(batch)
# You may write additional batches to the stream, provided that they have
# the same schema.
# Call "close" on the writer to indicate end-of-file/stream
writer$close()
# Then, close the connection--closing the IPC message does not close the file
file_obj$close()</pre>
```

# Now, we have a file we can read from. Same pattern: open file connection,

# Scalar

```
# then pass it to a RecordBatchReader
read_file_obj <- ReadableFile$create(tf)
reader <- RecordBatchFileReader$create(read_file_obj)
# RecordBatchFileReader knows how many batches it has (StreamReader does not)
reader$num_record_batches
# We could consume the Reader by calling $read_next_batch() until all are,
# consumed, or we can call $read_table() to pull them all into a Table
tab <- reader$read_table()
# Call as.data.frame to turn that Table into an R data.frame
df <- as.data.frame(tab)
# This should be the same data we sent
all.equal(df, chickwts, check.attributes = FALSE)
# Unlike the Writers, we don't have to close RecordBatchReaders,
# but we do still need to close the file connection
read_file_obj$close()
```

Scalar

Arrow scalars

#### Description

A Scalar holds a single value of an Arrow type.

Scanner

Scan the contents of a dataset

#### Description

A Scanner iterates over a Dataset's fragments and returns data according to given row filtering and column projection. A ScannerBuilder can help create one.

#### Factory

Scanner\$create() wraps the ScannerBuilder interface to make a Scanner. It takes the following arguments:

- dataset: A Dataset or arrow\_dplyr\_query object, as returned by the dplyr methods on Dataset.
- projection: A character vector of column names to select
- filter: A Expression to filter the scanned rows by, or TRUE (default) to keep all rows.
- use\_threads: logical: should scanning use multithreading? Default TRUE
- ...: Additional arguments, currently ignored

## Methods

ScannerBuilder has the following methods:

- \$Project(cols): Indicate that the scan should only return columns given by cols, a character vector of column names
- \$Filter(expr): Filter rows by an Expression.
- \$UseThreads(threads): logical: should the scan use multithreading? The method's default input is TRUE, but you must call the method to enable multithreading because the scanner default is FALSE.
- \$BatchSize(batch\_size): integer: Maximum row count of scanned record batches, default is 32K. If scanned record batches are overflowing memory then this method can be called to reduce their size.
- \$schema: Active binding, returns the Schema of the Dataset
- \$Finish(): Returns a Scanner

Scanner currently has a single method, \$ToTable(), which evaluates the query and returns an Arrow Table.

```
Schema
```

Schema class

## Description

A Schema is a list of Fields, which map names to Arrow data types. Create a Schema when you want to convert an R data.frame to Arrow but don't want to rely on the default mapping of R types to Arrow types, such as when you want to choose a specific numeric precision, or when creating a Dataset and you want to ensure a specific schema rather than inferring it from the various files.

Many Arrow objects, including Table and Dataset, have a \$schema method (active binding) that lets you access their schema.

#### Usage

schema(...)

# Arguments

named list of data types

#### Methods

- \$ToString(): convert to a string
- \$field(i): returns the field at index i (0-based)
- \$GetFieldByName(x): returns the field with name x
- \$WithMetadata(metadata): returns a new Schema with the key-value metadata set. Note that all list elements in metadata will be coerced to character.

## Table

## Active bindings

- \$names: returns the field names (called in names(Schema))
- \$num\_fields: returns the number of fields (called in length(Schema))
- \$fields: returns the list of Fields in the Schema, suitable for iterating over
- \$HasMetadata: logical: does this Schema have extra metadata?
- \$metadata: returns the key-value metadata as a named list. Modify or replace by assigning in (sch\$metadata <-new\_metadata). All list elements are coerced to string.

# Examples

```
df <- data.frame(col1 = 2:4, col2 = c(0.1, 0.3, 0.5))
tab1 <- Table$create(df)
tab1$schema
tab2 <- Table$create(df, schema = schema(col1 = int8(), col2 = float32()))
tab2$schema</pre>
```

Table

Table class

#### Description

A Table is a sequence of chunked arrays. They have a similar interface to record batches, but they can be composed from multiple record batches or chunked arrays.

#### Factory

The Table\$create() function takes the following arguments:

- ... arrays, chunked arrays, or R vectors, with names; alternatively, an unnamed series of record batches may also be provided, which will be stacked as rows in the table.
- schema a Schema, or NULL (the default) to infer the schema from the data in ...

#### S3 Methods and Usage

Tables are data-frame-like, and many methods you expect to work on a data.frame are implemented for Table. This includes [, [[, \$, names, dim, nrow, ncol, head, and tail. You can also pull the data from an Arrow table into R with as.data.frame(). See the examples.

A caveat about the \$ method: because Table is an R6 object, \$ is also used to access the object's methods (see below). Methods take precedence over the table's columns. So, tab\$Slice would return the "Slice" method function even if there were a column in the table called "Slice".

### **R6** Methods

In addition to the more R-friendly S3 methods, a Table object has the following R6 methods that map onto the underlying C++ methods:

- \$column(i): Extract a ChunkedArray by integer position from the table
- \$ColumnNames(): Get all column names (called by names(tab))
- \$GetColumnByName(name): Extract a ChunkedArray by string name
- \$field(i): Extract a Field from the table schema by integer position
- \$select(spec): Return a new table with a selection of columns. This supports the usual character, numeric, and logical selection methods as well as "tidy select" expressions.
- \$Slice(offset, length = NULL): Create a zero-copy view starting at the indicated integer offset and going for the given length, or to the end of the table if NULL, the default.
- \$Take(i): return an Table with rows at positions given by integers i. If i is an Arrow Array or ChunkedArray, it will be coerced to an R vector before taking.
- \$Filter(i, keep\_na = TRUE): return an Table with rows at positions where logical vector or Arrow boolean-type (Chunked)Array i is TRUE.
- \$serialize(output\_stream, ...): Write the table to the given OutputStream
- \$cast(target\_schema, safe = TRUE, options = cast\_options(safe)): Alter the schema of the record batch.

There are also some active bindings:

- \$num\_columns
- \$num\_rows
- \$schema
- \$metadata: Returns the key-value metadata of the Schema as a named list. Modify or replace by assigning in (tab\$metadata <-new\_metadata). All list elements are coerced to string.
- \$columns: Returns a list of ChunkedArrays

```
tab <- Table$create(name = rownames(mtcars), mtcars)
dim(tab)
dim(head(tab))
names(tab)
tab$mpg
tab[["cyl"]]
as.data.frame(tab[4:8, c("gear", "hp", "wt")])</pre>
```

type

# Description

infer the arrow Array type from an R vector

## Usage

type(x)

# Arguments

Х

# an R vector

# Value

an arrow logical type

unify\_schemas Combine and harmonize schemas

# Description

Combine and harmonize schemas

# Usage

unify\_schemas(..., schemas = list(...))

# Arguments

	Schemas to unify
schemas	Alternatively, a list of schemas

# Value

A Schema with the union of fields contained in the inputs

# Examples

```
## Not run:
a <- schema(b = double(), c = bool())
z <- schema(b = double(), k = utf8())
unify_schemas(a, z)
```

## End(Not run)

write\_arrow

#### Description

Apache Arrow defines two formats for serializing data for interprocess communication (IPC): a "stream" format and a "file" format, known as Feather. write\_ipc\_stream() and write\_feather() write those formats, respectively.

## Usage

write\_arrow(x, sink, ...)

write\_ipc\_stream(x, sink, ...)

#### Arguments

х	data.frame, RecordBatch, or Table
sink	A string file path or OutputStream
	extra parameters passed to write_feather().

## Details

write\_arrow(), a wrapper around write\_ipc\_stream() and write\_feather() with some nonstandard behavior, is deprecated. You should explicitly choose the function that will write the desired IPC format (stream or file) since either can be written to a file or OutputStream.

# Value

x, invisibly.

## See Also

write\_feather() for writing IPC files. write\_to\_raw() to serialize data to a buffer. Record-BatchWriter for a lower-level interface.

write\_feather

Write data in the Feather format

#### Description

Feather provides binary columnar serialization for data frames. It is designed to make reading and writing data frames efficient, and to make sharing data across data analysis languages easy. This function writes both the original, limited specification of the format and the version 2 specification, which is the Apache Arrow IPC file format.

write\_feather

# Usage

```
write_feather(
    x,
    sink,
    version = 2,
    chunk_size = 65536L,
    compression = c("default", "lz4", "uncompressed", "zstd"),
    compression_level = NULL
)
```

# Arguments

x	data.frame, RecordBatch, or Table	
sink	A string file path or OutputStream	
version	integer Feather file version. Version 2 is the current. Version 1 is the more limited legacy format.	
chunk_size	For V2 files, the number of rows that each chunk of data should have in the file. Use a smaller chunk_size when you need faster random row access. Default is 64K. This option is not supported for V1.	
compression	Name of compression codec to use, if any. Default is "lz4" if LZ4 is available in your build of the Arrow C++ library, otherwise "uncompressed". "zstd" is the other available codec and generally has better compression ratios in exchange for slower read and write performance See codec_is_available(). This op- tion is not supported for V1.	
compression_level		
	If compression is "zstd", you may specify an integer compression level. If omitted, the compression codec's default compression level is used.	

# Value

The input x, invisibly. Note that if sink is an OutputStream, the stream will be left open.

# See Also

RecordBatchWriter for lower-level access to writing Arrow IPC data.

```
tf <- tempfile()
on.exit(unlink(tf))
write_feather(mtcars, tf)</pre>
```

```
write_parquet
```

#### Description

Parquet is a columnar storage file format. This function enables you to write Parquet files from R.

#### Usage

```
write_parquet(
  х,
  sink,
  chunk_size = NULL,
  version = NULL,
  compression = NULL,
  compression_level = NULL,
  use_dictionary = NULL,
 write_statistics = NULL,
  data_page_size = NULL,
 properties = ParquetWriterProperties$create(x, version = version, compression =
  compression, compression_level = compression_level, use_dictionary = use_dictionary,
   write_statistics = write_statistics, data_page_size = data_page_size),
  use_deprecated_int96_timestamps = FALSE,
  coerce_timestamps = NULL,
  allow_truncated_timestamps = FALSE,
```

```
arrow_properties = ParquetArrowWriterProperties$create(use_deprecated_int96_timestamps
  = use_deprecated_int96_timestamps, coerce_timestamps = coerce_timestamps,
  allow_truncated_timestamps = allow_truncated_timestamps)
```

# )

# Arguments

х	An arrow::Table, or an object convertible to it.
sink	an arrow::io::OutputStream or a string which is interpreted as a file path
chunk_size	chunk size in number of rows. If NULL, the total number of rows is used.
version	parquet version, "1.0" or "2.0". Default "1.0"
compression	compression algorithm. Default "snappy". See details.
compression_level	
	compression level. Meaning depends on compression algorithm
use_dictionary	Specify if we should use dictionary encoding. Default TRUE
write_statistics	
	Specify if we should write statistics. Default TRUE
data_page_size	Set a target threshold for the approximate encoded size of data pages within a column chunk (in bytes). Default 1 MiB.

properties	properties for parquet writer, derived from arguments version, compression,
	compression_level, use_dictionary, write_statistics and data_page_size.
	You should not specify any of these arguments if you also provide a properties
	argument, as they will be ignored.

#### use\_deprecated\_int96\_timestamps

Write timestamps to INT96 Parquet format. Default FALSE.

#### coerce\_timestamps

Cast timestamps a particular resolution. Can be NULL, "ms" or "us". Default NULL (no casting)

#### allow\_truncated\_timestamps

Allow loss of data when coercing timestamps to a particular resolution. E.g. if microsecond or nanosecond data is lost when coercing to "ms", do not raise an exception

#### arrow\_properties

arrow specific writer properties, derived from arguments use\_deprecated\_int96\_timestamps, coerce\_timestamps and allow\_truncated\_timestamps You should not specify any of these arguments if you also provide a properties argument, as they will be ignored.

## Details

The parameters compression, compression\_level, use\_dictionary and write\_statistics support various patterns:

- The default NULL leaves the parameter unspecified, and the C++ library uses an appropriate default for each column (defaults listed above)
- A single, unnamed, value (e.g. a single string for compression) applies to all columns
- An unnamed vector, of the same size as the number of columns, to specify a value for each column, in positional order
- A named vector, to specify the value for the named columns, the default value for the setting is used when not supplied

The compression argument can be any of the following (case insensitive): "uncompressed", "snappy", "gzip", "brotli", "zstd", "lz4", "lzo" or "bz2". Only "uncompressed" is guaranteed to be available, but "snappy" and "gzip" are almost always included. See codec\_is\_available(). The default "snappy" is used if available, otherwise "uncompressed". To disable compression, set compression = "uncompressed". Note that "uncompressed" columns may still have dictionary encoding.

# Value

the input x invisibly.

```
tf1 <- tempfile(fileext = ".parquet")
write_parquet(data.frame(x = 1:5), tf1)</pre>
```

```
# using compression
if (codec_is_available("gzip")) {
   tf2 <- tempfile(fileext = ".gz.parquet")
   write_parquet(data.frame(x = 1:5), tf2, compression = "gzip", compression_level = 5)
}</pre>
```

write\_to\_raw

#### Write Arrow data to a raw vector

# Description

write\_ipc\_stream() and write\_feather() write data to a sink and return the data (data.frame, RecordBatch, or Table) they were given. This function wraps those so that you can serialize data to a buffer and access that buffer as a raw vector in R.

## Usage

```
write_to_raw(x, format = c("stream", "file"))
```

## Arguments

х	data.frame, RecordBatch, or Table
format	one of c("stream", "file"), indicating the IPC format to use

## Value

A raw vector containing the bytes of the IPC serialized data.

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