

# Package ‘hashmap’

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**Type** Package

**Title** The Faster Hash Map

**Version** 0.2.0

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**URL** <https://github.com/nathan-russell/hashmap>

**BugReports** <https://github.com/nathan-russell/hashmap/issues>

**Description** Provides a hash table class for fast  
key-value storage of atomic vector types.  
Internally, 'hashmap' makes extensive use of 'Rcpp', 'boost::variant',  
and 'boost::unordered\_map' to achieve high performance, type-safety,  
and versatility, while maintaining compliance with the C++98 standard.

**License** MIT + file LICENSE

**LazyData** Yes

**ByteCompile** TRUE

**Imports** Rcpp (>= 0.12.4)

**LinkingTo** Rcpp, BH

**Suggests** devtools, microbenchmark, testthat

**Depends** methods

**RcppModules** Hashmap

**Collate** 'hashmap.R' 'classes.R' 'Hashmap-class.R' 'RcppExports.R'  
'clone.R' 'load\_hashmap.R' 'merge.R' 'plugin.R'  
'save\_hashmap.R' 'zzz.R'

**RoxygenNote** 5.0.1

**NeedsCompilation** yes

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clone	<i>Clone a Hashmap</i>
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**Description**

clone creates a deep copy of a Hashmap so that modifications made to the cloned object do not affect the original object.

**Usage**

clone(x)

**Arguments**

x                    an object created by a call to hashmap.

**Details**

Since the actual cloning is done in C++, `y <- clone(x)` should be much more efficient than `y <- hashmap(x$keys(), x$values())`.

**Value**

a Hashmap identical to the input object.

**See Also**

[hashmap](#)

## Examples

```
x <- hashmap(letters[1:5], 1:5)

## shallow copy
y <- x
y[["a"]] <- 999

## original is affected
x[["a"]] == 999

z <- clone(x)
z[["c"]] <- 888

## original not affected
x[["c"]] == 888
```

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hashmap	<i>Atomic vector hash map</i>
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## Description

Create a new Hashmap instance

## Usage

```
hashmap(keys, values, ...)
```

## Arguments

keys	an atomic vector representing lookup keys
values	an atomic vector of values associated with keys in a pair-wise manner
...	other arguments passed to new when constructing the Hashmap instance

## Details

The following atomic vector types are currently supported for keys:

- integer
- numeric
- character
- Date
- POSIXct

The following atomic vector types are currently supported for values:

- logical

- integer
- numeric
- character
- complex
- Date
- POSIXct

### Value

a Hashmap object

### See Also

[Hashmap-class](#) for a more detailed discussion of available methods

### Examples

```
x <- replicate(10e3,
  paste0(sample(letters, 12, TRUE),
    collapse = ""))
)
y <- rnorm(length(x))
z <- sample(x, 100)

H <- hashmap(x, y)

all.equal(y[match(z, x)], H[[z]])

## Not run:
microbenchmark::microbenchmark(
  "R" = y[match(z, x)],
  "H" = H[[z]],
  times = 500L
)

## End(Not run)
```

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Hashmap-class

*Internal hash map class*

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### Description

A C++ class providing hash map functionality for atomic vectors

## Details

A Hashmap object (H) resulting from a call to `hashmap(keys, values)` provides the following methods accessible via `$method_name`:

- `keys()`: returns the keys of H.
- `values()`: returns the values of H.
- `cache_keys()`: caches an internal vector with the hash table's current keys resulting in very low overhead calls to `keys()`. For larger hash tables this has a significant effect. However, any calls to modifying functions (`clear`, `insert`, etc.) will invalidate the cached state.
- `cache_values()`: caches an internal vector with the hash table's current values resulting in very low overhead calls to `values()`. For larger hash tables this has a significant effect. However, any calls to modifying functions (`clear`, `insert`, etc.) will invalidate the cached state.
- `keys_cached()`: returns TRUE if the hash table's keys are currently cached, and FALSE otherwise.
- `values_cached()`: returns TRUE if the hash table's values are currently cached, and FALSE otherwise.
- `erase(remove_keys)`: deletes entries for elements that exist in the hash table, and ignores elements that do not.
- `clear()`: deletes all keys and values from H.
- `data()`: returns a named vector of values using the keys of H as names.
- `empty()`: returns TRUE if H is empty (e.g. immediately following a call to `clear`), else returns FALSE.
- `find(lookup_keys)`: returns the values associated with `lookup_keys` for existing key elements, and NA otherwise.
- `has_key(lookup_key)`: returns TRUE if `lookup_key` exists as a key in H and FALSE if it does not.
- `has_keys(lookup_keys)`: vectorized equivalent of `has_key`.
- `rehash(n_buckets)`: for the internal hash table, sets the number of buckets to at least `n` and the load factor to less than the max load factor.
- `bucket_count()`: returns the current number of buckets in the internal hash table.
- `hash_value(keys)`: compute hash values for the vector `keys` using the hash table's internal hash function. Note that `keys` need not exist in the hash table, but it must have the same type as the hash table's keys. This can be useful for investigating the efficacy of the object's hash function.
- `renew(new_keys, new_values)`: deletes current keys and values, and reinitialize H with `new_keys` and `new_values`, where `new_keys` and `new_values` are allowed to be different SEXP types than the original keys and values.
- `insert(more_keys, more_values)`: adds more key-value pairs to H, where existing key elements (`intersect(H$keys(), more_keys)`) will be updated with the corresponding elements in `more_values`, and non-existing key elements `setdiff(H$keys(), more_keys)` will be inserted with the corresponding elements in `more_values`.
- `size()`: returns the size (number of key-value pairs) of (held by) H.

Additionally, the following two convenience methods which do not require the use of \$:

- ``[[``: equivalent to `find(lookup_keys)`.
- ``[[<-``: equivalent to `insert(more_keys, more_values)`.

## Examples

```
x <- replicate(10e3,
  paste0(sample(letters, 12, TRUE),
    collapse = "")
)
y <- rnorm(length(x))
z <- sample(x, 100)

H <- hashmap(x, y)

H$empty()      #[1] FALSE
H$size()       #[1] 10000

## necessarily
any(duplicated(H$keys()))      #[1] FALSE

all.equal(H[[z]], H$find(z))  #[1] TRUE

## hash map ordering is random
all.equal(
  sort(H[[x]]),
  sort(H$values()))          #[1] TRUE

## a named vector
head(H$data())

## redundant, but TRUE
all.equal(
  H[[names(head(H$data()))]],
  unname(head(H$data()))))

## setting values
H2 <- hashmap(H$keys(), H$values())

all.equal(
  sort(H[[H2$keys()]]),
  sort(H2[[H$keys()]])      #[1] TRUE

H$insert("A", round(pi, 5))

H2[["A"]] <- round(pi, 5)

## still true
all.equal(
  sort(H[[H2$keys()]]),
  sort(H2[[H$keys()]])
```

```
## changing SEXPTYPE of key or value must be explicit
H3 <- hashmap(c("A", "B", "C"), c(1, 2, 3))

H3$size()      #[1] 3

H3$clear()
H3$size()      #[1] 0

## not allowed
class(try(H3[["D"]] <- "text", silent = TRUE)) #[1] "try-error"

## okay
H3$renew("D", "text")
H3$size()      #[1] 1
```

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internal-functions	<i>Hashmap internal functions</i>
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## Description

Hashmap internal functions

## Usage

```
.left_outer_join_impl(x, y)

.right_outer_join_impl(x, y)

.inner_join_impl(x, y)

.full_outer_join_impl(x, y)
```

## Arguments

x	an external pointer to a HashMap
y	an external pointer to a HashMap

## Details

These functions are intended for internal use only; do not call them directly.

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`load_hashmap`*Load Hashmaps*

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### Description

`load_hashmap` reads a file created by a call to [save\\_hashmap](#) and returns a Hashmap object.

### Usage

```
load_hashmap(file)
```

### Arguments

`file`                      the name of a file previously created by a call to `save_hashmap`.

### Details

The object returned will contain all of the same key-value pairs that were present in the original Hashmap at the time `save_hashmap` was called, but they are not guaranteed to be in the same order, due to rehashing.

### Value

A Hashmap object on success; an error on failure.

### See Also

[save\\_hashmap](#)

### Examples

```
H <- hashmap(sample(letters[1:10]), sample(1:10))
tf <- tempfile()

save_hashmap(H, tf)

H2 <- load_hashmap(tf)
all.equal(
  sort(H$find(H2$keys())),
  sort(H2$values())
)

all.equal(
  H$data.frame(),
  readRDS(tf)
)
```



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merge	<i>Merge two Hashmaps</i>
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## Description

merge method for Hashmap class

## Usage

```
## S3 method for class 'Rcpp_Hashmap'  
merge(x, y, type = c("inner", "left", "right", "full"),  
      ...)
```

## Arguments

x	an object created by a call to hashmap.
y	an object created by a call to hashmap.
type	a character string specifying the type of join, with partial argument matching (abbreviation) supported.
...	not used.

## Details

Valid arguments for type are:

- "inner": similar to all = FALSE in base::merge
- "left": similar to all.x = TRUE in base::merge
- "right": similar to all.y = TRUE in base::merge
- "full": similar to all = TRUE in base::merge

The default value for type is "inner".

## Value

a data.frame.

## Examples

```
hx <- hashmap(LETTERS[1:5], 1:5)  
hy <- hashmap(LETTERS[4:8], 4:8)  
  
## inner join  
merge(hx, hy)  
  
merge(  
  hx$data.frame(),  
  hy$data.frame(),
```

```
      by = "Keys",
      sort = FALSE
    )

## left join
merge(hx, hy, "left")

merge(
  hx$data.frame(),
  hy$data.frame(),
  by = "Keys",
  all.x = TRUE,
  sort = FALSE
)

## right join
merge(hx, hy, "right")

merge(
  hx$data.frame(),
  hy$data.frame(),
  by = "Keys",
  all.y = TRUE,
  sort = FALSE
)

## full outer join
merge(hx, hy, "full")

merge(
  hx$data.frame(),
  hy$data.frame(),
  by = "Keys",
  all = TRUE,
  sort = FALSE
)
```

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plot

*Plot method for Hashmap class*

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## Description

Plot method for Hashmap class

## Usage

```
## S3 method for class 'Rcpp_Hashmap'
plot(x, ...)
```

**Arguments**

x	an object created by a call to <code>hashmap</code>
...	arguments passed to <code>plot</code>

**Details**

A convenience function which simply calls `plot` using `x$keys()` and `x$values()` as plotting coordinates.

**Examples**

```
x <- hashmap(1:20, rnorm(20))
plot(x)
plot(x, type = 'p', pch = 20, col = 'red')

y <- hashmap(Sys.Date() + 1:20, rnorm(20))
plot(y, type = 'h', col = 'blue', lwd = 3)
```

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Rcpp_Hashmap-class	<i>Hashmap internal class</i>
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**Description**

Hashmap internal class

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save_hashmap	<i>Save Hashmaps</i>
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**Description**

`save_hashmap` writes a Hashmap's data to the specified file, which can be passed to [load\\_hashmap](#) at a later point in time to recreate the object.

**Usage**

```
save_hashmap(x, file, overwrite = TRUE, compress = FALSE)
```

**Arguments**

x	an object created by a call to <code>hashmap</code> .
file	a filename where the object's data will be saved.
overwrite	if TRUE (default) and file exists, it will be overwritten. If FALSE and file exists, an error is thrown.
compress	a logical value or the type of file compression to use; defaults to FALSE for better performance. See <code>?saveRDS</code> for details.

**Details**

Saving is done by calling `base::saveRDS` on the object's `data.frame` representation, `x$data.frame()`. Attempting to save an empty Hashmap results in an error.

**Value**

Nothing on success; an error on failure.

**See Also**

[load\\_hashmap](#), [saveRDS](#)

**Examples**

```
H <- hashmap(sample(letters[1:10]), sample(1:10))
tf <- tempfile()

save_hashmap(H, tf)

inherits(
  try(save_hashmap(H, tf, FALSE), silent = TRUE),
  "try-error"
)

H$insert("zzzzz", 123L)
save_hashmap(H, tf)

load_hashmap(tf)
```

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