Package 'sprof'

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Title Profil				
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Description	n Support utilitites			
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sprof-pa	ackage s	sprof: R profiling		
Description Profiling	ng, timing and optim	ization utilitites		
Details				
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readProf reads a profile file and returns a composite structure.

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Author(s)

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```

See Also

```
proftools.
profr.
```

readProf

Read Rprof Output files and Stack Logs

Description

Read a log of stack entries, such as the output of the Rprof function, and generate a more accessible representation.

Usage

```
readProf(filename = "Rprof.out", chunksize = 5000)
```

Arguments

filename Name of a file produced by Rprof(). chunksize Number of lines to read at a time.

Details

This function reads a log file of stacks, one stack snapshot per line, stack entries separated by space. As profiling output file could be very large, it is read in blocks of chunksize lines. Increasing chunksize will make the function run faster if sufficient memory is available.

Value

A list with components

firstline A verbatim copy of the first line of the input file. Typically this contains timing

or formatting information.

nodes A vector of node names. This may include stray entries from interperspersed

lines.

stacks A vector of unique stacks found in input, stored as verbatim copies.

stacksrenc A list of unique stacks in top down order (top first), encoded as vectors or refer-

ences to stacks.

data A vector encoding the data file as referenes to stacks.

freq A frequency table summarizing data.

Author(s)

G\"unther Sawitzki

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See Also

```
summaryRprof
```

The chapter on "Tidying and profiling R code" in "Writing R Extensions" (see the 'doc/manual' subdirectory of the R source tree).

Rprof is a sampling profiler.

tracemem traces copying of an object via the C function duplicate.

Rprofmem is a non-sampling memory-use profiler.

http://developer.r-project.org/memory-profiling.html

Examples

```
## Not run:
## Rprof() is not available on all platforms
profinterval <- 0.001
simruns <- 100

n <- 10000
x <- runif(n)
y0 <- 2+ 3 * x

sinknull <- textConnection(NULL, "w"); sink(sinknull)
Rprof(tmp <- tempfile(), interval = profinterval)
for (i in 1:simruns) {y <- y0 + rnorm(n); xxx<- summary(lm(y~x))}
Rprof()

Rprof_out <- readProf(tmp)

unlink(tmp)
sink(); close(sinknull)

## End(Not run)</pre>
```

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