# Package 'broadcast'

	January 29, 2025
	Simple Broadcasted and Type- Consistent Operations for Atomic and Recursive Arrays with Minimal Dependencies
Versio	n 0.0.0.9
	ption Implements simple broadcasted binding and binary operations, for atomic and recursive arrays.  Besides linking to 'Rcpp', broadcast' does not depend on, vendor, link to, or otherwise use any external libraries; broadcast' was essentially made from scratch and can be installed out-of-the-box.  The broadcasted implementations include, but are not limited to, the following.  1) Relational operations (like `==`, `!=`, `<`, `>`, `<=`, `>=`; can also take into account Machine precision);  2) Arithmetic operations (like `+`, `-`, `*`, `/`, `^`, `pmin()`, `pmax()`, integer modulo);  3) Boolean combiner operations (like `&`, ` `, `xor()`, ``nand");  4) String distance, (in)equality, and concatenation operations;  5) A Broadcasted implementation of `ifelse()`;  6) A Broadcasted apply-like function;  7) Binding arrays along any arbitrary axis, with broadcast support.  The broadcasted implementations strive to minimize computation time and memory usage (which is not just good for computer efficiency, but also for the environment).
	se MPL-2.0   file LICENSE
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Linkin	ngTo Rcpp
Roxyg	en list(markdown = TRUE)
Roxyg	enNote 7.3.1
Depen	<b>ds</b> R (>= $4.2.0$ )
_	ts Rcpp (>= 1.0.11), data.table
Sugges	sts tinytest, roxygen2, altdoc, kableExtra
Con	tents
	aaa00_broadcast_help

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aaa00\_broadcast\_help

broadcast: Simple Broadcasted Operations for Atomic and Recursive Arrays with Minimal Dependencies

### **Description**

broadcast:

Simple Broadcasted Binding and Binary Operations for Atomic and Recursive Arrays with Minimal Dependencies.

Implements simple broadcasted binding and binary operations, for atomic and recursive arrays.

Besides linking to 'Rcpp', 'broadcast' does not depend on, vendor, link to, or otherwise use any external libraries; 'broadcast' was essentially made from scratch and can be installed out-of-the-box

The broadcasted implementations include, but are not limited to, the following:

- 1. Relational operations (like ==, !=, <, >, <=, >=; can also take into account Machine precision);
- 2. Arithmetic operations (like +, -, \*, /, ^, pmin(), pmax(), integer modulo);
- 3. Boolean combiner operations (like &, |, xor(), "nand");
- 4. String distance, (in)equality, and concatenation operations;
- 5. A Broadcasted implementation of ifelse();
- 6. A Broadcasted apply-like function;
- 7. Binding arrays along any arbitrary axis, with broadcast support.

The broadcasted implementations strive to minimize computation time and memory usage (which is not just good for computer efficiency, but also for the environment).

#### **Getting Started**

An introduction and overview of the package can be found on the website.

#### **Methods and Functions**

# **Type Specific Binary Operations**

'broadcast' provides a set of functions for type-specific binary operations for broadcasted operations.

These functions use an API similar to the outer and sweep functions.

The following functions for type-specific binary operations are available:

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- bc.num: numeric arithmetic and relational operations;
- bc.b: Boolean combiner operations;
- bc.cplx: complex arithmetic and equality operations;
- bc.str: string equality, concatenation, and distance operations;
- bc.list: apply any 'R' function to 2 recursive arrays with broadcasting.

#### **General functions**

'broadcast' also comes with 2 general broadcasted functions:

- bc\_ifelse: Broadcasted version of ifelse.
- bcapply: Broadcasted apply-like function.

#### **Binding Implementations**

'broadcast' provides 3 binding implementations: bind\_mat, bind\_array, and bind\_dt.

#### Other functions

'broadcast' also provides type-casting functions, which preserve names and dimensions - convenient for arrays.

# Author(s)

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#### References

The badges shown in the documentation of this R-package were made using the services of: https://shields.io/

array\_recycle

Recycle Array Dimensions

### **Description**

The array\_recycle() function recycles array dimensions until the specified dimension sizes are reached, and returns the array.

The various broadcasting functions "recycle" an array virtually, meaning little to no additional memory is needed.

The array\_recycle() function, however, physically recycles an array (and thus actually occupies memory space).

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#### Usage

```
array_recycle(x, tdim)
```

### **Arguments**

x an atomic or recursive array or matrix.

tdim an integer vector, giving the target dimension to reach.

### Value

Returns the recycled array.

# **Examples**

```
x <- matrix(1:9, 3,3)
colnames(x) <- LETTERS[1:3]
rownames(x) <- letters[1:3]
names(x) <- month.abb[1:9]
print(x)
array_recycle(x, c(3,3,2)) # recycle to larger size</pre>
```

bc.b

Broadcasted Operations for Logical Arrays

# **Description**

The bc.b() function performs broadcasted operations on 2 logical arrays.

# Usage

```
bc.b(x, y, op)
```

# **Arguments**

x, y conformable atomic arrays of types logical, integer, or double.

op a single string, giving the operator.

Supported Boolean combiner operators: &, I, xor, nand. Supported relational operators: ==, !=, <, >, <=, >=.

#### Value

For the boolean combiner operators:

A logical array as a result of the broadcasted arithmetic operation.

For relational operators:

A logical array as a result of the broadcasted relational comparison.

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#### **Examples**

```
x.dim <- c(10:8)
x.len <- prod(x.dim)
x.data <- sample(c(TRUE, FALSE, NA), x.len, TRUE)
x <- array(x.data, x.dim)
y <- array(1:50, c(10,1,1))

bc.b(x, y, "&")
bc.b(x, y, "|")
bc.b(x, y, "xor")
bc.b(x, y, "nand")

bc.b(x, y, "==")
bc.b(x, y, "!=")
bc.b(x, y, "<")
bc.b(x, y, ">=")
```

bc.cplx

Broadcasted Operations for Complex Number Arrays

# Description

The bc.cplx() function performs broadcasted operations on 2 complex arrays.

Note that bc.cplx() uses more strict NA checks than base 'R':

If for an element of either x or y, either the real or imaginary part is NA or NaN, than the result of the operation for that element is necessarily NA.

# Usage

```
bc.cplx(x, y, op)
```

### **Arguments**

```
x, y conformable atomic arrays of typee complex.

op a single string, giving the operator.

Supported arithmetic operators: +, -, *, /.

Supported relational operators: ==, !=.
```

#### Value

For arithmetic operators:

A complex array as a result of the broadcasted arithmetic operation.

For relational operators:

A logical array as a result of the broadcasted relational comparison.

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#### **Examples**

```
x.dim <- c(10:8)
x.len <- prod(x.dim)
gen <- function() sample(c(rnorm(10), NA, NA, NaN, NaN, Inf, Inf, -Inf, -Inf))
x <- array(gen() + gen() * -1i, x.dim)
y <- array(gen() + gen() * -1i, c(10,1,1))

bc.cplx(x, y, "==")
bc.cplx(x, y, "!=")

bc.cplx(array(gen() + gen() * -1i), array(gen() + gen() * -1i), "==")
bc.cplx(array(gen() + gen() * -1i), array(gen() + gen() * -1i), "!=")

x <- gen() + gen() * -1i
y <- gen() + gen() * -1i
out <- bc.cplx(array(x), array(y), "*")
cbind(x, y, x*y, out)</pre>
```

bc.list

Broadcasted Operations for Recursive Arrays

#### **Description**

The bc.list() function performs broadcasted operations on 2 Recursive arrays.

### Usage

```
bc.list(x, y, f)
```

### **Arguments**

x, y conformable Recursive arrays (i.e. arrays of type list).

f a function that takes in exactly **2** arguments, and **returns** a result that can be stored in a single element of a list.

#### Value

A recursive array.

```
x.dim <- c(c(10, 2,2))
x.len <- prod(x.dim)

gen <- function(n) sample(list(letters, month.abb, 1:10), n, TRUE)

x <- array(gen(10), x.dim)
y <- array(gen(10), c(10,1,1))
```

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```
bc.list(
    x, y,
    \((x, y)c(length(x) == length(y), typeof(x) == typeof(y)))
```

bc.num

Broadcasted Operations for Numeric Arrays

### **Description**

The bc.num() function performs broadcasted operations on 2 numeric arrays.

### Usage

```
bc.num(x, y, op, prec = sqrt(.Machine$double.eps))
```

#### **Arguments**

x, y conformable atomic arrays of types logical, integer, or double.

op a single string, giving the operator.

Supported arithmetic operators: +, -, \*, /, ^, pmin, pmax.

Supported relational operators: ==, !=, <, >, <=, >=, d==, d!=, d<, d>, d<=, d>=.

prec a single number between 0 and 0.1, giving the machine precision to use.

Only relevant for the following operators:

d==, d!=, d<, d>, d<=, d>=

See the d==, d!=, d<, d>, d<=, d>= operators from the 'tinycodet' package for

details.

#### Value

For arithmetic operators:

A numeric array as a result of the broadcasted arithmetic operation.

For relational operators:

A logical array as a result of the broadcasted relational comparison.

```
x.dim <- c(10:8)
x.len <- prod(x.dim)
x.data <- sample(c(NA, 1.1:1000.1), x.len, TRUE)
x <- array(x.data, x.dim)
y <- array(1:50, c(10,1,1))
bc.num(x, y, "+")
bc.num(x, y, "-")</pre>
```

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```
bc.num(x, y, "*")
bc.num(x, y, "/")
bc.num(x, y, "^")

bc.num(x, y, "==")
bc.num(x, y, "!=")
bc.num(x, y, "<")
bc.num(x, y, ">")
bc.num(x, y, ">=")
bc.num(x, y, ">=")
```

bc.str

Broadcasted Operations for Character/String Arrays

# Description

The bc.str() function performs broadcasted operations on 2 character/string arrays. bc.chr() is an alias for bc.str().

# Usage

```
bc.str(x, y, op)
bc.chr(x, y, op)
```

# **Arguments**

x, y conformable atomic arrays of typee character.

op a single string, giving the operator.
Supported concatenation operators: +.
Supported relational operators: ==, !=.
Supported distance operators: levenshtein.

### Value

For concatenation operation:

A character array as a result of the broadcasted concatenation operation.

For relational operation:

A logical array as a result of the broadcasted relational comparison.

For distance operation:

An integer array as a result of the broadcasted relational comparison.

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#### **Examples**

```
# string concatenation:
x <- array(letters, c(10, 2, 1))
y <- array(letters, c(10,1,1))
bc.str(x, y, "+")

# string (in)equality:
bc.str(array(letters), array(letters), "==")
bc.str(array(letters), array(letters), "!=")

# string distance (Levenshtein):
x <- array(month.name, c(12, 1))
y <- array(month.abb, c(1, 12))
out <- bc.str(x, y, "levenshtein")
dimnames(out) <- list(month.name, month.abb)
print(out)</pre>
```

bcapply

Apply a Function to 2 Broadcasted Arrays

#### **Description**

The bcapply() function applies a function to 2 arrays with broadcasting.

# Usage

```
bcapply(x, y, f, v = "list")
```

# **Arguments**

x, y conformable atomic or recursive arrays.

f a function that takes in exactly 2 arguments, and **returns** a result that can be

stored in a single element of a recursive or atomic array.

v a single string, giving the scalar type for a single iteration.

If NULL or "list" (default), the result will be a recursive array.

If it is certain that, for every iteration, f() always results in a single atomic

**scalar**, the user can specify the type in v to pre-allocate the result.

Pre-allocating the results leads to slightly faster and more memory efficient

code.

NOTE: Incorrectly specifying v leads to undefined behaviour.

#### Value

An atomic or recursive array with dimensions  $bc_dim(x, y)$ .

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### **Examples**

```
x.dim <- c(c(10, 2,2))
x.len <- prod(x.dim)

gen <- function(n) sample(list(letters, month.abb, 1:10), n, TRUE)

x <- array(gen(10), x.dim)
y <- array(gen(10), c(10,1,1))

f <- function(x, y) list(x, y)
bcapply(x, y, f)</pre>
```

bc\_dim

Predict Broadcasted dimensions

# Description

 $bc_dim(x, y)$  gives the dimensions an array would have, as the result of an broadcasted binary element-wise operation between 2 arrays x and y.

### Usage

```
bc_dim(x, y)
```

# Arguments

x, y

an atomic array or matrix.

# Value

Returns the recycled array.

```
x.dim <- c(10:8)
x.len <- prod(x.dim)
x.data <- sample(c(TRUE, FALSE, NA), x.len, TRUE)
x <- array(x.data, x.dim)
y <- array(1:50, c(10,1,1))

dim(bc.b(x, y, "&")) == bc_dim(x, y)
dim(bc.b(x, y, "|")) == bc_dim(x, y)</pre>
```

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bc\_ifelse

Broadcasted Ifelse

#### **Description**

The bc\_ifelse() S4 generic method performs a broadcasted form of ifelse.

# Usage

```
bc_ifelse(cond, yes, no)
```

# **Arguments**

cond logical vector or array with the length equal to prod(bc\_dim(yes, no)). yes, no conformable arrays of the same type.

All atomic types (see atomic) are supported.

Recursive arrays of type list are also supported.

since bc\_ifelse() is an S4 generic, it can be extended to support special array

classes.

#### Value

The ouput, here referred to as out, will be an array of the same type as yes and no. After broadcasting yes against no, given any element index i, the following will hold for the output:

- when cond[i] == TRUE, out[i] is yes[i];
- when cond[i] == FALSE, out[i] is no[i];
- when cond[i] is NA, out[i] is NA when yes and no are atomic, and out[i] is list(NULL) when yes and no are recursive.

```
x.dim <- c(c(10, 2,2))
x.len <- prod(x.dim)

gen <- function(n) sample(list(letters, month.abb, 1:10), n, TRUE)

x <- array(gen(10), x.dim)
y <- array(gen(10), c(10,1,1))

cond <- bc.list(
    x, y,
    \((x, y)c(length(x) == length(y) && typeof(x) == typeof(y))
) |> as_bool()

bc_ifelse(cond, yes = x, no = y)
```

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bind

Dimensional Binding of Objects

# Description

The bind\_implementations provide dimensional binding functionalities.

The following implementations are available:

- bind\_mat() binds dimensionless (atomic/recursive) vectors and (atomic/recursive) matrices row- or column-wise.
   Allows for recycling.
- bind\_array() binds (atomic/recursive) arrays and (atomic/recursive) matrices. Allows for broadcasting.
- bind\_dt() binds data.tables and other data.frame-like objects.

  This function is only available if the 'data.table' package is installed.

  Returns a data.table.

  Faster than do.call(cbind, ...) or do.call(rbind, ...) for regular data.frame objects.

Note that the naming convention of the binding implementations here is "bind\_" followed by the **resulting class** (abbreviated).

I.e. bind\_mat returns a matrix, but can bind both matrices and vectors.

And bind\_array **returns** an array, but can bind both arrays and matrices.

And bind\_dt **returns** a data.table, but can bind not only data.tables, but also most other data.frame-like objects.

#### Usage

```
bind_mat(input, along, name_deparse = TRUE, comnames_from = 1L)
bind_array(
  input,
  along,
  max_bc = 1L,
  name_along = TRUE,
  comnames_from = 1L,
  name_flat = FALSE
)
bind_dt(input, along, ...)
```

### **Arguments**

input

a list of only the appropriate objects.

If input is named, its names will be used for the names of dimension along of the output, as far as possible.

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along

a single integer, indicating the dimension along which to bind the dimensions.

I.e. use along = 1 for row-binding, along = 2 for column-binding, etc.

For arrays, additional flexibility is available:

• Specifying along = 0 will bind the arrays on a new dimension before the first, making along the new first dimension.

• Specifying along = n+1, with n being the last available dimension, will create an additional dimension (n+1) and bind the arrays along that new dimension

name\_deparse

Boolean, for bind\_mat().

Indicates if dimension along should be named. Uses the naming method from rbind/cbind itself.

comnames\_from

either integer scalar or NULL, for bind\_mat() and bind\_array().

Indicates which object in input should be used for naming the shared dimen-

sion.

If NULL, no communal names will be given.

For example:

When binding columns of matrices, the matrices will share the same rownames.

Using comnames\_from = 10 will then result in bind\_array() using rownames(input[[10]])

for the rownames of the output.

max\_bc integer, for bind\_array.

Specify here the number of dimensions that are allowed to be broadcasted when

binding arrays.

If max\_bc = 0L, **no** broadcasting will be allowed at all.

name\_along Boolean, for bind\_array().

Indicates if dimension along should be named.

name\_flat Boolean, for bind\_array().

Indicates if flat indices should be named.

Note that setting this to TRUE will reduce performance considerably.

for performance: set to FALSE

... arguments to be passed to rbindlist.

# **Details**

The API of bind\_array() is inspired by the fantastic abind::abind function by Tony Plare & Richard Heiberger (2016).

But bind\_array() differs considerably from abind::abind in the following ways:

- bind\_array() differs from abind::abind in that it can handle recursive arrays properly (the abind::abind function would unlist everything to atomic arrays, ruining the structure).
- bind\_array() allows for broadcasting, while abind::abind does not support broadcasting.
- bind\_array() is generally faster than abind::abind, as bind\_array() relies heavily on 'C' and 'C++' code.
- unlike abind: :abind, bind\_array() only binds (atomic/recursive) arrays and matrices. bind\_array()does not attempt to convert things to arrays when they are not arrays, but will give an error instead.

This saves computation time and prevents unexpected results.

• bind\_array() has more streamlined naming options, compared to abind::abind.

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bind\_mat() is a modified version of rbind/cbind.

The primary differences is that  $bind_mat()$  gives an error when fractional recycling is attempted (like binding 1:3 with 1:10).

#### Value

The bound object.

# bind\_mat ====

#### References

Plate T, Heiberger R (2016). *abind: Combine Multidimensional Arrays*. R package version 1.4-5, https://CRAN.R-project.org/package=abind.

```
# bind_array ====
# here, atomic and recursive arrays are mixed,
# resulting in recursive arrays
# creating the arrays
  lapply(1:3, \(x)sample(c(TRUE, FALSE, NA))),
  lapply(1:3, \x) sample(1:10)),
  lapply(1:3, \xspace (x)rnorm(10)),
  lapply(1:3, \x) sample(letters))
x \leftarrow matrix(x, 4, 3, byrow = TRUE)
dimnames(x) <- list(letters[1:4], LETTERS[1:3])</pre>
print(x)
y <- matrix(1:12, 4, 3)
print(y)
# binding the arrays
input \leftarrow list(x = x, y = y)
bind_array(input, along = 0L) # binds on new dimension before first
bind_array(input, along = 1L) # binds on first dimension
bind_array(input, along = 2L)
bind_array(input, along = 3L) # bind on new dimension after last
# binding, wwith empty arrays
emptyarray <- array(numeric(0L), c(0L, 3L))</pre>
dimnames(emptyarray) <- list(NULL, paste("empty", 1:3))</pre>
print(emptyarray)
input <- list(x = x, y = emptyarray)
bind_array(input, along = 1L, comnames_from = 2L) # row-bind
```

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```
# here, atomic and recursive matrices are mixed,
# resulting in a recursive matrix
x <- c(
  lapply(1:3, \(x)sample(c(TRUE, FALSE, NA))),
  lapply(1:3, \x) sample(1:10)),
 lapply(1:3, \xspace (x)rnorm(10)),
 lapply(1:3, \(x)sample(letters))
x \leftarrow matrix(x, 4, 3, byrow = TRUE)
dimnames(x) <- list(letters[1:4], LETTERS[1:3])</pre>
print(x)
y <- matrix(1:12, 4, 3)
print(y)
bind_mat(list(x = x, y = y), 2L)
# bind_dt ====
x <- data.frame(a = 1:12, b = month.abb) # data.frame
y <- data.table::data.table(a = 1:12, b = month.abb) # data.table
bind_dt(list(x = x, y = y), 2L) # column bind
bind_dt(list(x = x, y = y), 1L) # row bind
```

typecast

Atomic and List Type Casting With Names and Dimensions Preserved

#### **Description**

Type casting usually strips away attributes of objects.

The functions provided here preserve dimensions, dimnames, and names, which may be more convenient for arrays and array-like objects.

The functions are as follows:

- as\_bool(): converts object to atomic type logical (TRUE, FALSE, NA).
- as\_int(): converts object to atomic type integer.
- as\_dbl(): converts object to atomic type double (AKA numeric).
- as\_chr(): converts object to atomic type character.
- as\_cplx(): converts object to atomic type complex.
- as\_raw(): converts object to atomic type raw.

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```
• as_list(): converts object to recursive type list.

as_num() is an alias for as_dbl().

as_str() is an alias for as_chr().
```

See also typeof.

# Usage

```
as_bool(x, ...)
as_int(x, ...)
as_int(x, ...)
as_dbl(x, ...)
as_num(x, ...)
as_chr(x, ...)
as_str(x, ...)
as_cplx(x, ...)
as_raw(x, ...)
as_list(x, ...)
```

# **Arguments**

x an R object.

... further arguments passed to or from other methods.

#### Value

The converted object.

```
# matrix example ====
x <- matrix(sample(-1:28), ncol = 5)
colnames(x) <- month.name[1:5]
rownames(x) <- month.abb[1:6]
names(x) <- c(letters[1:20], LETTERS[1:10])
print(x)

as_bool(x)
as_int(x)
as_dbl(x)
as_chr(x)</pre>
```

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as\_raw(x)

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