

# Package: crossmap (via r-universe)

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**Title** Apply Functions to All Combinations of List Elements

**Version** 0.4.0

**Description** Provides an extension to the 'purrr' family of mapping functions to apply a function to each combination of elements in a list of inputs. Also includes functions for automatically detecting output type in mapping functions, finding every combination of elements of lists or rows of data frames, and applying multiple models to multiple subsets of a dataset.

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**URL** <https://crossmap.rossellhayes.com>,  
<https://github.com/rossellhayes/crossmap>

**BugReports** <https://github.com/rossellhayes/crossmap/issues>

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autonames	<i>Automatically generate names for vectors</i>
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## Description

Automatically generate names for vectors

## Usage

```
autonames(x, ..., trimws = TRUE)
```

## Arguments

x	A vector
...	Additional arguments passed to <code>format()</code>
trimws	Whether to trim whitespace surrounding automatically formatted names. Defaults to TRUE.

## Value

Returns the names of a named vector and the elements of an unnamed vector formatted as characters.

## Examples

```
autonames(c(a = "apple", b = "banana", c = "cantaloupe"))
autonames(c("apple", "banana", "cantaloupe"))
```

```
autonames(10^(1:4))
autonames(10^(1:4), big.mark = ",")
autonames(10^(1:4), scientific = TRUE)
```

cross\_fit

*Cross map a model across multiple formulas, subsets, and weights***Description**

Applies a modeling function to every combination of a set of formulas and a set of data subsets.

**Usage**

```
cross_fit(
  data,
  formulas,
  cols = NULL,
  weights = NULL,
  clusters = NULL,
  families = NULL,
  fn = lm,
  fn_args = list(),
  tidy = tidy_glance,
  tidy_args = list(),
  errors = c("stop", "warn")
)
```

**Arguments**

data	A data frame
formulas	A list of formulas to apply to each subset of the data. If named, these names will be used in the <code>model</code> column of the output. Otherwise, the formulas will be converted to strings in the <code>model</code> column.
cols	Columns to subset the data. Can be any expression supported by <code>&lt;tidy-select&gt;</code> . If <code>NULL</code> , the data is not subset into columns. Defaults to <code>NULL</code> .
weights	A list of columns passed to <code>weights</code> in <code>fn</code> . If one of the elements is <code>NULL</code> or <code>NA</code> , that model will not be weighted. Defaults to <code>NULL</code> .
clusters	A list of columns passed to <code>clusters</code> if supported by <code>fn</code> . If one of the elements is <code>NULL</code> or <code>NA</code> , that model will not be clustered. Defaults to <code>NULL</code> .
families	A list of <code>glm</code> model families passed to <code>family</code> if supported by <code>fn</code> . Defaults to <code>gaussian("identity")</code> , the equivalent of <code>lm()</code> . See <code>family</code> for examples.
fn	The modeling function. Either an unquoted function name or a <code>purrr</code> -style lambda function with two arguments. To use multiple modeling functions, see <code>cross_fit_glm()</code> . Defaults to <code>lm</code> .
fn_args	A list of additional arguments to <code>fn</code> .
tidy	A logical or function to use to tidy model output into <code>data.frame</code> columns. If <code>TRUE</code> , uses the default tidying function: <code>tidy_glance()</code> . If <code>FALSE</code> , <code>NA</code> , or <code>NULL</code> , the untidied model output will be returned in a list column named <code>fit</code> . An

	alternative function can be specified with an unquoted function name or a <code>purrr</code> -style lambda function with one argument (see usage with <code>broom::tidy(conf.int = TRUE)</code> in examples). Defaults to <code>tidy_glance</code> .
<code>tidy_args</code>	A list of additional arguments to the <code>tidy</code> function
<code>errors</code>	If "stop", the default, the function will stop and return an error if any subset produces an error. If "warn", the function will produce a warning for subsets that produce an error and return results for all subsets that do not.

### Value

A tibble with a column for the model formula, columns for subsets, columns for the model family and type (if applicable), columns for the weights and clusters (if applicable), and columns of tidy model output or a list column of models (if `tidy = FALSE`)

### See Also

[cross\\_fit\\_glm\(\)](#) to map a model across multiple model types.

[cross\\_fit\\_robust\(\)](#) to map robust linear models.

[xmap\(\)](#) to apply any function to combinations of inputs.

### Examples

```
cross_fit(mtcars, mpg ~ wt, cyl)
cross_fit(mtcars, list(mpg ~ wt, mpg ~ hp), cyl)
cross_fit(mtcars, list(wt = mpg ~ wt, hp = mpg ~ hp), cyl)

cross_fit(mtcars, list(mpg ~ wt, mpg ~ hp), c(cyl, vs))
cross_fit(mtcars, list(mpg ~ wt, mpg ~ hp), dplyr::starts_with("c"))

cross_fit(mtcars, list(hp = mpg ~ hp), cyl, weights = wt)
cross_fit(mtcars, list(hp = mpg ~ hp), cyl, weights = c(wt, NA))

cross_fit(
  mtcars, list(vs ~ cyl, vs ~ hp), am,
  fn = glm, fn_args = list(family = binomial(link = logit))
)
cross_fit(
  mtcars, list(vs ~ cyl, vs ~ hp), am,
  fn = ~ glm(.x, .y, family = binomial(link = logit))
)

cross_fit(mtcars, list(mpg ~ wt, mpg ~ hp), cyl, tidy = FALSE)
cross_fit(mtcars, list(mpg ~ wt, mpg ~ hp), cyl, tidy_args = c(conf.int = TRUE))

cross_fit(mtcars, list(mpg ~ wt, mpg ~ hp), cyl, tidy = broom::tidy)
cross_fit(
  mtcars, list(mpg ~ wt, mpg ~ hp), cyl,
  tidy = ~ broom::tidy(., conf.int = TRUE)
)
```

cross\_fit\_glm

*Cross fit generalized linear models***Description**

Cross fit generalized linear models

**Usage**

```
cross_fit_glm(
  data,
  formulas,
  cols = NULL,
  weights = NULL,
  families = gaussian(link = identity),
  fn_args = list(),
  tidy = tidy_glance,
  tidy_args = list(),
  errors = c("stop", "warn")
)
```

**Arguments**

data	A data frame
formulas	A list of formulas to apply to each subset of the data. If named, these names will be used in the model column of the output. Otherwise, the formulas will be converted to strings in the model column.
cols	Columns to subset the data. Can be any expression supported by <code>&lt;tidy-select&gt;</code> . If <code>NULL</code> , the data is not subset into columns. Defaults to <code>NULL</code> .
weights	A list of columns passed to <code>weights</code> in <code>fn</code> . If one of the elements is <code>NULL</code> or <code>NA</code> , that model will not be weighted. Defaults to <code>NULL</code> .
families	A list of <code>glm</code> model families. Defaults to <code>gaussian("identity")</code> , the equivalent of <code>lm()</code> . See <code>family</code> for examples.
fn_args	A list of additional arguments to <code>glm()</code> .
tidy	A logical or function to use to tidy model output into <code>data.frame</code> columns. If <code>TRUE</code> , uses the default tidying function: <code>tidy_glance()</code> . If <code>FALSE</code> , <code>NA</code> , or <code>NULL</code> , the untidied model output will be returned in a list column named <code>fit</code> . An alternative function can be specified with an unquoted function name or a <code>purrr</code> -style lambda function with one argument (see usage with <code>broom::tidy(conf.int = TRUE)</code> in examples). Defaults to <code>tidy_glance</code> .
tidy_args	A list of additional arguments to the <code>tidy</code> function
errors	If <code>"stop"</code> , the default, the function will stop and return an error if any subset produces an error. If <code>"warn"</code> , the function will produce a warning for subsets that produce an error and return results for all subsets that do not.

**Value**

A tibble with a column for the model formula, columns for subsets, columns for the model family and type, columns for the weights (if applicable), and columns of tidy model output or a list column of models (if `tidy = FALSE`)

**See Also**

[cross\\_fit\(\)](#) to use any modeling function.

**Examples**

```
cross_fit_glm(
  data      = mtcars,
  formulas = list(am ~ gear, am ~ cyl),
  cols     = vs,
  families = list(gaussian("identity"), binomial("logit"))
)
```

---

cross_fit_robust	<i>Cross fit robust linear models</i>
------------------	---------------------------------------

---

**Description**

Cross fit robust linear models

**Usage**

```
cross_fit_robust(
  data,
  formulas,
  cols = NULL,
  weights = NULL,
  clusters = NULL,
  fn_args = list(),
  tidy = tidy_glance,
  tidy_args = list(),
  errors = c("stop", "warn")
)
```

**Arguments**

<code>data</code>	A data frame
<code>formulas</code>	A list of formulas to apply to each subset of the data. If named, these names will be used in the <code>model</code> column of the output. Otherwise, the formulas will be converted to strings in the <code>model</code> column.
<code>cols</code>	Columns to subset the data. Can be any expression supported by <code>&lt;tidy-select&gt;</code> . If <code>NULL</code> , the data is not subset into columns. Defaults to <code>NULL</code> .

weights	A list of columns passed to weights in fn. If one of the elements is <code>NULL</code> or <code>NA</code> , that model will not be weighted. Defaults to <code>NULL</code> .
clusters	A list of columns passed to clusters. If one of the elements is <code>NULL</code> or <code>NA</code> , that model will not be clustered. Defaults to <code>NULL</code> .
fn_args	A list of additional arguments to <code>estimatr::lm_robust()</code> .
tidy	A logical or function to use to tidy model output into data.frame columns. If <code>TRUE</code> , uses the default tidying function: <code>tidy_glance()</code> . If <code>FALSE</code> , <code>NA</code> , or <code>NULL</code> , the untidied model output will be returned in a list column named <code>fit</code> . An alternative function can be specified with an unquoted function name or a <code>purrr</code> -style lambda function with one argument (see usage with <code>broom::tidy(conf.int = TRUE)</code> in examples). Defaults to <code>tidy_glance</code> .
tidy_args	A list of additional arguments to the tidy function
errors	If "stop", the default, the function will stop and return an error if any subset produces an error. If "warn", the function will produce a warning for subsets that produce an error and return results for all subsets that do not.

**Value**

A tibble with a column for the model formula, columns for subsets, columns for the weights and clusters (if applicable), and columns of tidy model output or a list column of models (if `tidy = FALSE`)

**See Also**

[cross\\_fit\(\)](#) to use any modeling function.

**Examples**

```
cross_fit_robust(mtcars, mpg ~ wt, cyl, clusters = list(NULL, am))
```

---

cross\_join

*Crossing join*

---

**Description**

Adds columns from a set of data frames, creating all combinations of their rows

**Usage**

```
cross_join(..., copy = FALSE)
```

**Arguments**

- ... [Data frames](#) or a [list](#) of data frames – including data frame extensions (e.g. [tibbles](#)) and lazy data frames (e.g. from [dbplyr](#) or [dtplyr](#)). [NULL](#) inputs are silently ignored.
- copy If inputs are not from the same data source, and copy is TRUE, then they will be copied into the same src as the first input. This allows you to join tables across srcs, but it is a potentially expensive operation so you must opt into it.

**Value**

An object of the same type as the first input. The order of the rows and columns of the first input is preserved as much as possible. The output has the following properties:

- Rows from each input will be duplicated.
- Output columns include all columns from each input. If columns have the same name, suffixes are added to disambiguate.
- Groups are taken from the first input.

**See Also**

[cross\\_list\(\)](#) to find combinations of elements of vectors and lists.

**Examples**

```
fruits <- dplyr::tibble(
  fruit = c("apple", "banana", "cantaloupe"),
  color = c("red", "yellow", "orange")
)

desserts <- dplyr::tibble(
  dessert = c("cupcake", "muffin", "streudel"),
  makes = c(8, 6, 1)
)

cross_join(fruits, desserts)
cross_join(list(fruits, desserts))
cross_join(rep(list(fruits), 3))
```

---

cross\_list

*List all combinations of values*

---

**Description**

List all combinations of values

**Usage**

```
cross_list(...)
```

```
cross_tbl(...)
```

**Arguments**

... Inputs or a [list](#) of inputs. [NULL](#) inputs are silently ignored.

**Value**

A [list](#) for `cross_list()` or [tibble](#) for `cross_tbl()`. Names will match the names of the inputs. Unnamed inputs will be left unnamed for `cross_list()` and automatically named for `cross_tbl()`.

**See Also**

[cross\\_join\(\)](#) to find combinations of data frame rows.

[purrr::cross\(\)](#) for an implementation that results in a differently formatted list.

[expand.grid\(\)](#) for an implementation that results in a [data.frame](#).

**Examples**

```
fruits <- c("apple", "banana", "cantaloupe")
desserts <- c("cupcake", "muffin", "streudel")

cross_list(list(fruits, desserts))
cross_list(fruits, desserts)
cross_tbl(fruits, desserts)

cross_list(list(fruit = fruits, dessert = desserts))
cross_list(fruit = fruits, dessert = desserts)
cross_tbl(fruit = fruits, dessert = desserts)
```

---

future\_map\_vec

*Parallelized mapping functions that automatically determine type*

---

**Description**

These functions work exactly the same as [map\\_vec\(\)](#), [map2\\_vec\(\)](#), [pmap\\_vec\(\)](#), [imap\\_vec\(\)](#) and [xmap\\_vec\(\)](#), but allow you to map in parallel.

**Usage**

```
future_map_vec(  
  .x,  
  .f,  
  ...,  
  .class = NULL,  
  .progress = FALSE,  
  .options = furrr::furrr_options()  
)
```

```
future_map2_vec(  
  .x,  
  .y,  
  .f,  
  ...,  
  .class = NULL,  
  .progress = FALSE,  
  .options = furrr::furrr_options()  
)
```

```
future_pmap_vec(  
  .l,  
  .f,  
  ...,  
  .class = NULL,  
  .progress = FALSE,  
  .options = furrr::furrr_options()  
)
```

```
future_imap_vec(  
  .x,  
  .f,  
  ...,  
  .class = NULL,  
  .progress = FALSE,  
  .options = furrr::furrr_options()  
)
```

```
future_xmap_vec(  
  .l,  
  .f,  
  ...,  
  .class = NULL,  
  .progress = FALSE,  
  .options = furrr::furrr_options()  
)
```

**Arguments**

<code>.x</code>	A list or atomic vector.
<code>.f</code>	A function, formula, or vector (not necessarily atomic). If a <b>function</b> , it is used as is. If a <b>formula</b> , e.g. <code>~ .x + 2</code> , it is converted to a function. There are three ways to refer to the arguments: <ul style="list-style-type: none"> <li>• For a single argument function, use <code>.</code></li> <li>• For a two argument function, use <code>.x</code> and <code>.y</code></li> <li>• For more arguments, use <code>..1</code>, <code>..2</code>, <code>..3</code> etc</li> </ul> This syntax allows you to create very compact anonymous functions. If <b>character vector</b> , <b>numeric vector</b> , or <b>list</b> , it is converted to an extractor function. Character vectors index by name and numeric vectors index by position; use a list to index by position and name at different levels. If a component is not present, the value of <code>.default</code> will be returned.
<code>...</code>	Additional arguments passed on to <code>.f</code>
<code>.class</code>	If <code>.class</code> is specified, all
<code>.progress</code>	A single logical. Should a progress bar be displayed? Only works with multisession, multicore, and multiprocess futures. Note that if a multicore/multisession future falls back to sequential, then a progress bar will not be displayed. <b>Warning:</b> The <code>.progress</code> argument will be deprecated and removed in a future version of <code>furrr</code> in favor of using the more robust <code>progressr</code> package.
<code>.options</code>	The future specific options to use with the workers. This must be the result from a call to <code>furrr_options()</code> .
<code>.y</code>	A vector the same length as <code>.x</code> . Vectors of length 1 will be recycled.
<code>.l</code>	A list of vectors, such as a data frame. The length of <code>.l</code> determines the number of arguments that <code>.f</code> will be called with. List names will be used if present.

**Value**

Equivalent to `map_vec()`, `map2_vec()`, `pmap_vec()`, `imap_vec()` and `xmap_vec()`

**See Also**

The original functions: `furrr::future_map()`, `furrr::future_map2()`, `furrr::future_pmap()`, `furrr::future_imap()` and `future_xmap()`

Non-parallelized equivalents: `map_vec()`, `map2_vec()`, `pmap_vec()`, `imap_vec()` and `xmap_vec()`

**Examples**

```
fruits <- c("apple", "banana", "carrot", "durian", "eggplant")
desserts <- c("bread", "cake", "cupcake", "streudel", "muffin")
x <- sample(5)
y <- sample(5)
z <- sample(5)
names(z) <- fruits
```

```

future_map_vec(x, ~ . ^ 2)
future_map_vec(fruits, paste0, "s")

future_map2_vec(x, y, ~ .x + .y)
future_map2_vec(fruits, desserts, paste)

future_pmap_vec(list(x, y, z), sum)
future_pmap_vec(list(x, fruits, desserts), paste)

future_imap_vec(x, ~ .x + .y)
future_imap_vec(x, ~ paste0(.y, ": ", .x))
future_imap_vec(z, paste)

future_xmap_vec(list(x, y), ~ .x * .y)
future_xmap_vec(list(fruits, desserts), paste)

```

---

future\_xmap

*Map over each combination of list elements simultaneously via futures*


---

### Description

These functions work exactly the same as `xmap()` functions, but allow you to run the map in parallel using `future::future()`

### Usage

```
future_xmap(.l, .f, ..., .progress = FALSE, .options = furrr::furrr_options())
```

```

future_xmap_chr(
  .l,
  .f,
  ...,
  .progress = FALSE,
  .options = furrr::furrr_options()
)

```

```

future_xmap_dbl(
  .l,
  .f,
  ...,
  .progress = FALSE,
  .options = furrr::furrr_options()
)

```

```

future_xmap_dfc(
  .l,
  .f,

```

```

    ...,
    .progress = FALSE,
    .options = furrr::furrr_options()
  )

future_xmap_dfr(
  .l,
  .f,
  ...,
  .id = NULL,
  .progress = FALSE,
  .options = furrr::furrr_options()
)

future_xmap_int(
  .l,
  .f,
  ...,
  .progress = FALSE,
  .options = furrr::furrr_options()
)

future_xmap_lgl(
  .l,
  .f,
  ...,
  .progress = FALSE,
  .options = furrr::furrr_options()
)

future_xmap_raw(
  .l,
  .f,
  ...,
  .progress = FALSE,
  .options = furrr::furrr_options()
)

future_xwalk(.l, .f, ..., .progress = FALSE, .options = furrr::furrr_options())

```

### Arguments

- .l            A list of vectors, such as a data frame. The length of .l determines the number of arguments that .f will be called with. List names will be used if present.
- .f            A function, formula, or vector (not necessarily atomic).  
               If a **function**, it is used as is.  
               If a **formula**, e.g.  $\sim .x + 2$ , it is converted to a function. There are three ways to refer to the arguments:

- For a single argument function, use `.`
- For a two argument function, use `.x` and `.y`
- For more arguments, use `..1`, `..2`, `..3` etc

This syntax allows you to create very compact anonymous functions.

If **character vector**, **numeric vector**, or **list**, it is converted to an extractor function. Character vectors index by name and numeric vectors index by position; use a list to index by position and name at different levels. If a component is not present, the value of `.default` will be returned.

<code>...</code>	Additional arguments passed on to <code>.f</code>
<code>.progress</code>	A single logical. Should a progress bar be displayed? Only works with multisession, multicore, and multiprocess futures. Note that if a multicore/multisession future falls back to sequential, then a progress bar will not be displayed. <b>Warning:</b> The <code>.progress</code> argument will be deprecated and removed in a future version of <code>furrr</code> in favor of using the more robust <code>progressr</code> package.
<code>.options</code>	The future specific options to use with the workers. This must be the result from a call to <code>furrr_options()</code> .
<code>.id</code>	Either a string or NULL. If a string, the output will contain a variable with that name, storing either the name (if <code>.x</code> is named) or the index (if <code>.x</code> is unnamed) of the input. If NULL, the default, no variable will be created. Only applies to <code>_dfr</code> variant.

## Value

An atomic vector, list, or data frame, depending on the suffix. Atomic vectors and lists will be named if the first element of `.l` is named.

If all input is length 0, the output will be length 0. If any input is length 1, it will be recycled to the length of the longest.

## See Also

`xmap()` to run functions without parallel processing.

`future_xmap_vec()` to automatically determine output type.

`future_xmap_mat()` and `future_xmap_arr()` to return results in a matrix or array.

`furrr::future_map()`, `furrr::future_map2()`, and `furrr::future_pmap()` for other parallelized mapping functions.

## Examples

```
future_xmap(list(1:5, 1:5), ~ .y * .x)
future_xmap_dbl(list(1:5, 1:5), ~ .y * .x)
future_xmap_chr(list(1:5, 1:5), ~ paste(.y, "*", .x, "=", .y * .x))
```

```
apples_and_bananas <- list(
  x = c("apples", "bananas"),
  pattern = "a",
  replacement = c("oo", "ee")
```

```

)

future_xmap_chr(apples_and_bananas, gsub)

formulas <- list(mpg ~ wt, mpg ~ hp)
subsets <- split(mtcars, mtcars$cyl)

future_xmap(list(subsets, formulas), ~ lm(.y, data = .x))

```

---

future\_xmap\_mat      *Parallelized cross map returning a matrix or array*

---

## Description

Parallelized cross map returning a matrix or array

## Usage

```

future_xmap_mat(
  .l,
  .f,
  ...,
  .names = TRUE,
  .progress = FALSE,
  .options = furrr::furrr_options()
)

future_xmap_arr(
  .l,
  .f,
  ...,
  .names = TRUE,
  .progress = FALSE,
  .options = furrr::furrr_options()
)

```

## Arguments

- .l**      A list of vectors, such as a data frame. The length of **.l** determines the number of arguments that **.f** will be called with. List names will be used if present.
- .f**      A function, formula, or vector (not necessarily atomic).  
 If a **function**, it is used as is.  
 If a **formula**, e.g. `~ .x + 2`, it is converted to a function. There are three ways to refer to the arguments:
- For a single argument function, use `.`
  - For a two argument function, use `.x` and `.y`
  - For more arguments, use `..1`, `..2`, `..3` etc

This syntax allows you to create very compact anonymous functions.

If **character vector**, **numeric vector**, or **list**, it is converted to an extractor function. Character vectors index by name and numeric vectors index by position; use a list to index by position and name at different levels. If a component is not present, the value of `.default` will be returned.

<code>...</code>	Additional arguments passed on to <code>.f</code>
<code>.names</code>	A logical indicating whether to give names to the dimensions of the matrix or array. If inputs are named, the names are used. If inputs are unnamed, the elements of the input are used as names. Defaults to <code>TRUE</code> .
<code>.progress</code>	A single logical. Should a progress bar be displayed? Only works with multisession, multicore, and multiprocess futures. Note that if a multicore/multisession future falls back to sequential, then a progress bar will not be displayed. <b>Warning:</b> The <code>.progress</code> argument will be deprecated and removed in a future version of <code>furrr</code> in favor of using the more robust <code>progressr</code> package.
<code>.options</code>	The future specific options to use with the workers. This must be the result from a call to <code>furrr_options()</code> .

### Value

A matrix (for `future_xmap_mat()`) or array (for `future_xmap_arr()`) with dimensions matching the lengths of each input in `.l`.

### See Also

Unparallelized versions: `xmap_mat()` and `xmap_arr()`

`future_xmap_vec()` to return a vector.

`future_xmap()` for the underlying functions.

### Examples

```
future_xmap_mat(list(1:3, 1:3), ~ ..1 * ..2)

fruits <- c(a = "apple", b = "banana", c = "cantaloupe")
future_xmap_mat(list(1:3, fruits), paste)
future_xmap_mat(list(1:3, fruits), paste, .names = FALSE)

future_xmap_arr(list(1:3, 1:3, 1:3), ~ ..1 * ..2 * ..3)
```

---

map\_vec

*Mapping functions that automatically determine type*

---

### Description

These functions work exactly the same as typed variants of `purrr::map()`, `purrr::map2()`, `purrr::pmap()`, `purrr::imap()` and `xmap()` (e.g. `purrr::map_chr()`), but automatically determine the type.

**Usage**

```
map_vec(.x, .f, ..., .class = NULL)

map2_vec(.x, .y, .f, ..., .class = NULL)

pmap_vec(.l, .f, ..., .class = NULL)

imap_vec(.x, .f, ..., .class = NULL)

xmap_vec(.l, .f, ..., .class = NULL)
```

**Arguments**

.x	A list or atomic vector.
.f	A function, formula, or vector (not necessarily atomic). If a <b>function</b> , it is used as is. If a <b>formula</b> , e.g. $\sim .x + 2$ , it is converted to a function. There are three ways to refer to the arguments: <ul style="list-style-type: none"> <li>• For a single argument function, use <code>.</code></li> <li>• For a two argument function, use <code>.x</code> and <code>.y</code></li> <li>• For more arguments, use <code>..1</code>, <code>..2</code>, <code>..3</code> etc</li> </ul> This syntax allows you to create very compact anonymous functions. If <b>character vector</b> , <b>numeric vector</b> , or <b>list</b> , it is converted to an extractor function. Character vectors index by name and numeric vectors index by position; use a list to index by position and name at different levels. If a component is not present, the value of <code>.default</code> will be returned.
...	Additional arguments passed on to <code>.f</code>
.class	If <code>.class</code> is specified, all
.y	A vector the same length as <code>.x</code> . Vectors of length 1 will be recycled.
.l	A list of vectors, such as a data frame. The length of <code>.l</code> determines the number of arguments that <code>.f</code> will be called with. List names will be used if present.

**Value**

Equivalent to the typed variants of `purrr::map()`, `purrr::map2()`, `purrr::pmap()`, `purrr::imap()` and `xmap()` with the type automatically determined.

If the output contains multiple types, the type is determined from the highest type of the components in the hierarchy `raw < logical < integer < double < complex < character < list` (as in `c()`).

If the output contains elements that cannot be coerced to vectors (e.g. lists), the output will be a list.

**See Also**

The original functions: `purrr::map()`, `purrr::map2()`, `purrr::pmap()`, `purrr::imap()` and `xmap()`

Parallelized equivalents: `future_map_vec()`, `future_map2_vec()`, `future_pmap_vec()`, `future_imap_vec()` and `future_xmap_vec()`

**Examples**

```

fruits <- c("apple", "banana", "cantaloupe", "durian", "eggplant")
desserts <- c("bread", "cake", "cupcake", "muffin", "streudel")
x <- sample(5)
y <- sample(5)
z <- sample(5)
names(z) <- fruits

map_vec(x, ~ . ^ 2)
map_vec(fruits, paste0, "s")

map2_vec(x, y, ~ .x + .y)
map2_vec(fruits, desserts, paste)

pmap_vec(list(x, y, z), sum)
pmap_vec(list(x, fruits, desserts), paste)

imap_vec(x, ~ .x + .y)
imap_vec(x, ~ paste0(.y, ": ", .x))
imap_vec(z, paste)

xmap_vec(list(x, y), ~ .x * .y)
xmap_vec(list(fruits, desserts), paste)

```

---

tidy\_glance

*Turn an object into a tidy tibble with glance information*


---

**Description**

Apply both `generics::tidy()` and `generics::glance()` to an object and return a single `tibble` with both sets of information.

**Usage**

```
tidy_glance(x, ..., tidy_args = list(), glance_args = list())
```

**Arguments**

<code>x</code>	An object to be converted into a tidy <code>tibble</code> .
<code>...</code>	Additional arguments passed to <code>generics::tidy()</code> and <code>generics::glance()</code> . Arguments are passed to both methods, but should be ignored by the inapplicable method. For example, if called on an <code>lm</code> object, <code>conf.int</code> will affect <code>generics::tidy()</code> but not <code>generics::glance()</code> .
<code>tidy_args</code>	A list of additional arguments passed only to <code>generics::tidy()</code> .
<code>glance_args</code>	A list of additional arguments passed only to <code>generics::glance()</code> .

**Value**

A [tibble](#) with columns and rows from `generics::tidy()` and columns of repeated rows from `generics::glance()`.

Column names that appear in both the tidy data and glance data will be disambiguated by appending "model." to the glance column names.

**Examples**

```
mod <- lm(mpg ~ wt + qsec, data = mtcars)
tidy_glance(mod)
tidy_glance(mod, conf.int = TRUE)
tidy_glance(mod, tidy_args = list(conf.int = TRUE))
```

---

xmap

*Map over each combination of list elements*


---

**Description**

These functions are variants of `purrr::pmap()` that iterate over each combination of elements in a list.

**Usage**

```
xmap(.l, .f, ...)
xmap_chr(.l, .f, ...)
xmap_dbl(.l, .f, ...)
xmap_dfc(.l, .f, ...)
xmap_dfr(.l, .f, ..., .id = NULL)
xmap_int(.l, .f, ...)
xmap_lgl(.l, .f, ...)
xmap_raw(.l, .f, ...)
xwalk(.l, .f, ...)
```

**Arguments**

`.l` A list of vectors, such as a data frame. The length of `.l` determines the number of arguments that `.f` will be called with. List names will be used if present.

<code>.f</code>	A function, formula, or vector (not necessarily atomic). If a <b>function</b> , it is used as is. If a <b>formula</b> , e.g. $\sim .x + 2$ , it is converted to a function. There are three ways to refer to the arguments: <ul style="list-style-type: none"> <li>• For a single argument function, use <code>.</code></li> <li>• For a two argument function, use <code>.x</code> and <code>.y</code></li> <li>• For more arguments, use <code>. . 1</code>, <code>. . 2</code>, <code>. . 3</code> etc</li> </ul> This syntax allows you to create very compact anonymous functions. If <b>character vector</b> , <b>numeric vector</b> , or <b>list</b> , it is converted to an extractor function. Character vectors index by name and numeric vectors index by position; use a list to index by position and name at different levels. If a component is not present, the value of <code>.default</code> will be returned.
<code>...</code>	Additional arguments passed on to <code>.f</code>
<code>.id</code>	Either a string or NULL. If a string, the output will contain a variable with that name, storing either the name (if <code>.x</code> is named) or the index (if <code>.x</code> is unnamed) of the input. If NULL, the default, no variable will be created. Only applies to <code>_dfr</code> variant.

## Details

Typed variants return a vector of the specified type. To automatically determine type, try `xmap_vec()`.

To return results as a matrix or array, try `xmap_mat()` and `xmap_arr()`.

Note that a data frame is a very important special case, in which case `xmap()` and `xwalk()` apply the function `.f` to each row. `xmap_dfr()` and `xmap_dfc()` return data frames created by row-binding and column-binding respectively.

## Value

An atomic vector, list, or data frame, depending on the suffix. Atomic vectors and lists will be named if the first element of `.l` is named.

If all input is length 0, the output will be length 0. If any input is length 1, it will be recycled to the length of the longest.

## See Also

`xmap_vec()` to automatically determine output type.

`xmap_mat()` and `xmap_arr()` to return results in a matrix or array.

`future_xmap()` to run `xmap` functions with parallel processing.

`cross_fit()` to apply multiple models to multiple subsets of data.

`cross_list()` to find combinations of list elements.

`purrr::map()`, `purrr::map2()`, and `purrr::pmap()` for other mapping functions.

**Examples**

```
xmap(list(1:5, 1:5), ~ .y * .x)
xmap_dbl(list(1:5, 1:5), ~ .y * .x)
xmap_chr(list(1:5, 1:5), ~ paste(.y, "*", .x, "=", .y * .x))

apples_and_bananas <- list(
  x = c("apples", "bananas"),
  pattern = "a",
  replacement = c("oo", "ee")
)

xmap_chr(apples_and_bananas, gsub)

formulas <- list(mpg ~ wt, mpg ~ hp)
subsets <- split(mtcars, mtcars$cyl)

xmap(list(subsets, formulas), ~ lm(.y, data = .x))
xmap(list(data = subsets, formula = formulas), lm)
```

xmap\_mat

*Return a table applying a function to all combinations of list elements***Description**

Return a table applying a function to all combinations of list elements

**Usage**

```
xmap_mat(.l, .f, ..., .names = TRUE)
```

```
xmap_arr(.l, .f, ..., .names = TRUE)
```

**Arguments**

**.l** A list of vectors, such as a data frame. The length of **.l** determines the number of arguments that **.f** will be called with. List names will be used if present.

**.f** A function, formula, or vector (not necessarily atomic).

If a **function**, it is used as is.

If a **formula**, e.g. `~ .x + 2`, it is converted to a function. There are three ways to refer to the arguments:

- For a single argument function, use `.`
- For a two argument function, use `.x` and `.y`
- For more arguments, use `..1`, `..2`, `..3` etc

This syntax allows you to create very compact anonymous functions.

If **character vector**, **numeric vector**, or **list**, it is converted to an extractor function. Character vectors index by name and numeric vectors index by position; use a list to index by position and name at different levels. If a component is not present, the value of `.default` will be returned.

... Additional arguments passed on to `.f`

`.names` A logical indicating whether to give names to the dimensions of the matrix or array. If inputs are named, the names are used. If inputs are unnamed, the elements of the input are used as names. Defaults to `TRUE`.

### Value

A matrix (for `xmap_mat()`) or array (for `xmap_arr()`) with dimensions equal to the lengths of each input in `.l`.

### See Also

[future\\_xmap\\_mat\(\)](#) and [future\\_xmap\\_arr\(\)](#) to run functions in parallel.

[xmap\\_vec\(\)](#) to return a vector.

[xmap\(\)](#) for the underlying functions.

### Examples

```
xmap_mat(list(1:3, 1:3), ~ ..1 * ..2)
```

```
fruits <- c(a = "apple", b = "banana", c = "cantaloupe")
```

```
xmap_mat(list(1:3, fruits), paste)
```

```
xmap_mat(list(1:3, fruits), paste, .names = FALSE)
```

```
xmap_arr(list(1:3, 1:3, 1:3), ~ ..1 * ..2 * ..3)
```

---

xpluck

*Get one or more elements deep within a nested data structure*

---

### Description

`xpluck()` provides an alternative to [purrr::pluck\(\)](#). Unlike [purrr::pluck\(\)](#), `xpluck()` allows you to extract multiple indices at each nesting level.

### Usage

```
xpluck(.x, ..., .default = NULL)
```

### Arguments

`.x` A [list](#) or [vector](#)

... A list of accessors for indexing into the object. Can be positive integers, negative integers (to index from the right), strings (to index into names) or missing (to keep all elements at a given level).

Unlike [purrr::pluck\(\)](#), each accessor may be a vector to extract multiple elements.

`.default` Value to use if target is `NULL` or absent.

**Value**

A [list](#) or [vector](#).

**Examples**

```
obj1 <- list("a", list(1, elt = "foo"))
obj2 <- list("b", list(2, elt = "bar"))
x <- list(obj1, obj2)
```

```
xpluck(x, 1:2, 2)
xpluck(x, , 2)
```

```
xpluck(x, , 2, 1)
xpluck(x, , 2, 2)
xpluck(x, , 2, 1:2)
```

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