## Package 'Rdsm'

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Title Threads Environment for R

**Description** Provides a threads-type programming environment for R. The package gives the R programmer the clearer, more concise shared memory world view, and in some cases gives superior performance as well. In addition, it enables parallel processing on very large, out-of-core matrices.

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Imports bigmemory (>= 4.0.0), parallel Suggests synchronicity LazyLoad no License GPL (>= 2) Repository CRAN

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Rdsm-package

#### Description

This package provides a parallel shared-memory programming paradigm for R, very similar to threaded programming in C/C++. This enables the programmer to write simpler, clearer code. Furthermore, in some applications this package produces significantly faster code, compared to versions written for other parallel R libraries. It also allows placing very large matrices in secondary storage, while treating them as being in shared memory.

#### Details

Package:	Rdsm
Type:	Package
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List of functions:

Built-in variables accessible by the threads, at the worker nodes:

## 13

```
myinfo$nwrkrs: total number of threads
myinfo$id: this thread's ID number
```

To run, set up a cluster via the **parallel** package); we'll refer to the R process from which this is done as the manager; the processes running in the cluster will be called workers. Create the application's shared variables from the manager, using mgrmakevar(). Launch the worker threads, again from the manager, by the **parallel** call clusterEvalQ() or clusterCall(). One typically codes so that the results are in shared variables. See examples below, and more in the examples/ directory in this distribution.

The shared variables are read to/written by any of the workers and the manager. In fact, while an **Rdsm** application is running, other R processes on the same machine (or a different machine sharing the same file system, if the variables are filebacked) can access the shared variables. See the file ExternalAccess.txt in the doc/ directory.

**Rdsm** uses the **bigmemory** library to store its shared variables. Though the latter can work on a (physical) cluster of several machines sharing a file system, **Rdsm** does not run on such systems at this time.

Further documentation in the doc/ directory.

#### Author(s)

Norm Matloff <matloff@cs.ucdavis.edu>

#### See Also

mgrinit, mgrmakevar, mgrmakelock, barr, rdsmlock, rdsmunlock, getidxs, getmatrix

#### Examples

```
library(parallel)
c2 <- makeCluster(2) # form 2-thread Snow cluster
mgrinit(c2) # initialize Rdsm
mgrmakevar(c2,"m",2,2) # make a 2x2 shared matrix
m[,] <- 3 # 2x2 matrix of all 3s</pre>
# example of shared memory:
# at each thread, set id to Rdsm built-in ID variable for that thread
clusterEvalQ(c2,id <- myinfo$id)</pre>
clusterEvalQ(c2,m[1,id] <- id^2) # assignment executed by each thread
m[,] # top row of m should now be (1,4)
# matrix multiplication; the product u %*% v is computed, product
# placed in w
# note again: mmul() call will be executed by each thread
mmul <- function(u,v,w) {</pre>
   require(parallel)
   # decide which rows of u this thread will work on
   myidxs <- splitIndices(nrow(u),myinfo$nwrkrs)[[myinfo$id]]</pre>
   # multiply this thread's part of u with v, placing the product in the
```

```
# corresponding part of w
w[myidxs,] <- u[myidxs,] %*% v[,]
invisible(0)
}
# create test matrices
mgrmakevar(c2,"a",6,2)
mgrmakevar(c2,"b",2,6)
mgrmakevar(c2,"c",6,6)
# give them values
a[,] <- 1:12
b[,] <- 1 # all 1s
clusterExport(c2,"mmul") # send mmul() to the threads
clusterEvalQ(c2,mmul(a,b,c)) # run the threads
c[,] # check results</pre>
```

barr

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Barrier operation.

## Description

Standard barrier operation.

#### Usage

barr()

### Details

Standard barrier operation, to ensure that work done by one thread is ready before other threads make use of it. When a thread executes barr(), it will block until all threads have executed it.

#### Author(s)

Norm Matloff

getidxs

Parallelizing work assignment.

## Description

Assigns to an **Rdsm** thread its portion of a set of indices, for the purpose of partitioning work to the threads.

#### Usage

getidxs(m)

#### getmatrix

#### Arguments

т

The sequence 1:m will be partitioned, and one portion will be assigned to the calling thread.

## Details

The range 1:m will be partitioned into r subranges, the i-th of which will be used by thread i to determine which work that thread has been assigned.

## Value

The subrange assigned to the invoking thread.

#### Author(s)

Norm Matloff

getmatrix

Referencing a matrix via different forms.

## Description

Returns a matrix, whether requested via an R matrix, bigmemory object, bigmemory descriptor or quoted name form.

#### Usage

getmatrix(m)

## Arguments

m

Specification of the matrix, as either an R matrix, bigmemory object, bigmemory descriptor or quoted name.

## Details

This utility function enables writing general **Rdsm** code, specifying a matrix via different forms.

## Value

The requested matrix.

#### Author(s)

Norm Matloff

loadex

## Examples

```
library(parallel)
c2 <- makeCluster(2)
mgrmakevar(c2,"u",2,2)  # u is a 2x2 bigmemory matrix
u[] <- 8  # fill u with 8s
u[]  # prints a 2x2 matrix of 8s
v <- getmatrix(u)  # get u and assign it to v
# u and v are both addresses, pointing to the same memory location
v[]  # prints all 8s
v[2,1] <- 3
v[]  # prints three 8s and a 3
u[]  # prints three 8s and a 3
w <- getmatrix("u")  # w will also be a copy of u
w[]  # same as u</pre>
```

loadex

Sources example files

#### Description

Facilitates learning a new package, by sourcing any specified example file for an installed package.

#### Usage

loadex(pkg,exfile=NA,subdir="examples")

#### Arguments

pkg	Package name, quoted.
exfile	Desired example file name, if any, quoted.
subdir	Subdirectory name of the examples directory in the package.

#### Details

Loads the example file exfile, from the subdir directory within the tree wherepkg is installed. If exfile = NA, the names of the files are returned, without any loading.

## Value

See the case exfile = NA above.

#### Author(s)

Norm Matloff

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makebarr

## Description

Creates an Rdsm barrier.

## Usage

makebarr(cls,boost=F,barrback=F)

## Arguments

cls	The <b>snow</b> cluster.
boost	Locks type. See mgrinit.
barrback	If TRUE, the count/sense variables related to the barrier will be placed in back-
	ing store.

## Details

Run this from the manager (the R process from which you create the cluster) if you need a barrier. Only one barrier is allowed in an **Rdsm** program (but multiple calls are allowed). It is accessible from application code only through barr().

## Author(s)

Norm Matloff

mgrinit

Initialize Rdsm

## Description

Initializes Rdsm on the given parallel cluster.

## Usage

```
mgrinit(cls,boost=F,barrback=F)
```

## Arguments

cls	The <b>parallel</b> cluster.
boost	Lock functions. If TRUE, boostlock() and boostunlock() are used; otherwise backlock() and backunlock().
barrback	If TRUE, the count/sense variables associated with the barrier will be placed in backing store.

## Details

Run this from the manager (the R process from which you create the cluster), before creating the shared variables with mgrmakevar. The initialization need be done only once for the life of the cluster.

If you put shared variables in backing store (barrback = TRUE in mgrmakevar()), or if you are on a Windows platform, you must have boost = FALSE.

## Author(s)

Norm Matloff

mgrmakelock Create an Rdsm lock.

## Description

Creates an Rdsm lock.

#### Usage

mgrmakelock(cls,lockname,boost=F)

#### Arguments

cls	The <b>parallel</b> cluster.
lockname	Name of the lock, quoted.
boost	If TRUE, boost locks will be used.

## Details

Run this from the manager (the R process from which you create the cluster) if you need a lock. The lock is created, lockable/unlockable by all threads. If boost is TRUE, The variable will be of class boost.mutex; see the library **synchronicity** for details.

## Author(s)

Norm Matloff

mgrmakevar

## Description

Creates an Rdsm shared variable.

## Usage

mgrmakevar(cls,varname,nr,nc,vartype="double",fs=FALSE,mgrcpy=TRUE,savedesc=TRUE)

## Arguments

cls	The <b>parallel</b> cluster.
varname	Name of the shared variable, quoted. (The variable must be a matrix, though it could be $1x1$ etc.)
nr	Number of rows in the variable.
nc	Number of columns in the variable.
vartype	Atomic R type of the variable, quoted, "double" by default.
fs	Place in backing store? FALSE by default.
mgrcpy	Place a copy of the shared variable on the manager node? TRUE by default.
savedesc	Save the <b>bigmemory</b> descriptor for this variable on disk.

#### Details

Run this from the manager (the R process from which you create the cluster). The shared variable will be created, readable/writable from all threads. The variable will be of class big.matrix; see the library **bigmemory** for details.

#### Author(s)

Norm Matloff

rdsmlock

Lock/unlock operations.

## Description

Lock/unlock operations to avoid race conditions among the threads.

## Usage

```
rdsmlock(lck)
rdsmunlock(lck)
```

#### Arguments

lck

Lock name, quoted.

## Details

Standard lock/unlock operations from the threaded coding world. When one thread executes rdsmlock(), any other thread attempting to do so will block until the first thread executes rdsmunlock(). If a thread does rdsmlock() on an unlocked lock, the thread call immediately returns and the thread continues.

These functions are set in the call to mgrinit() via the argument boost to either boostlock and boostunlock() or backlock and backunlock(), depending on whether you set boost to TRUE or FALSE. respectively.

Code should be written so that locks are used as sparingly as possible, since they detract from performance.

#### Author(s)

Norm Matloff

## Examples

```
## Not run:
# unreliable function
s <- function(n) {</pre>
   for (i in 1:n) {
      tot[1,1] <- tot[1,1] + 1</pre>
   }
}
library(parallel)
c2 <- makeCluster(2)</pre>
clusterExport(c2,"s")
mgrinit(c2)
mgrmakevar(c2,"tot",1,1)
tot[1,1] <- 0
clusterEvalQ(c2,s(1000))
tot[1,1] # should be 2000, but likely far from it
s1 <- function(n) {</pre>
   require(Rdsm)
   for (i in 1:n) {
      rdsmlock("totlock")
      tot[1,1] <- tot[1,1] + 1
      rdsmunlock("totlock")
   }
}
mgrmakelock(c2,"totlock")
tot[1,1] <- 0
```

#### readsync

clusterExport(c2,"s1")
clusterEvalQ(c2,s1(1000))
tot[1,1] # will print out 2000, the correct number
## End(Not run)

readsync

Syncing file-backed variables.

#### Description

Actions to propagate changes to file-backed Rdsm variables across a shared file system.

#### Usage

readsync(varname)
writesync(varname)

#### Arguments

varname Name of the **Rdsm** variable, quoted.

## Details

This feature should be considered experimental, with poor performance and portability.

Suppose we have an **Rdsm** variable x which is in backing store (fs = TRUE in call to mgrmakevar()), and that we on a shared file system. (These functions are not needed if all threads are on the same machine.) When one **Rdsm** thread writes to x, the question here is when the updated value for x is visible to other **Rdsm** threads.

The answer may depend on the underlying file system. The functions readsync() and writesync() force the updates across the network. Normally one would call readsync() following rdsmlock() and call writesync() just before calling rdsmunlock().

These should work on systems with "close-to-open cache coherency," as with the Network File System (NFS). On some systems, these functions should be unnecessary.

## Value

None.

#### Author(s)

Norm Matloff

## Examples

```
library(parallel)
c2 <- makeCluster(2)
mgrinit(c2)
mgrmakevar(c2,"x",1,1,fs=TRUE)
clusterEvalQ(c2,me <- myinfo$id)
clusterEvalQ(c2,if (me==1) x[1,1] <- 3)
# force update on network
clusterEvalQ(c2,if (me==1) writesync("x"))
clusterEvalQ(c2,if (me==2) readsync("x"))
clusterEvalQ(c2,if (me==2) x[1,1]) # should be 3
clusterEvalQ(c2,if (me==2) x[1,1] <- 8)
clusterEvalQ(c2,if (me==2) writesync("x"))
clusterEvalQ(c2,if (me==1) readsync("x"))
clusterEvalQ(c2,x[1,1]) # both should yield 8
```

stoprdsm

Barrier operation.

## Description

Standard barrier operation.

## Usage

stoprdsm(cls)

#### Arguments

cls Cluster from the **parallel** package.

## Details

Shuts down the given parallel cluster, and removes any .desc files that had been created.

## Author(s)

Norm Matloff

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