as.ref  

coercing to reference

Description

This function returns a reference to its argument.

Usage

as.ref(obj)

Arguments

obj an object existing in the current environment/frame

Value

an object of class "ref"

Author(s)

Jens Oehlschlägel

See Also

ref, deref

Examples

v <- 1
r <- as.ref(v)
r
deref(r)

deref  
dereferencing references

Description

This functions allow to access a referenced object. deref(ref) returns the object, and deref(ref) <- value assigns to the referenced object.
deref

Usage

deref(ref)
deref(ref) <- value  
# the following does not pass R CMD CHECK
# deref<-ref, value)
# deref(ref)[1] <- value  # subsetted assignment appears to be inefficient in S+.

Arguments

ref a reference as returned by ref or as.ref
value a value to be assigned to the reference

Details
deref and deref<- provide convenient access to objects in other environments/frames. In fact they are wrappers to get and assign. However, convenient does not necessarily means efficient. If performance is an issue, the direct use of new.env, substitute and eval may give better results. See the examples below.

Value
deref returns the referenced object.
"deref<-" returns a reference to the modified object, see ref.

Note

Subsetted assignment appears to be inefficient in S+. Note the use of substitute in the examples.

Author(s)

Jens Oehlschlägel

References

Writing R Extensions

See Also

ref, as.ref, get, assign, substitute, eval

Examples

# Simple usage example
x <- cbind(1:5, 1:5)  # take some object
rx <- as.ref(x)      # wrap it into a reference
deref(rx)            # read it through the reference
deref(rx) <- rbind(1:5, 1:5)  # replace the object in the reference by another one
deref(rx)[1,]        # read part of the object
deref(rx)[1,] <- 5:1  # replace part of the object
deref(rx)            # see the change
cat("For performance test examples see ?deref\n")

## Not run:
## Performance test examples showing actually passing by reference
# define test matrix size of nmatrix by nmatrix
nmatrix <- 1000
# you might want to use less loops in S+
# you might want more in R versions before 1.8
nloop <- 10

# Performance test using ref
t1 <- function(){  # outer function
  m <- matrix(nrow=nmatrix, ncol=nmatrix)
a <- as.ref(m)
t2(a)
m[1,1]
}

# subsetting deref is slower (by factor 75 slower since R 1.8 compared to previous versions
# , and much, much slower in S+ ...
t2 <- function(ref(){
cat("timing", timing.wrapper(
    for(i in 1:nloop)
      deref(ref)[1,1] <- i
  ), "\n")
})
if (is.R())gc()
t1()

# ... than using substitute
t2 <- function(ref(){
  obj <- as.name(ref$name)
  loc <- ref$loc
  cat("timing", timing.wrapper(
    for(i in 1:nloop)
      eval(substitute(x[1,1] <- i, list(x=obj, i=i)), loc)
  ), "\n")
})
if (is.R())gc()
t1()

# Performance test using Object (R only)
# see Henrik Bengtsson package(oo)
Object <- function()
  this <- list(env.=new.env());
  class(this) <- "Object";
  this;
"
$.Object" <- function(this, name){
  get(name, envir=unclass(this)$env.);
}
"$<-.Object" <- function(this, name, value){
  assign(name, value, envir=unclass(this)$env.);
  this;
}
HanoiTower

\)
# outer function
t1 <- function()
  o <- Object()
  o$M <- matrix(nrow=nmatrix, ncol=nmatrix)
t2(o)
o$M[1,1]
\)
# subsetting o$M is slower ...
t2 <- function(o){
cat("timing", timing.wrapper,
  for(i in 1:nloop)
o$M[1,1] <- i
), "); "}
if (is.R())gc()
t1()
# ... than using substitute
t2 <- function(o){
  env <- unclass(o)$env.
cat("timing", timing.wrapper,
  for(i in 1:nloop)
eval(substitute(m[1,1] <- i, list(i=i)), env)
), "); "}
if (is.R())gc()
t1()

## End(Not run)

---

HanoiTower  
application example for references

**Description**

This is an example for using references in S (R/S+) with package ref. HanoiTower implements a recursive algorithm solving the Hanoi tower problem. It is implemented such that the recursion can be done either by passing the HanoiTower by reference or by value to the workhorse function move.HanoiTower. Furthermore you can choose whether recursion should use Recall or should directly call move.HanoiTower. As the HanoiTower object is not too big, it can be extended by some garbage MBBytes, that will demonstrate the advantage of passing references instead of values. The deeper we recurse, the more memory we waist by passing values (and the more memory we save by passing references). Functions move.HanoiTower and print.HanoiTower are internal (not intended to be called by the user directly).

**Usage**

HanoiTower(n = 5)
Arguments

n          number of slices
parameter.mode one of "reference" or "value" deciding how to pass the HanoiTower object
recursion.mode one of "recall" or "direct" deciding how to call recursively
garbage    no. of bytes to add to the HanoiTower size
print       TRUE print the HanoiTower changes
plot        FALSE not to plot the HanoiTower changes
sleep       no. of seconds to wait between HanoiTower changes for better monitoring of progress

Details

The Hanoi Tower problem can be described as follows: you have n slices of increasing size placed on one of three locations a, b, c such that the biggest slice is at the bottom, the next biggest slice on top of it and so forth with the smallest slice as the top of the tower. Your task is to move all slices from one stick to the other, but you are only allowed to move one slice at a time and you may never put a bigger slice on top of a smaller one. The recursive solution is: to move n slices from a to c you just need to do three steps: move n-1 slices to b, move the biggest slice to c and move n-1 slices from b to c. If n equals 1, just move from a to c.

Value

invisible()

Author(s)

Jens Oehlschlägel

See Also

ref, Recall

Examples

HanoiTower(n=2)

## Not run:
# small memory examples
HanoiTowerDemoBytes <- 0
if (is.R())
  gc()
HanoiTower(
  parameter.mode = "reference",
  recursion.mode = "direct",
  garbage = HanoiTowerDemoBytes)
if (is.R())
  gc()
HanoiTower(
  parameter.mode = "reference",
  recursion.mode = "recall",
  garbage = HanoiTowerDemoBytes)
if (is.R())
  gc()
HanoiTower(
  parameter.mode = "value",
  recursion.mode = "direct",
  garbage = HanoiTowerDemoBytes)
if (is.R())
  gc()
HanoiTower(
  parameter.mode = "value",
  recursion.mode = "recall",
  garbage = HanoiTowerDemoBytes)
rm(HanoiTowerDemoBytes)

# Big memory examples
HanoiTowerDemoBytes <- 100000
if (is.R())
  gc()
HanoiTower(
  parameter.mode = "reference",
  recursion.mode = "direct",
  garbage = HanoiTowerDemoBytes)
if (is.R())
  gc()
HanoiTower(
  parameter.mode = "reference",
  recursion.mode = "recall",
  garbage = HanoiTowerDemoBytes)
if (is.R())
  gc()
HanoiTower(
  parameter.mode = "value",
  recursion.mode = "direct",
  garbage = HanoiTowerDemoBytes)
if (is.R())
gc()
HanoiTower(
    parameter.mode = "value",
    recursion.mode = "recall",
    garbage = HanoiTowerDemoBytes
)
rm(HanoiTowerDemoBytes)

## End(Not run)

### is.ref checking (for) references

**Description**

is.ref checks whether an object inherits from class "ref".  
exists.ref checks whether an referenced object exists.

**Usage**

```r
is.ref(x)  
exists.ref(ref)
```

**Arguments**

- `x`  
an object that might be a reference
- `ref`  
a reference as returned from `ref` or `as.ref`

**Value**

logical scalar

**Author(s)**

Jens Oehlschlägel

**See Also**

`ref`, `exists`, `inherits`, `class`

**Examples**

```r
v <- 1
good.r <- as.ref(v)
bad.r <- ref("NonExistingObject")
is.ref(v)
is.ref(good.r)
is.ref(bad.r)
```
### optimal.index

creating standardized, memory optimized index for subsetting

**Description**

Function `optimal.index` converts an index specification of type \{logical, integer, -integer, character\} into one of \{integer, -integer\} whatever is smaller. Function `need.index` returns TRUE if the index does represent a subset (and thus indexing is needed). Function `posi.index` returns positive integers representing the (sub)set.

**Usage**

```r
optimal.index(i, n=length(i.names), i.names = names(i), i.previous = NULL, strict = TRUE)
need.index(oi)
posi.index(oi)
```

**Arguments**

- `i` the original one-dimensional index
- `n` length of the indexed dimension (potential `iMax` if `i` where integer), not necessary if `names= ` is given
- `i.names` if `i` is character then `names= ` represents the names of the indexed dimension
- `i.previous` if `i.previous= ` is given, the returned index represents \( x[i.previous][i] \) rather than \( x[i] \) = \( x[optimal.index] \)
- `strict` set to FALSE to allow for NAs and duplicated index values, but see details
- `oi` a return value of `optimal.index`

**Details**

When `strict=TRUE` it is expected that `i` does not contain NAs and no duplicated index values. Then

\[
\text{identical}(x[i], x[optimal.index(i, n=length(x), i.names=names(x))$i]) = \text{TRUE}.
\]

When `strict=FALSE` `i` may contain NAs and/or duplicated index values. In this case length optimisation is not performed and `optimal.index` always returns positive integers.

**Value**

`optimal.index` returns the index `oi` with attributes `n=n` and `ni=length(x[optimal.index])` (which is `n-length(i)` when `i` is negative). `need.index` returns a logical scalar `posi.index` returns a vector of positive integers (or integer(0)).

**Note**

`need.index(NULL)` is defined and returns `FALSE`. This allows a function to have an optional parameter `oi=NULL` and to determine the need of subsetting in one request.
Author(s)
Jens Oehlschlägel

See Also
\texttt{refdata}
please ignore the following unpublished links: ids2index, shift.index, startstop2index

Examples

```r
1 <- letters
names(l) <- letters
stopifnot((i <- 1:3 ; identical(l[i], l[optimal.index(i, n=length(l))])))
stopifnot((i <- -(4:26) ; identical(l[i], l[optimal.index(i, n=length(l))])))
stopifnot((i <- c(rep(TRUE, 3), rep(FALSE, 23))
  identical(l[i], l[optimal.index(i, n=length(l))))))
stopifnot((i <- c("a", "b", "c")
  identical(l[i], l[optimal.index(i, i.names=names(l))))))
old.options <- options(show.error.messages=FALSE)
stopifnot(inherits(try(optimal.index(c(1:3, 3), n=length(l))), "try-error"))
options(old.options)
stopifnot((i <- c(1:3, 3, NA)
  identical(l[i], l[optimal.index(i, n=length(l), strict=FALSE))))
stopifnot((i <- c(-4:26), -26)
  identical(l[i], l[optimal.index(i, n=length(l), strict=FALSE))))
stopifnot((i <- c(rep(TRUE, 3), rep(FALSE, 23), TRUE, FALSE, NA)
  identical(l[i], l[optimal.index(i, n=length(l), strict=FALSE))))
stopifnot((i <- c("a", "b", "c", "a", NA)
  identical(l[i], l[optimal.index(i, i.names=names(l), strict=FALSE))))
rm(l)
```

---

Description

Package \texttt{ref} implements references for S (R/S+). Function \texttt{ref} creates references. For a memory efficient wrapper to matrixes and data.frames which allows nested subsetting see \texttt{refdata}

Usage

\texttt{ref(name, loc = parent.frame())}

Arguments

- \texttt{name} name of an (existing) object to be referenced
- \texttt{loc} location of the referenced object, i.e. an environment in R or a frame in S+

---

\texttt{ref} creating references
Details

In S (R/S+) parameters are passed by value and not by reference. When passing big objects, e.g. in recursive algorithms, this can quickly eat up memory. The functions of package ref allow to pass references in function calls. The implementation is purely S and should work in R and S+. Existence of the referenced object is not checked by function ref. Usually as.ref is more convenient and secure to use. There is also a print method for references.

Value

a list with

name name of the referenced object
loc location of the referenced object, i.e. an environment in R or a frame in S+
and class "ref"

WARNING

Usually functions in S have no side-effects except for the main effect of returning something. Working with references circumvents this programming style and can have considerable side-effects. You are using it at your own risk.

R 1.8 WARNING

Changing parts of referenced objects has been slowed down by order of magnitudes since R version 1.8, see performance test examples on the help page for deref. Hopefully the old performance can be restored in future versions.

S+ WARNING

Package ref should generally work under R and S+. However, when changing very small parts of referenced objects, using references under S+ might be inefficient (very slow with high temporary memory requirements).

Historical remarks

This package goes back to an idea submitted April 9th 1997 and code offered on August 17th 1997 on s-news. The idea of implementing references in S triggered an intense discussion on s-news. The status reached in 1997 can be summarized as follows:

1. **advantage** passing by reference can save memory compared to passing by value
2. **disadvantage** passing by reference is more dangerous than passing by value
3. **however** the implementation is purely in S, thus rather channels existing danger than adding new danger
4. **restriction** assigning to a subitted part of a referenced object was inefficient in S+ (was S+ version 3)
Due to the last restriction the code was never submitted as a mature library. Now in 2003 we have a stable version of R and astonishingly assigning to a sub)setted part of a referenced object can be implemented efficient. This shows what a great job the R core developers have done. In the current version the set of functions for references was dramatically simplified, the main differences to 1997 being the following:

1. **no idempotence** `deref` and `deref<-` now are a simple function and no longer are methods. This decision was made due to performance reasons. As a consequence, `deref()` no longer is idempotent: one has to know whether an object is a reference. Function `is.ref` provides a test.

2. **no write protection** The 1997 suggestion included a write protection attribute of references, allowing for read only references and allowing for references that could only be changed by functions that know the access code. Reasons for this: there is no need forreadonly references (due to copy on modify) and oop provides better mechanisms for security.

3. **no static variables** The suggestion made in 1997 did include an implementation of static variables realized as special cases of references with a naming convention which reduced the risk of name collisions in the 1997 practice of assigning to frame 0. Now R has namespaces and the oop approach of Henrik Bengtsson using environments is to be preferred over relatively global static objects.

**Note**

Using this type of references is fine for prototyping in a non-object oriented programming style. For bigger projects and safer programming you should consider the approach suggested by Henrik Bengtsson at [http://www.maths.lth.se/help/R/ImplementingReferences](http://www.maths.lth.se/help/R/ImplementingReferences) (announced to be released as package "oo" or "classes")

**Author(s)**

Jens Oehlschlägel

**See Also**

`as.ref`, `deref`, `deref<-`, `exists.ref`, `is.ref`, `print.ref`, `HanoiTower`

**Examples**

```r
v <- 1
r <- ref("v")
r
deref(r)
cat("For more examples see ?deref\n")
```
Description

Function `refdata` creates objects of class `refdata` which behave not totally unlike matrices or data.frames but allow for much more memory efficient handling.

Usage

```r
# -- usage for R CMD CHECK, see below for human readable version ---------------
refdata(x)
derefdata(x)
derefdata(x) <- value
  ## S3 method for class 'refdata'
x[i = NULL, j = NULL, drop = FALSE, ref = FALSE]
  ## S3 replacement method for class 'refdata'
x[i = NULL, j = NULL, ref = FALSE] <- value
  ## S3 method for class 'refdata'
dim(x)
  ## S3 method for class 'refdata'
dimnames(x)
  ## S3 method for class 'refdata'
row.names(x)
  ## S3 method for class 'refdata'
names(x)

# -- most important usage for human beings -------------------------------
# rd <- refdata(x)          # create reference
# derefdata(rd)            # retrieve original data
# derefdata(rd) <- value    # modify original data
# rd[]                     # get all (current) data
# rd[i, j]                 # get part of data
# rd[i, j, ref=TRUE]       # get new reference on part of data
# rd[i, j] <- value        # modify / create local copy
# rd[i, j, ref=TRUE] <- value # modify original data (respecting subsetting history)
# dim(rd)                  # dim of (subsetted) data
# dimnames(rd)             # dimnames of (subsetted) data
```

Arguments

- `x`: a matrix or data.frame or any other 2-dimensional object that has operators "[" and "[<-" defined
- `i`: row index
- `j`: col index
ref

FALSE by default. In subsetting: FALSE returns data, TRUE returns new ref-data object. In assignments: FALSE modifies a local copy and returns a refdata object embedding it, TRUE modifies the original.

drop

FALSE by default, i.e. returned data have always a dimension attribute. TRUE drops dimension in some cases, the exact result depends on whether a matrix or data.frame is embedded

value

some value to be assigned

Details

Refdata objects store 2D-data in one environment and index information in another environment. Derived refdata objects usually share the data environment but not the index environment.

The index information is stored in a standardized and memory efficient form generated by `optimal.index`. Thus refdata objects can be copied and subsetted and even modified without duplicating the data in memory.

Empty square bracket subsetting (`rd[]`) returns the data, square bracket subsetting (`rd[i, j]`) returns subsets of the data as expected.

An additional argument (`rd[i, j, ref=TRUE]`) allows to get a reference that stores the subsetting indices. Such a reference behaves transparently as if a smaller matrix/data.frame would be stored and can be subsetted again recursively. With ref=TRUE indices are always interpreted as row/col indices, i.e. `x[i]` and `x[cbind(i, j)]` are undefined (and raise stop errors).

Standard square bracket assignment (`rd[i, j] <- value`) creates a reference to a locally modified copy of the (potentially subsetted) data.

An additional argument (`rd[i, j, ref=TRUE] <- value`) allows to modify the original data, properly recognizing the subsetting history.

A method `dim(refdata)` returns the dim of the (indexed) data.

A `dimnames(refdata)` returns the dimnames of the (indexed) data.

Value

an object of class refdata (appended to class attributes of data), which is an empty list with two attributes

dat the environment where the data x and its dimension dim is stored

ind the environment where the indexes i, j and the effective subset size ni, nj is stored

Note

The refdata code is currently R only (not implemented for S+).

Please note the following differences to matrices and dataframes:

- `x[]` you need to write `x[]` instead of `x` in order to get all current data
- `drop=FALSE` by default `drop=FALSE` which gives consistent behaviour for matrices and data.frames.

You can use the `$- or `[[`-operator to extract single column vectors which are granted to be of a consistent data type. However, currently `$` and `[[` are only wrappers to `[]`. They might be performance tuned in later versions.
refdata

x[i] single index subsetting is not defined, use x[][i] instead, but beware of differences between matrices and dataframes

x[cbind()] matrix index subsetting is not defined, use x[][cbind(i, j)] instead

ref=TRUE parameter ref needs to be used sensibly to exploit the advantages of refdata objects

Author(s)
Jens Oehlschlägel

See Also
Extract, matrix, data.frame, optimal.index, ref

Examples

## Simple usage Example
x <- cbind(1:5, 5:1) # take a matrix or data frame
rx <- refdata(x) # wrap it into an refdata object
rx # see the autoprinting
rm(x) # delete original to save memory
rx[] # extract all data
rx[-1, ] # extract part of data
rx2 <- rx[-1, , ref=TRUE] # create refdata object referencing part of data
# (only index, no data is duplicated)
rx2 # compare autoprinting
rx2[] # extract 'all' data
rx2[-1, ] # extract part of (part of) data
cat("for more examples look the help pages\n")

## Not run:
# Memory saving demos
square.matrix.size <- 1000
recursion.depth.limit <- 10
non.referenced.matrix <- matrix(1:(square.matrix.size*square.matrix.size),
, nrow=square.matrix.size, ncol=square.matrix.size)
rownames(non.referenced.matrix) <- paste("a", seq(length=square.matrix.size), sep="")
colnames(non.referenced.matrix) <- paste("b", seq(length=square.matrix.size), sep="")
referenced.matrix <- refdata(non.referenced.matrix)
recurse.nonref <- function(m, depth.limit=10){
  x <- m[1,1] # need read access here to create local copy
gc()
cat("depth.limit=", depth.limit, " memory.size=", memsize.wrapper(), "\n", sep="")
if (depth.limit)
  Recall(m[-1, -1, drop=FALSE], depth.limit=depth.limit-1)
invisible()
}
recurse.ref <- function(m, depth.limit=10){
  x <- m[1,1] # read access, otherwise nothing happens
gc()
cat("depth.limit=", depth.limit, " memory.size=", memsize.wrapper(), "\n", sep="")
if (depth.limit)
Recall(m[-1, -1, ref=TRUE], depth.limit=depth.limit-1)
invisible()
}
gc()
memsic.wrapper()
recurse.ref(referenced.matrix, recursion.depth.limit)
gc()
memsic.wrapper()
recurse.nonref(non.referenced.matrix, recursion.depth.limit)
gc()
memsic.wrapper()
rm(recurse.nonref, recurse.ref, non.referenced.matrix,
  referenced.matrix, square.matrix.size, recursion.depth.limit)

## End(Not run)
cat("for even more examples look at regression.test.refdata()\n")
regression.test.refdata() # testing correctness of refdata functionality
Index

+Topic **manip**
  - optimal.index, 9
  - refdata, 13

+Topic **programming**
  - as.ref, 2
  - deref, 2
  - HanoiTower, 5
  - is.ref, 8
  - ref, 10
  - refdata, 13

+Topic **utilities**
  - optimal.index, 9
  - [.refdata(refdata), 13
  - [<-refdata(refdata), 13
  - [[.refdata(refdata), 13
  - [[<-refdata(refdata), 13
  - $.refdata(refdata), 13
  - $<-refdata(refdata), 13
  - as.ref, 2, 3, 8, 11, 12
  - assign, 3
  - class, 8
  - data.frame, 14, 15
  - deref, 2, 11, 12
  - deref<- (deref), 2
  - derefdata(refdata), 13
  - derefdata<- (refdata), 13
  - dim, 14
  - dim.reffdata(refdata), 13
  - dimnames, 14
  - dimnames.reffdata(refdata), 13
  - eval, 3
  - exists, 8
  - exists.ref, 12
  - exists.ref (is.ref), 8
  - Extract, 15
  - get, 3

HanoiTower, 5, 12

inherits, 8
is.ref, 8, 12

matrix, 14, 15
move.HanoiTower (HanoiTower), 5

names.refdata(refdata), 13
need.index (optimal.index), 9
new.env, 3

optimal.index, 9, 14, 15

plot.HanoiTower (HanoiTower), 5
posix.index (optimal.index), 9
print.HanoiTower (HanoiTower), 5
print.ref, 12
print.ref (ref), 10
print.refdata(refdata), 13

Recall, 5, 6
ref, 2, 3, 6, 8, 10, 15
refdata, 10, 13
row.names.refdata(refdata), 13

substitute, 3