

# Package: Ipaper (via r-universe)

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**Type** Package

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**License** GPL-3

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## Contents

add_dn . . . . .	3
apply_3d . . . . .	3

apply_col . . . . .	4
array_3dTo2d . . . . .	5
array2dt . . . . .	6
char2script . . . . .	7
chunk . . . . .	7
clamp . . . . .	8
cut_plevels . . . . .	8
date_ym . . . . .	9
dir.show . . . . .	9
dir2 . . . . .	10
dt_day2year . . . . .	10
dt_ddply . . . . .	11
dt_tools . . . . .	12
export . . . . .	13
file_ext . . . . .	13
file_size . . . . .	14
flipud . . . . .	14
fprintf . . . . .	14
fread_dir . . . . .	15
getwd_clip . . . . .	15
git_tools . . . . .	16
github . . . . .	16
group_map2 . . . . .	16
grouped_list . . . . .	17
ifelse2 . . . . .	17
import . . . . .	18
install_gitee . . . . .	18
key_blind . . . . .	19
killCluster . . . . .	19
label_tag . . . . .	19
listk . . . . .	20
llply . . . . .	20
melt_list . . . . .	21
mkdir . . . . .	22
obj_size . . . . .	22
parse_deg . . . . .	23
path.mnt . . . . .	23
pdf_view . . . . .	23
print.data.table . . . . .	24
pro_map . . . . .	24
progressively . . . . .	25
read_excel . . . . .	26
read_xlsx . . . . .	26
read_xlsx2list . . . . .	26
runningId . . . . .	27
set_dim . . . . .	27
set_jupyter . . . . .	28
slope_arr . . . . .	28

`add_dn` 3

`which.na` . . . . . 29  
`write_fig` . . . . . 29  
`write_sheet` . . . . . 31

**Index** 33

---

`add_dn` *Add n-day flag*

---

### Description

To aggregated data into n-day (e.g. 8-day, 16-day) like MODIS product, a n-day flag is need.

### Usage

```
add_dn(d, days = 8)
```

### Arguments

`d` data.frame or data.table  
`days` Integer number or vector, can't have duplicated value.

### Examples

```
date <- seq.Date(as.Date("2010-01-01"), as.Date("2010-12-31"), by = "day")  
d <- data.frame(date)  
dnew <- add_dn(d, days = c(8, 16))
```

---

`apply_3d` *apply function for 3d array*

---

### Description

NA values will be removed automatically

### Usage

```
apply_3d(  
  array,  
  dim = 3,  
  FUN = rowMeans2,  
  by = NULL,  
  scale = 1,  
  na.rm = TRUE,  
  ...  
)
```

**Arguments**

array	A 3d array
dim	giving the subscripts to split up data by.
FUN	function, should only be row applied function, e.g. <code>matrixStats::rowMeans2</code> , <code>matrixStats::rowMins</code> , <code>matrixStats::rowRanges</code> . Because 3d array will be convert to matrix first, with the aggregated dim in the last dimension.
by	<ul style="list-style-type: none"> <li>• If not provided (NULL), the aggregated dim will be disappear. For example, daily precipitation <code>[nrow, ncol, 31-days]</code> aggregate into monthly <code>[nrow, ncol]</code>.</li> <li>• If provided, by should be equal to the aggregated dim. For example, daily precipitation <code>[nrow, ncol, 365-days]</code> aggregate into monthly <code>[nrow, ncol, 12-months]</code>. In that situation, by should be equal to 365, and be <code>format(date, '%Y%m')</code>.</li> </ul>
scale	in the same length of by, or a const value, <code>value_returned = FUN(x)*scale</code> . This parameter is designed for converting monthly to yearly, meanwhile multiply days in month. Currently, same group should have the same scale factor. Otherwise, only the first is used.

**See Also**

[apply\\_row](#) `matrixStats::rowRanges`

**Examples**

```
set.seed(1)
size <- c(10, 8, 31)
arr <- array(rnorm(10 * 8 * 31), dim = size)

by <- c(rep(1, 10), rep(2, 21))
r2 <- apply_3d(arr, 3, by = by, FUN = rowMeans)

## Not run:
arr_yearly <- apply_3d(arr, by = year(dates), scale = days_in_month(dates))

## End(Not run)
```

---

`apply_col`

*apply\_col*

---

**Description**

- `apply_col`: aggregate by col, return a `[ngrp, ncol]` matrix
- `apply_row`: aggregate by row, return a `[nrow, ngrp]` matrix

**Usage**

```
apply_col(mat, by, FUN = colMeans2, scale = 1, ...)
```

```
apply_row(mat, by, FUN = rowMeans2, scale = 1, ...)
```

**Arguments**

mat                   matrix, [nrow, ncol]  
 by                    integer vector, with the dim of [ntime]  
 scale                 in the same length of by, or a const value, value\_returned = FUN(x)\*scale.  
 This parameter is designed for converting monthly to yearly, meanwhile multiply days in month. Currently, same group should have the same scale factor. Otherwise, only the first is used.

**Details**

For example, setting the dimension of mat is [ngrid, ntime], if you want to aggregate by time, apply\_row should be used here; if you want to aggregate by region (grids), apply\_col should be used.

**Note**

This function also suits for big.matrix object.

**Examples**

```
mat <- matrix(rnorm(4 * 6), 4, 6)
mat_bycol <- apply_col(mat, c(1, 1, 2, 2), colMeans)
mat_byrow <- apply_row(mat, c(1, 1, 2, 2, 3, 3), rowMeans)
```

---

array_3dTo2d	<i>array_3dTo2d</i>
--------------	---------------------

---

**Description**

array\_3dTo2d

**Usage**

```
array_3dTo2d(array, I_grid = NULL)

array_2dTo3d(array, I_grid = NULL, dim)
```

**Arguments**

array                 array with the dimension of [nlon, nlat, ntime]  
 I\_grid                subindex of [nrow, ncol]  
 dim                   [nrow, ncol]

**Value**

[nlat\*nlon, ntime]

---

array2dt	<i>Convert array to data.table</i>
----------	------------------------------------

---

### Description

Convert array to data.table

### Usage

```
array2dt(arr, dimnames)

dt2array(dt, value_col = "value")
```

### Arguments

arr	input array
dimnames	list of dimension names, e.g., dimnames(arr)

### Value

data.table

### Examples

```
library(Ipaper)

arr <- array(1:6,
  dim = c(2, 3),
  dimnames = list(
    site = c("A", "B"),
    date = c("d1", "d2", "d3")
    # var = c("v1", "v2", "v3", "v4")
  )
)

# array -> dt
dt <- array2dt(arr, dimnames(arr))
print(dt)

# dt -> array
arr2 <- dt2array(dt)
all.equal(arr, arr2)
```

---

char2script	<i>generate R script of character vector</i>
-------------	--

---

**Description**

generate R script of character vector

**Usage**

```
char2script(x, collapse = "\", verbose = TRUE)
```

```
code_ChrVec(x, collapse = "\", verbose = TRUE)
```

**Arguments**

x character vector, data.frame or list.

collapse an optional character string to separate the results. Not NA\_character\_.

---

chunk	<i>chunk</i>
-------	--------------

---

**Description**

chunk

**Usage**

```
chunk(x, nchunk = 6)
```

**References**

<https://stackoverflow.com/questions/3318333/split-a-vector-into-chunks-in-r>

---

clamp	<i>clamp</i>
-------	--------------

---

**Description**

clamp values in the range of lims

**Usage**

```
clamp(x, lims = c(0, 1), fill.na = FALSE)
```

```
clamp_min(x, value = 0)
```

```
clamp_max(x, value = 0)
```

**Arguments**

x	Numeric vector
lims	limits
fill.na	If true, values of lims are set to NA; else, values are just constrained in the range of lims.

**Examples**

```
clamp(1:10, lims = c(4, 7), fill.na = TRUE)
```

---

cut_plevels	<i>cut_plevels</i>
-------------	--------------------

---

**Description**

cut\_plevels

**Usage**

```
cut_plevels(x, pvalue = c(0.01, 0.05, 0.1), verbose = FALSE)
```

**Arguments**

x	numeric vector
pvalue	p <= x%, means its significant at x% level

**Examples**

```
x <- c(-0.09, -0.4, 0.04, 0.15)
cut_plevels(x, verbose = TRUE)
```

---

date_ym	<i>The corresponding date of the first day of month</i>
---------	---

---

**Description**

The corresponding date of the first day of month

**Usage**

```
date_ym(date)
```

```
date_y(date)
```

```
date_yj(year, doy)
```

```
date_ydn(year, dn, delta = 8)
```

---

dir.show	<i>Open directory in Explorer</i>
----------	-----------------------------------

---

**Description**

open assign path in windows explorer, and default path is current directory. This function is only designed for windows system.

**Usage**

```
dir.show(path = getwd(), verbose = FALSE)
```

**Arguments**

path	the path you want to open
------	---------------------------

---

dir2	<i>dir2</i>
------	-------------

---

**Description**

dir2

**Usage**

```
dir2(path = ".", pattern = NULL, full.names = TRUE, ...)
```

**Arguments**

path	a character vector of full path names; the default corresponds to the working directory, <code>getwd()</code> . Tilde expansion (see <code>path.expand</code> ) is performed. Missing values will be ignored. Elements with a marked encoding will be converted to the native encoding (and if that fails, considered non-existent).
pattern	an optional <a href="#">regular expression</a> . Only file names which match the regular expression will be returned.
full.names	a logical value. If TRUE, the directory path is prepended to the file names to give a relative file path. If FALSE, the file names (rather than paths) are returned.
...	other parameters to <code>base::dir()</code>

**See Also**[base::dir\(\)](#)


---

dt_day2year	<i>dt_day2year</i>
-------------	--------------------

---

**Description**

dt\_day2year

**Usage**

```
dt_day2year(
  dat,
  nmiss_day_per_mon = 3,
  nmiss_MonPerYear = 0,
  nmin_year = 55,
  ...
)
```

**Arguments**

dat	A <code>data.table</code> , at least with the columns of <code>c("site", "date", "value")</code>
-----	--

---

dt_ddply	<i>Split data.table, apply function, and return results in a data.table.</i>
----------	--

---

## Description

For each subset of a `data.table`, apply function then combine results into a `data.table`.

## Usage

```
dt_ddply(  
  .data,  
  .variables,  
  .f = NULL,  
  ...,  
  .progress = "none",  
  .drop = TRUE,  
  .parallel = FALSE  
)
```

```
dt_ldply(  
  .data,  
  .f = NULL,  
  ...,  
  .progress = "none",  
  .parallel = FALSE,  
  .id = NA  
)
```

```
dt_dlply(  
  .data,  
  .variables,  
  .f = NULL,  
  ...,  
  .progress = "none",  
  .drop = TRUE,  
  .parallel = FALSE  
)
```

## Arguments

<code>.data</code>	data frame to be processed
<code>.variables</code>	variables to split data frame by, as <a href="#">as.quoted</a> variables, a formula or character vector
<code>.f</code>	A function, specified in one of the following ways: <ul style="list-style-type: none"><li>• A named function, e.g. <code>mean</code>.</li><li>• An anonymous function, e.g. <code>\(x) x + 1</code> or <code>function(x) x + 1</code>.</li></ul>

- A formula, e.g. `~ .x + 1`. Use `.x` to refer to the first argument. No longer recommended.
- A string, integer, or list, e.g. `"idx"`, `1`, or `list("idx", 1)` which are shorthand for `\(x) pluck(x, "idx")`, `\(x) pluck(x, 1)`, and `\(x) pluck(x, "idx", 1)` respectively. Optionally supply `.default` to set a default value if the indexed element is NULL or does not exist.

#### [Experimental]

Wrap a function with `in_parallel()` to declare that it should be performed in parallel. See `in_parallel()` for more details. Use of `...` is not permitted in this context.

<code>...</code>	other arguments passed on to <code>.fun</code>
<code>.progress</code>	name of the progress bar to use, see <code>create_progress_bar</code>
<code>.drop</code>	should combinations of variables that do not appear in the input data be preserved (FALSE) or dropped (TRUE, default)
<code>.parallel</code>	if TRUE, apply function in parallel, using parallel backend provided by <code>foreach</code>

#### Examples

```
dt <- data.table(x = 1:10, y = 1:5)
dt_dply(dt, .(y), ~.[which.max(x)])
dt_ddply(dt, .(y), ~ top_n(., 1, x))
```

---

dt\_tools

*data.frame* manipulating function by `dplyr::across`

---

#### Description

data.frame manipulating function by `dplyr::across`

#### Usage

```
dt_round(d, digits = 2)

dt_chr2num(d, fun = as.numeric)
```

---

export	<i>export data</i>
--------	--------------------

---

**Description**

export data

**Usage**

```
export(x, path, ..., nthreads = 6)
```

**Details**

Support rda, rds, fst, csv, qs

---

file_ext	<i>get file name and extension</i>
----------	------------------------------------

---

**Description**

get file name and extension

**Usage**

```
file_ext(file)
```

```
file_name(file)
```

**Arguments**

file	file path
------	-----------

**Examples**

```
file_name("./a.pdf")  
file_ext("./a.pdf")
```

---

file_size	<i>file_size</i>
-----------	------------------

---

**Description**

file\_size

**Usage**

file\_size(file)

**Arguments**

file	file path
------	-----------

---

flipud	<i>flipud and fliplr</i>
--------	--------------------------

---

**Description**

flipud and fliplr

**Usage**

flipud(x, ...)

fliplr(x)

---

fprintf	<i>fprintf Print sprintf result into console, just like C style fprintf function</i>
---------	--

---

**Description**

fprintf Print sprintf result into console, just like C style fprintf function

**Usage**

fprintf(fmt, ...)

**Arguments**

fmt	a character vector of format strings, each of up to 8192 bytes.
...	other parameters will be passed to sprintf

**Examples**

```
cat(sprintf("%s\n", "Hello phenofit!"))
```

---

fread_dir	<i>fread_dir</i>
-----------	------------------

---

**Description**

fread\_dir

**Usage**

fread\_dir(indir, pattern = "\*.csv", ..., .progress = "text", list2df = FALSE)

**Arguments**... others to `data.table::fread()`

---

getwd_clip	<i>GET or SET working directory</i>
------------	-------------------------------------

---

**Description**

- getwd\_clip: get directory path in clipboard, same as getwd function
- setwd\_clip: set directory path in clipboard, same as setwd function

**Usage**

getwd\_clip()

setwd\_clip()

**Note**

Only works in windows

**References**

1. <https://stackoverflow.com/questions/10959521/how-to-write-to-clipboard-on-ubuntu-linux-in-r>

**Examples**

```
## Not run:  
getwd_clip()  
setwd_clip()  
dir.show()  
  
## End(Not run)
```

---

git\_tools

*git tools*

---

**Description**

git tools

**Usage**

```
git_push(f = FALSE)
```

```
git_commit_amend()
```

```
git_commit(title = Sys.time())
```

```
git_set_remote(url)
```

---

github

*github*

---

**Description**

github

**Usage**

```
github(path = getwd())
```

---

group\_map2

*grouped\_list*

---

**Description**

grouped\_list

**Usage**

```
group_map2(df, .f, result.name = "data", ..., .keep = FALSE, .progress = FALSE)
```

---

grouped_list	<i>grouped_list</i>
--------------	---------------------

---

**Description**

grouped\_list

**Usage**

```
grouped_list(data, group)
```

**Arguments**

data	A list object, length(data) = nrow(group), with elements in the same order as group rows
group	A data.frame or data.table object

---

ifelse2	<i>ifelse2</i>
---------	----------------

---

**Description**

ternary operator just like java test ? yes : no

**Usage**

```
ifelse2(test, yes, no)
```

**Arguments**

test	an object which can be coerced to logical mode.
yes	return values for true elements of test.
no	return values for false elements of test.

**Examples**

```
x <- ifelse2(TRUE, 1:4, 1:10)
```

---

import	<i>import data to R</i>
--------	-------------------------

---

**Description**

import data to R

**Usage**

```
import(path, ..., nthreads = 6)
```

**Arguments**

nthreads	The number of threads to use when reading data (the initial value is 1L). When TBB is not available, values greater than 1 emit a warning and fall back to 1.
----------	---

**Details**

Support rda, rds, fst, csv, qs2

---

install_gitee	<i>Attempts to install a package directly from gitee.</i>
---------------	---

---

**Description**

Attempts to install a package directly from gitee.

**Usage**

```
install_gitee(repo)
```

**Arguments**

Repository	address in the format username/repo[/subdir][@ref #pull]. Alternatively, you can specify subdir and/or ref using the respective parameters (see below); if both is specified, the values in repo take precedence.
------------	---

**Examples**

```
## Not run:
install_gitee("adv-r/Ipaper")

## End(Not run)
```

---

key_blind	<i>blind shortcuts to rstudio addin</i>
-----------	---

---

**Description**

blind shortcuts to rstudio addin

**Usage**

```
key_blind()
```

---

killCluster	<i>killCluster</i>
-------------	--------------------

---

**Description**

killCluster

**Usage**

```
killCluster()
```

---

label_tag	<i>label_tag</i>
-----------	------------------

---

**Description**

label\_tag

**Usage**

```
label_tag(labels, tag = TRUE, expression = TRUE, letter_begin = 1)
```

```
char2expr(labels)
```

**Arguments**

labels	character vector or expression vector
--------	---------------------------------------

tag	boolean
-----	---------

**Examples**

```
label_tag(1:5)
```

```
char2expr(1:5)
```

---

listk	<i>listk</i>
-------	--------------

---

**Description**

listk

**Usage**

listk(...)

**Arguments**

... objects, possibly named.

**Examples**

```

a <- 1
b <- 1:2
c <- 1:3
l1 <- listk(a, b, c)
l2 <- listk(a, b, c = 1:3)
l3 <- listk(a = 1, b = 1:2, c = 1:3)
l4 <- listk(1, 1:2, c)

```

---

lply	<i>plyr function in purrr style</i>
------	-------------------------------------

---

**Description**

plyr function in purrr style

**Usage**

lply(.data, .f = NULL, .progress = "none", .parallel = FALSE, ...)

ldply(.data, .f = NULL, ...)

laply(.data, .f = NULL, ...)

map\_simplify(.data, .f = NULL, ...)

**Arguments**

.data	list to be processed
.progress	name of the progress bar to use, see <a href="#">create_progress_bar</a>
.parallel	if TRUE, apply function in parallel, using parallel backend provided by foreach
...	other arguments passed on to .f

**References**

1. <https://github.com/TylerGrantSmith/purrrgress>

**Examples**

```
x <- list(a = 1:10, beta = exp(-3:3), logic = c(TRUE, FALSE, FALSE, TRUE))
llply(x, mean, .progress = "text")
llply(x, ~ mean(.x), .progress = TRUE)
llply(x, quantile, probs = 1:3 / 4)
```

---

melt_list	<i>melt_list</i>
-----------	------------------

---

**Description**

melt\_list

**Usage**

```
melt_list(list, ..., na.rm = TRUE, str2num = TRUE, str2factor = TRUE)

melt_tree(x, names, ...)
```

**Arguments**

list	A list object, with the same colnames data.frame in every element.
...	other parameters to melt
na.rm	Boolean
var.name	vector of id variables. Can be integer (variable position) or string (variable name). If blank, will use all non-measured variables.

**References**

1. <https://stackoverflow.com/questions/15673550/why-is-rbindlist-better-than-rbind>

**Examples**

```
# data.frame
df <- data.frame(year = 2010, day = 1:3, month = 1, site = "A")
l <- list(a = df, b = df)

melt_list(l, 'id') %>% as_tibble()
melt_list(l, 'id' = 1) %>% as_tibble()
melt_list(set_names(l, NULL), 'id') %>% as_tibble()

# data.table
df <- data.table::data.table(year = 2010, day = 1:3, month = 1, site = "A")
```

```
l <- list(a = df, b = df)
melt_list(l, "id")
melt_list(l, id = c("a", "b"))
# int `id` is much more efficient
melt_list(l, id = c(1, 2))
```

---

mkdir	<i>mkdir</i>
-------	--------------

---

### Description

mkdir

### Usage

```
mkdir(path)

file_mv(files, outdir)

file_cp(files, outdir)

check_dir(path)
```

### Arguments

path	character vectors
------	-------------------

---

obj_size	<i>obj_size</i>
----------	-----------------

---

### Description

Get object size in unit

### Usage

```
obj_size(obj, unit = "Mb")

obj.size(obj, unit = "Mb")
```

### Arguments

obj	Object
unit	"Kb", "Mb" or "Gb"

### Examples

```
obj_size(1:100)
```

---

parse_deg	<i>parse_deg</i>
-----------	------------------

---

**Description**

- \*: zero or more
- +: one or more
- ?: zero or one

**Usage**

```
parse_deg(xs)
```

---

path.mnt	<i>convert windows path to wsl</i>
----------	------------------------------------

---

**Description**

In windows system, conversion will not be applied.

**Usage**

```
path.mnt(path)
```

```
path_mnt(path)
```

**Arguments**

path	character vector of file paths.
------	---------------------------------

---

pdf_view	<i>pdf_view</i>
----------	-----------------

---

**Description**

```
pdf_view
```

**Usage**

```
pdf_view(file, ...)
```

```
SumatraPDF(path = getwd(), verbose = FALSE)
```

```
evince(path = getwd(), verbose = FALSE)
```

**Arguments**

file                    the path of pdf file

**Note**

not work in wsl

---

print.data.table        *print.data.table*

---

**Description**

print.data.table

**Usage**

```
## S3 method for class 'data.table'
print(d, n = NULL, ..., maxrows = 1e+06)
```

**Examples**

```
d <- data.table(1:100)
options(datatable.print.nrow = 100)
print(d)
```

---

pro\_map                    *Modified purrr functions with progress bar*

---

**Description**

Modified purrr functions with progress bar

**Usage**

```
pro_map(.x, .f, ..., .progress = FALSE)
```

**Arguments**

.x                    A list or atomic vector.

.f                    A function, specified in one of the following ways:

- A named function, e.g. mean.
- An anonymous function, e.g.  $\backslash(x) \ x + 1$  or `function(x) x + 1`.
- A formula, e.g.  $\sim .x + 1$ . Use `.x` to refer to the first argument. No longer recommended.

- A string, integer, or list, e.g. "idx", 1, or list("idx", 1) which are shorthand for `\(x) pluck(x, "idx")`, `\(x) pluck(x, 1)`, and `\(x) pluck(x, "idx", 1)` respectively. Optionally supply `.default` to set a default value if the indexed element is NULL or does not exist.

### [Experimental]

Wrap a function with `in_parallel()` to declare that it should be performed in parallel. See `in_parallel()` for more details. Use of `...` is not permitted in this context.

<code>...</code>	<p>Additional arguments passed on to the mapped function.</p> <p>We now generally recommend against using <code>...</code> to pass additional (constant) arguments to <code>.f</code>. Instead use a shorthand anonymous function:</p> <pre># Instead of x  &gt; map(f, 1, 2, collapse = ",") # do: x  &gt; map(\(x) f(x, 1, 2, collapse = ","))</pre> <p>This makes it easier to understand which arguments belong to which function and will tend to yield better error messages.</p>
<code>.progress</code>	<p>Whether to show a progress bar. Use TRUE to turn on a basic progress bar, use a string to give it a name, or see <a href="#">progress_bars</a> for more details.</p>

---

progressively

*Helper function for generating progress bar functions*

---

## Description

Helper function for generating progress bar functions

## Usage

```
progressively(
  .mapper,
  .farg = NULL,
  .xarg = NULL,
  .yarg = NULL,
  .larg = NULL,
  .atarg = NULL,
  .parg = NULL
)
```

---

read_excel	<i>read csv or xls/xlsx file</i>
------------	----------------------------------

---

**Description**

read csv or xls/xlsx file

**Usage**

```
read_excel(f, ...)
```

**Arguments**

... others to fun, one of [data.table::fread\(\)](#), [readxl::read\\_xls\(\)](#), [read\\_xlsx\(\)](#)

---

read_xlsx	<i>read_xlsx</i>
-----------	------------------

---

**Description**

read\_xlsx

**Usage**

```
read_xlsx(file, sheet = 1, ...)
```

---

read_xlsx2list	<i>read_xlsx2list</i>
----------------	-----------------------

---

**Description**

If excel file hava many sheets, this function also works.

**Usage**

```
read_xlsx2list(file, ...)
```

**Arguments**

file xlsx or xls file path  
 ... other parameters to [readxl::read\\_excel\(\)](#)

---

runningId	<i>print the running ID in the console</i>
-----------	--

---

**Description**

print the running ID in the console

**Usage**

```
runningId(i, step = 1, N, prefix = "")
```

**Arguments**

i	the running Id.
step	how long of print step.
N	The number of total iteration
prefix	prefix string

**Examples**

```
for (i in 1:10) {  
  runningId(i, prefix = "phenofit")  
}
```

---

set_dim	<i>Set dimensions of an Object</i>
---------	------------------------------------

---

**Description**

Set dimensions of an Object

**Usage**

```
set_dim(x, dim)
```

**Arguments**

x	an R object, for example a matrix, array or data frame.
---	---

**See Also**

[base::dim](#)

**Examples**

```
x <- 1:12  
set_dim(x, c(3, 4))
```

---

set_jupyter	<i>set_jupyter</i>
-------------	--------------------

---

**Description**

set\_jupyter

**Usage**

```
set_jupyter(width = 10, height = 6, res = 200)
```

```
set_dirRoot(verbose = TRUE, dir = getwd())
```

**References**

1. <https://stackoverflow.com/questions/42729049/how-to-change-the-size-of-r-plots-in-jupyter>

---

slope_arr	<i>slope_arr</i>
-----------	------------------

---

**Description**

slope\_arr

**Usage**

```
slope_arr(arr, FUN = rtrend::slope_mk, return.list = FALSE)
```

**Arguments**

arr a matrix ([ngrid, ntime]) or array ([nlon, nlat, ntime]).

FUN slope functions, see [rtrend::slope\\_mk\(\)](#).

return.list boolean,

- TRUE: list(slope, pvalue) will be return
- FALSE: a array, with the dim of [nx, ny, 2].

**Value**

t, A 3d array, with the dim of [nx, ny, 2].

- t[, , 1]: slope
- t[, , 2]: pvalue

---

which.na	<i>which functions</i>
----------	------------------------

---

**Description**

which functions

**Usage**

which.na(x)

which.notna(x)

which.isnull(x)

which.null(x)

which.notnull(x)

which.notempty(x)

which.empty(x)

which.dup(x)

**Arguments**

x                    numeric vector

---

write_fig	<i>write figures into disk</i>
-----------	--------------------------------

---

**Description**

write figure to pdf, tif, png, jpg, svg, or emf, according to file suffix.

**Usage**

```
write_fig(  
  p,  
  file = "Rplot.pdf",  
  width = 10,  
  height = 5,  
  devices = NULL,  
  res = 300,
```

```

    show = TRUE,
    use.cairo_pdf = TRUE,
    use.file_show = FALSE
  )

```

### Arguments

<code>p</code>	could be one of <code>grid</code> , <code>ggplot</code> or <code>plot</code> expression
<code>file</code>	file path of output figure
<code>width</code>	the width of the device in inches.
<code>height</code>	the height of the device in inches.
<code>devices</code>	can be <code>c("pdf", "tif", "tiff", "png", "jpg", "svg", "emf")</code> . If not specified, devices will be determined according to the postfix of file. The default type is pdf.
<code>res</code>	The nominal resolution in ppi which will be recorded in the bitmap file, if a positive integer. Also used for units other than the default. If not specified, taken as 72 ppi to set the size of text and line widths.
<code>show</code>	Boolean. Whether show file after finished writing?
<code>use.cairo_pdf</code>	This parameter is for pdf type. whether to use <code>grDevices::cairo_pdf</code> ? <code>cairo_pdf</code> supports self defined font, but can not create multiple page pdf.
<code>use.file_show</code>	boolean. If true, <code>file.show</code> will be used mandatorily.

### See Also

[grDevices::cairo\\_pdf\(\)](#), [grDevices::png\(\)](#), [Cairo::Cairo\(\)](#)

### Examples

```

## Not run:
library(ggplot2)
p <- ggplot(mpg, aes(class, hwy))
p1 <- p + geom_boxplot2()

## ggplot version
write_fig(p1, "Fig. 1. ggplot.pdf", show = TRUE)
# pdf_view("Fig. 1. ggplot.pdf")
write_fig(p1, "Fig. 1. ggplot", show = TRUE)
write_fig(p1, "Fig. 1. ggplot.pdf", show = TRUE,
  devices = c("jpg", "png", "svg", "pdf", "tif", "emf"))

## lattice
x <- seq(pi/4, 5 * pi, length.out = 100)
y <- seq(pi/4, 5 * pi, length.out = 100)
r <- as.vector(sqrt(outer(x^2, y^2, "+")))
grid <- expand.grid(x=x, y=y)
grid$z <- cos(r^2) * exp(-r/(pi^3))
p <- levelplot(z~x*y, grid, cuts = 50, scales=list(log="e"), xlab="",
  ylab="", main="Weird Function", sub="with log scales",
  colorkey = FALSE, region = TRUE)

```

```

write_fig(p, "fig_lattice", show = TRUE)

## grid expression
g <- grid::circleGrob()
write_fig(g, "fig_grid", show = TRUE)

## R expression
write_fig({
  rx <- range(x <- 10*1:nrow(volcano))
  ry <- range(y <- 10*1:ncol(volcano))
  ry <- ry + c(-1, 1) * (diff(rx) - diff(ry))/2
  tcol <- terrain.colors(12)
  # par(opar);
  # opar <- par(pty = "s", bg = "lightcyan")
  plot(x = 0, y = 0, type = "n", xlim = rx, ylim = ry, xlab = "", ylab = "")
  u <- par("usr")
  rect(u[1], u[3], u[2], u[4], col = tcol[8], border = "red")
  contour(x, y, volcano, col = tcol[2], lty = "solid", add = TRUE,
          vfont = c("sans serif", "plain"))
  title("A Topographic Map of Maunga Whau", font = 4)
  abline(h = 200*0:4, v = 200*0:4, col = "lightgray", lty = 2, lwd = 0.1)
}, "fig_expr.pdf")

# quote expression also works
expr = quote({
  rx <- range(x <- 10*1:nrow(volcano))
  ry <- range(y <- 10*1:ncol(volcano))
  ry <- ry + c(-1, 1) * (diff(rx) - diff(ry))/2
  tcol <- terrain.colors(12)
  # par(opar);
  # opar <- par(pty = "s", bg = "lightcyan")
  plot(x = 0, y = 0, type = "n", xlim = rx, ylim = ry, xlab = "", ylab = "")
  u <- par("usr")
  rect(u[1], u[3], u[2], u[4], col = tcol[8], border = "red")
  contour(x, y, volcano, col = tcol[2], lty = "solid", add = TRUE,
          vfont = c("sans serif", "plain"))
  title("A Topographic Map of Maunga Whau", font = 4)
  abline(h = 200*0:4, v = 200*0:4, col = "lightgray", lty = 2, lwd = 0.1)
})
write_fig(expr, "fig_expr2.pdf")

## End(Not run)

```

---

write\_sheet

*write\_sheet*


---

## Description

Write a single data frame into one sheet of a Workbook. `file` can be a `wbWorkbook` object or a file path string. When a file path is given, the workbook is loaded (or created if the file does not exist), the sheet is written, and the file is saved back automatically.

Write a list of data frames into a Workbook, one sheet per element. `file` can be a `wbWorkbook` object or a file path string. When a file path is given, the workbook is loaded (or created if absent), written, and saved back automatically. The file extension is normalised to `.xlsx`.

### Usage

```
write_sheet(d, file, sheetName, overwrite = FALSE)

write_sheets(lst, file, .progress = "none", show = FALSE, overwrite_wb = TRUE)

write_list2xlsx(
  lst,
  file,
  .progress = "none",
  show = FALSE,
  overwrite_wb = TRUE
)
```

### Arguments

<code>d</code>	A data frame.
<code>file</code>	A file path string or an existing <code>wbWorkbook</code> object.
<code>sheetName</code>	Sheet name string.
<code>overwrite</code>	if TRUE and <code>sheetName</code> already exists, replace it.
<code>lst</code>	List of data frames. Names are used as sheet names.
<code>.progress</code>	name of the progress bar to use, see <code>create_progress_bar</code> .
<code>show</code>	open the file after saving (only applies when <code>file</code> is a path).
<code>overwrite_wb</code>	if TRUE (default) always create a fresh workbook when <code>file</code> is a path, even if the file already exists. Set to FALSE to load and append to an existing file.

### Value

The (modified) `wbWorkbook` object, invisibly.

The (modified) `wbWorkbook` object, invisibly.

# Index

add\_dn, 3  
apply\_3d, 3  
apply\_col, 4  
apply\_row, 4  
apply\_row (apply\_col), 4  
array2dt, 6  
array\_2dTo3d (array\_3dTo2d), 5  
array\_3dTo2d, 5  
as.quoted, 11  
  
base::dim, 27  
base::dir(), 10  
  
Cairo::Cairo(), 30  
char2expr (label\_tag), 19  
char2script, 7  
check\_dir (mkdir), 22  
chunk, 7  
clamp, 8  
clamp\_max (clamp), 8  
clamp\_min (clamp), 8  
code\_chrVec (char2script), 7  
create\_progress\_bar, 12, 20  
cut\_plevels, 8  
  
data.table::fread(), 15, 26  
date\_y (date\_ym), 9  
date\_ydn (date\_ym), 9  
date\_yj (date\_ym), 9  
date\_ym, 9  
dir.show, 9  
dir2, 10  
dt2array (array2dt), 6  
dt\_chr2num (dt\_tools), 12  
dt\_day2year, 10  
dt\_ddply, 11  
dt\_dply (dt\_ddply), 11  
dt\_ldply (dt\_ddply), 11  
dt\_round (dt\_tools), 12  
dt\_tools, 12  
  
evince (pdf\_view), 23  
export, 13  
  
file\_cp (mkdir), 22  
file\_ext, 13  
file\_mv (mkdir), 22  
file\_name (file\_ext), 13  
file\_size, 14  
fliplr (flipud), 14  
flipud, 14  
fprintf, 14  
fread\_dir, 15  
  
getwd, 10  
getwd\_clip, 15  
git\_commit (git\_tools), 16  
git\_commit\_amend (git\_tools), 16  
git\_push (git\_tools), 16  
git\_set\_remote (git\_tools), 16  
git\_tools, 16  
github, 16  
grDevices::cairo\_pdf(), 30  
grDevices::png(), 30  
group\_map2, 16  
grouped\_list, 17  
  
ifelse2, 17  
import, 18  
in\_parallel(), 12, 25  
install\_gitee, 18  
  
key\_blind, 19  
killCluster, 19  
  
label\_tag, 19  
lapply (llply), 20  
ldply (llply), 20  
listk, 20  
llply, 20  
  
map\_simplify (llply), 20

matrixStats::rowRanges, 4  
melt\_list, 21  
melt\_tree (melt\_list), 21  
mkdir, 22

obj.size (obj\_size), 22  
obj\_size, 22

parse\_deg, 23  
path.expand, 10  
path.mnt, 23  
path\_mnt (path.mnt), 23  
pdf\_view, 23  
print.data.table, 24  
pro\_map, 24  
progress\_bars, 25  
progressively, 25

read\_excel, 26  
read\_xlsx, 26  
read\_xlsx(), 26  
read\_xlsx2list, 26  
readxl::read\_excel(), 26  
readxl::read\_xls(), 26  
regular expression, 10  
rtrend::slope\_mk(), 28  
runningId, 27

set\_dim, 27  
set\_dirRoot (set\_jupyter), 28  
set\_jupyter, 28  
setwd\_clip (getwd\_clip), 15  
slope\_arr, 28  
SumatraPDF (pdf\_view), 23

which.dup (which.na), 29  
which.empty (which.na), 29  
which.isnull (which.na), 29  
which.na, 29  
which.notempty (which.na), 29  
which.notna (which.na), 29  
which.notnull (which.na), 29  
which.null (which.na), 29  
write\_fig, 29  
write\_list2xlsx (write\_sheet), 31  
write\_sheet, 31  
write\_sheets (write\_sheet), 31