

# Package ‘profr’

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**Title** An Alternative Display for Profiling Information

**Version** 0.3.3

**Description** An alternative data structure and visual rendering  
for the profiling information generated by Rprof.

**License** MIT + file LICENSE

**URL** <https://github.com/hadley/profr>

**BugReports** <https://github.com/hadley/profr/issues>

**Imports** plyr, stringr

**Suggests** ggplot2

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 6.1.1

**NeedsCompilation** no

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**Repository** CRAN

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<code>ggplot.profr</code>	<i>Visualise profiling data with ggplot2. Visualise profiling data stored in a profr data.frame.</i>
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### Description

This will plot the call tree of the specified stop watch object. If you only want a small part, you will need to subset the object

### Usage

```
ggplot.profr(data, ..., minlabel = 0.1, angle = 0)
```

### Arguments

<code>data</code>	profile output to plot
<code>...</code>	other arguments passed on to <a href="#">ggplot</a>
<code>minlabel</code>	minimum percent of time for function to get a label
<code>angle</code>	function label angle

### See Also

[plot.profr](#)

### Examples

```
if (require("ggplot2")) {  
  ggplot(nesting_prof)  
  ggplot(reshape_prof)  
}
```

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<code>parse_rprof</code>	<i>Parse Rprof output.</i>
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### Description

Parses the output of [Rprof](#) into an alternative format described in [profr](#). This produces a flat data frame, which is somewhat easier to summarise and visualise.

### Usage

```
parse_rprof(path, interval = 0.02)
```

**Arguments**

path                    path to [Rprof](#) output  
interval                real-time interval between samples (in seconds)

**Value**

[data.frame](#) of class `profr`

**See Also**

[profr](#) for profiling and parsing

**Examples**

```
nesting_ex <- system.file("samples", "nesting.rprof", package="profr")
nesting <- parse_rprof(nesting_ex)

reshape_ex <- system.file("samples", "reshape.rprof", package="profr")
diamonds <- parse_rprof(reshape_ex)
```

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plot.profr	<i>Visualise profiling data with base graphics. Visualise profiling data stored in a profr data.frame.</i>
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**Description**

If you only want a small part of the total call tree, you will need to subset the object as demonstrated by the example.

**Usage**

```
## S3 method for class 'profr'
plot(x, ..., minlabel = 0.1, angle = 0)
```

**Arguments**

x                        profile output to plot  
...                        other arguments passed on to [plot.default](#)  
minlabel                minimum percent of time for function to get a label  
angle                    function label angle

**See Also**

[ggplot.profr](#)

**Examples**

```
plot(nesting_prof)
plot(reshape_prof)
```

---

profr

*Profile the performance of a function call.*

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## Description

This is a wrapper around [Rprof](#) that provides results in an alternative data structure, a `data.frame`. The columns of the `data.frame` are:

## Usage

```
profr(expr, interval = 0.02, quiet = TRUE)
```

## Arguments

<code>expr</code>	expression to profile
<code>interval</code>	interval between samples (in seconds)
<code>quiet</code>	should output be discarded?

## Details

**f** name of function

**level** level in call stack

**time** total time (seconds) spent in function

**start** time at which control entered function

**end** time at which control exited function

**leaf** TRUE if the function is a terminal node in the call tree, i.e. didn't call any other functions

**source** guess at the package that the function came from

## Value

`data.frame` of class `profr`

## See Also

[parse\\_rprof](#) to parse standalone `Rprof` file, [plot.profr](#) and [ggplot.profr](#) to visualise the profiling data

## Examples

```
## Not run:
glm_ex <- profr({Sys.sleep(1); example(glm)}, 0.01)
head(glm_ex)
summary(glm_ex)
plot(glm_ex)

## End(Not run)
```

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`sample-data`*Sample profiling datasets*

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**Description**

These two datasets illustrate the results of running `parse_rprof` on the sample `Rprof` output stored in the `samples` directory. The output was generated by the code in `samples/generate.r`.

**Usage**`nesting_prof``reshape_prof`**Format**

a data frame

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