

# Package ‘cursory’

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**Type** Package

**Title** A Cursory Look at Variables

**Version** 1.0.0

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**Description** Provides functions for quickly looking at summary statistics for variables in a data set.

**License** MIT + file LICENSE

**Encoding** UTF-8

**LazyData** true

**Language** en-US

**RoxygenNote** 6.1.1

**Imports** dplyr, methods, pkgcond, purrr, rlang, tibble, tidymargins, tidy, tidyselect

**Suggests** covr, forcats, datasets, dbplyr, testthat, testextra, DBI, RSQLite

**URL** <https://github.com/halpo/cursory>

**BugReports** <https://github.com/halpo/cursory/issues>

**NeedsCompilation** no

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add_class	<i>Add to object helpers</i>
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### Description

These function make using pipe functions easier when altering an object.

### Usage

```
add_class(x, new)
```

```
set_class(x, new)
```

```
add_comment(x, new)
```

```
set_comment(x, new)
```

### Arguments

x	object to alter
new	new characteristic

### Functions

- set\_class: Overwrite the class
- add\_comment: Add a comment
- set\_comment: Overwrite the comment

### Examples

```
# set versions will replace anything already there.
a <- set_class(TRUE, 'example')
a <- set_comment(a, 'this is an example of setting a comment')

# Add version will add to `c()` anything there.
b <- add_class(a, 'ex123')
b <- add_comment(b, 'but you can also add more information')

class(a)
class(b)

comment(a)
comment(b)
```

---

ci *Confidence interval structure*

---

**Description**

Create a confidence-interval object.

**Usage**

```
ci(estimate, lower, upper, confidence = 0.95, ...)
```

**Arguments**

estimate	The Estimate
lower	Lower bound
upper	Upper bound.
confidence	confidence level
...	other information such as

**Value**

A confidence-interval object.

**Examples**

```
x <- ci(est=0, low=-1, upp=1)
format(x)
format(x, span='---')

y <- ci(1, 0, 2, span=',')
c(x,y)
```

---

cursory *Cursory Functions*

---

**Description**

Cursory functions act like the dplyr `summarize_(all|at|if)` functions with an important difference, they put the variable name in a column and for each function passed in it puts the value in it's own column.

- `cursory_all()` is the analog of `dplyr::summarize_all()`
- `cursory_at()` is the analog of `dplyr::summarize_at()`
- `cursory_if()` is the analog of `dplyr::summarize_if()`

## Usage

```
cursory_all(.tbl, .funcs, ..., var.name = "Variable")  
cursory_at(.tbl, .vars, .funcs, ..., var.name = "Variable")  
cursory_if(.tbl, .predicate, .funcs, ..., var.name = "Variable")
```

## Arguments

<code>.tbl</code>	A <code>tbl</code> object.
<code>.funcs</code>	A function <code>fun</code> , a quosure style lambda <code>~ fun(.)</code> or a list of either form.
<code>...</code>	Additional arguments for the function calls in <code>.funcs</code> . These are evaluated only once, with <a href="#">tidy dots</a> support.
<code>var.name</code>	Name of the column with variable names.
<code>.vars</code>	A list of columns generated by <code>vars()</code> , a character vector of column names, a numeric vector of column positions, or <code>NULL</code> .
<code>.predicate</code>	A predicate function to be applied to the columns or a logical vector. The variables for which <code>.predicate</code> is or returns <code>TRUE</code> are selected. This argument is passed to <code>rlang::as_function()</code> and thus supports quosure-style lambda functions and strings representing function names.

## Value

A [tibble](#) with columns from the groups, the `var.name` column, and columns corresponding to each of function from `.funcs`.

## Examples

```
library(dplyr)  
data(iris)  
  
## basic summary statistics for each variable in a data frame.  
cursory_all(group_by(iris, Species), lst(mean, sd))  
  
## summary statistics for only numeric variables.  
cursory_if(iris, is.numeric, lst(mean, sd))  
  
## summary statistics for specific variables.  
cursory_at(iris, vars(ends_with("Length")), lst(Variance = var))
```

---

dontrepeat                      *Indicate that when printing repeat values should be hidden.*

---

**Description**

Indicate that when printing repeat values should be hidden.

**Usage**

```
dontrepeat(x, replace.with = "")
```

**Arguments**

x                                  a vector  
 replace.with            what to replace the value with.

**Examples**

```
library(dplyr)
library(tibble)
x <- cursory_all(group_by(iris, Species), lst(mean, sd))
x <- as.data.frame(arrange(x, Species))
print(x)
x[[1]] <- dontrepeat(x[[1]], replace.with='')
print(x)
```

---

Npct                                  *Combine count with a percent*

---

**Description**

Combine count with a percent

**Usage**

```
Npct(x, n, ...)
```

## S4 method for signature 'logical,missing'

```
Npct(x, n, ...)
```

## S4 method for signature 'logical,logical'

```
Npct(x, n, ...)
```

## S4 method for signature 'integer,integer'

```
Npct(x, n, ...)
```

## S4 method for signature 'numeric,numeric'

```
Npct(x, n, ...)
```

**Arguments**

x                    count or object to count  
 n                    see methods.  
 ...                  formatting arguments for formatting the percent. See format.percent.

**Value**

A character vector formatted with number and percent of successes.

**Methods (by class)**

- x = logical, n = missing: Count and give percent of TRUE from a logical vector.
- x = logical, n = logical: Count and percent of a logical filtered by a second logical.
- x = integer, n = integer: Provided with count(s) of cases and total(s)
- x = numeric, n = numeric: Provided the actual count and the percent.

**Examples**

```
# Vector of cases only.
Npct(c(TRUE, FALSE, TRUE))

# Cases with indices
Npct( c(TRUE,FALSE,TRUE,FALSE,TRUE), c(TRUE,TRUE,TRUE,FALSE,FALSE))

# Successes/Total
Npct(2L, 3L)

# Count + percent directly, count must be integerish.
Npct(2, 2/3)
```

---

percent

*Designate a numeric vector as a percent.*

---

**Description**

The percentage is stored as a formatted string with the original value as an attribute. The formatted value is what will most often be needed but but allows for the original value to be recovered when the attribute is not stripped off.

**Usage**

```
percent(x, ...)

pct(x, places = attr(x, "places") %||% getOption("percent::places", 2),
    threshold = attr(x, "threshold") %||% getOption("percent::threshold",
    1 * 10^-places), ...)
```

**Arguments**

x	a numeric object indicating a percentage.
...	additional formatting arguments.
places	Places to show after the decimal point.
threshold	The minimum absolute percentage to show.

**Value**

- For `pct()` a string formatted for a percent.
- For `percent()` the same as `pct` but is classed as a 'percent' and includes attributes for the raw value.

**Functions**

- `pct`: Format a number as a percent.

**Examples**

```
pct(2/3)           #<-- no class
(x<- percent(2/3)) #<-- has class
as.numeric(x)
```

---

table\_1

---

*Create a Dataset Summary Table (I.E. Table 1)*


---

**Description**

This helps creates demographics or summary table for a dataset, the eponymous "table 1". Given a data set a key for columns, describe the differences across the provided factor variables between the levels of key.

**Usage**

```
table_1(.data, key, .vars = vars(everything()), ...)

table_1_dispatcher(.data, var, name, key)

table_1_summarise(.data, var, name, key)

table_1_summarize(.data, var, name, key)
```

**Arguments**

<code>.data</code>	a dataset
<code>key</code>	the comparison variable, such as case/control.
<code>.vars</code>	a lazy list of variables to include in the description.
<code>...</code>	passed on to other methods.
<code>var</code>	Variable identifier, used to dispatch
<code>name</code>	name of the variable

**Details**

`table_1_summarise` and `table_1_dispatcher` dispatch on the data type of the variable identified by `var` in `.data`

**Value**

The result is a [table](#) which includes:

- A 'Variable' column, can be renamed with the `var.name` argument

**Examples**

```
table_1(iris, Species)
table_1(CO2, Plant)
```



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