

Package ‘DelayedArray’

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Title A unified framework for working transparently with on-disk and in-memory array-like datasets

Description Wrapping an array-like object (typically an on-disk object) in a DelayedArray object allows one to perform common array operations on it without loading the object in memory. In order to reduce memory usage and optimize performance, operations on the object are either delayed or executed using a block processing mechanism. Note that this also works on in-memory array-like objects like DataFrame objects (typically with Rle columns), Matrix objects, ordinary arrays and, data frames.

biocViews Infrastructure, DataRepresentation, Annotation, GenomeAnnotation

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BugReports <https://github.com/Bioconductor/DelayedArray/issues>

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SparseArraySeed-utils.R read_sparse_block.R
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AutoBlock-global-settings	
<i>Control the geometry of automatic blocks</i>	

Description

A family of utilities to control the automatic block size (or length) and shape.

Usage

```
getAutoBlockSize()
setAutoBlockSize(size=1e8)

getAutoBlockLength(type)

getAutoBlockShape()
setAutoBlockShape(shape=c("hypercube",
                           "scale",
                           "first-dim-grows-first",
                           "last-dim-grows-first"))
```

Arguments

size	The <i>auto block size</i> (automatic block size) in bytes. Note that, except when the type of the array data is "character" or "list", the size of a block is its length multiplied by the size of an array element. For example, a block of 500 x 1000 x 500 doubles has a length of 250 million elements and a size of 2 Gb (each double occupies 8 bytes of memory). The <i>auto block size</i> is set to 100 Mb at package startup and can be reset anytime to this value by calling <code>setAutoBlockSize()</code> with no argument.
type	A string specifying the type of the array data.
shape	A string specifying the <i>auto block shape</i> (automatic block shape). See makeCappedVolumeBox for a description of the supported shapes. The <i>auto block shape</i> is set to "hypercube" at package startup and can be reset anytime to this value by calling <code>setAutoBlockShape()</code> with no argument.

Details

block size != *block length*

block length = number of array elements in a block (i.e. `prod(dim(block))`).

block size = *block length* * size of the individual elements in memory.

For example, for an integer array, *block size* (in bytes) is going to be 4 x *block length*. For a numeric array `x` (i.e. `type(x) == "double"`), it's going to be 8 x *block length*.

In its current form, block processing in the **DelayedArray** package must decide the geometry of the blocks before starting the walk on the blocks. It does this based on several criteria. Two of them are:

- The *auto block size*: maximum size (in bytes) of a block once loaded in memory.
- The `type()` of the array (e.g. integer, double, complex, etc...)

The *auto block size* setting and `type(x)` control the maximum length of the blocks. Other criteria control their shape. So for example if you set the *auto block size* to 8GB, this will cap the length of the blocks to `2e9` if your `DelayedArray` object `x` is of type integer, and to `1e9` if it's of type double.

Note that this simple relationship between *block size* and *block length* assumes that blocks are loaded in memory as ordinary (a.k.a. dense) matrices or arrays. With sparse blocks, all bets are off. But the max block length is always taken to be the *auto block size* divided by `get_type_size(type())` whether the blocks are going to be loaded as dense or sparse arrays. If they are going to be loaded as sparse arrays, their memory footprint is very likely to be smaller than if they were loaded as dense arrays so this is safe (although probably not optimal).

It's important to keep in mind that the *auto block size* setting is a simple way for the user to put a cap on the memory footprint of the blocks. Nothing more. In particular it doesn't control the maximum amount of memory used by the block processing algorithm. Other variables can impact dramatically memory usage like parallelization (where more than one block is loaded in memory at any given time), what the algorithm is doing with the blocks (e.g. something like `blockApply(x, identity)` will actually load the entire array data in memory), what delayed operations are on `x`, etc... It would be awesome to have a way to control the maximum amount of memory used by a block processing algorithm as a whole but we don't know how to do that.

Value

`getAutoBlockSize`: The current *auto block size* in bytes as a single numeric value.

`setAutoBlockSize`: The new *auto block size* in bytes as an invisible single numeric value.

`getAutoBlockLength`: The *auto block length* as a single integer value.

`getAutoBlockShape`: The current *auto block shape* as a single string.

`setAutoBlockShape`: The new *auto block shape* as an invisible single string.

See Also

- [defaultAutoGrid](#) and family to create automatic grids to use for block processing of array-like objects.
- [blockApply](#) and family for convenient block processing of an array-like object.
- The [makeCappedVolumeBox](#) utility to make *capped volume boxes*.

Examples

```
getAutoBlockSize()

getAutoBlockLength("double")
getAutoBlockLength("integer")
getAutoBlockLength("logical")
getAutoBlockLength("raw")

m <- matrix(runif(600), ncol=12)
setAutoBlockSize(140)
getAutoBlockLength(type(m))
defaultAutoGrid(m)
lengths(defaultAutoGrid(m))
dims(defaultAutoGrid(m))

getAutoBlockShape()
setAutoBlockShape("scale")
defaultAutoGrid(m)
lengths(defaultAutoGrid(m))
dims(defaultAutoGrid(m))

## Reset the auto block size and shape to factory settings:
setAutoBlockSize()
setAutoBlockShape()
```

AutoGrid	<i>Create automatic grids to use for block processing of array-like objects</i>
----------	---

Description

We provide various utility functions to create grids that can be used for block processing of array-like objects:

- `defaultAutoGrid()` is the default *automatic grid maker*. It creates a grid that is suitable for block processing of the array-like object passed to it.
- `rowAutoGrid()` and `colAutoGrid()` are more specialized *automatic grid makers*, for the 2-dimensional case. They can be used to create a grid where the blocks are made of full rows or full columns, respectively.
- `defaultSinkAutoGrid()` is a specialized version of `defaultAutoGrid()` for creating a grid that is suitable for writing to a [RealizationSink](#) derivative while walking on it.

Usage

```
defaultAutoGrid(x, block.length=NULL, chunk.grid=NULL, block.shape=NULL)

## Two specialized "automatic grid makers" for the 2-dimensional case:
rowAutoGrid(x, nrow=NULL, block.length=NULL)
```

```
colAutoGrid(x, ncol=NULL, block.length=NULL)

## Replace default automatic grid maker with user-defined one:
getAutoGridMaker()
setAutoGridMaker(GRIDMAKER="defaultAutoGrid")

## A specialized version of defaultAutoGrid() to create an automatic
## grid on a RealizationSink derivative:
defaultSinkAutoGrid(sink, block.length=NULL, chunk.grid=NULL)
```

Arguments

x	An array-like or matrix-like object for defaultAutoGrid. A matrix-like object for rowAutoGrid and colAutoGrid.
block.length	The length of the blocks i.e. the number of array elements per block. By default the automatic block length (returned by <code>getAutoBlockLength(type(x))</code> , or <code>getAutoBlockLength(type(sink))</code> in the case of <code>defaultSinkAutoGrid()</code>) is used. Depending on how much memory is available on your machine, you might want to increase (or decrease) the automatic block length by adjusting the automatic block size with <code>setAutoBlockSize()</code> .
chunk.grid	The grid of physical chunks. By default <code>chunkGrid(x)</code> (or <code>chunkGrid(sink)</code> in the case of <code>defaultSinkAutoGrid()</code>) is used.
block.shape	A string specifying the shape of the blocks. See makeCappedVolumeBox for a description of the supported shapes. By default <code>getAutoBlockShape()</code> is used.
nrow	The number of rows of the blocks. The bottommost blocks might have less. See examples below.
ncol	The number of columns of the blocks. The rightmost blocks might have less. See examples below.
GRIDMAKER	The function to use as <i>automatic grid maker</i> , that is, the function that will be used by <code>blockApply()</code> and <code>blockReduce()</code> to make a grid when no grid is supplied via their grid argument. The function will be called on array-like object x and must return an ArrayGrid object, say grid, that is compatible with x i.e. such that <code>refdim(grid)</code> is identical to <code>dim(x)</code> . GRIDMAKER can be specified as a function or as a single string naming a function. It can be a user-defined function or a pre-defined grid maker like <code>defaultAutoGrid</code> , <code>rowAutoGrid</code> , or <code>colAutoGrid</code> . The <i>automatic grid maker</i> is set to <code>defaultAutoGrid</code> at package startup and can be reset anytime to this value by calling <code>setAutoGridMaker()</code> with no argument.
sink	A RealizationSink derivative.

Details

By default, primary block processing functions `blockApply()` and `blockReduce()` use the grid returned by `defaultAutoGrid(x)` to walk on the blocks of array-like object x. This can be changed with `setAutoGridMaker()`.

By default `sinkApply()` uses the grid returned by `defaultSinkAutoGrid(sink)` to walk on the viewports of `RealizationSink` derivative sink and write to them.

Value

`defaultAutoGrid`: An `ArrayGrid` object on reference array `x`. The grid elements define the blocks that will be used to process `x` by block. The grid is *optimal* in the sense that:

1. It's *compatible* with the grid of physical chunks a.k.a. *chunk grid*. This means that, when the chunk grid is known (i.e. when `chunkGrid(x)` is not NULL or `chunk.grid` is supplied), every block in the grid contains one or more *full* chunks. In other words, chunks never cross block boundaries.
2. Its *resolution* is such that the blocks have a length that is as close as possible to (but does not exceed) `block.length`. An exception is made when some chunks already have a length that is $\geq$ `block.length`, in which case the returned grid is the same as the chunk grid.

Note that the returned grid is regular (i.e. is a `RegularArrayGrid` object) unless the chunk grid is not regular (i.e. is an `ArbitraryArrayGrid` object).

`rowAutoGrid`: A `RegularArrayGrid` object on reference array `x` where the grid elements define blocks made of full rows of `x`.

`colAutoGrid`: A `RegularArrayGrid` object on reference array `x` where the grid elements define blocks made of full columns of `x`.

`defaultSinkAutoGrid`: Like `defaultAutoGrid` except that `defaultSinkAutoGrid` always returns a grid with a "first-dim-grows-first" shape (note that, unlike the former, the latter has no `block.shape` argument). The advantage of using a grid with a "first-dim-grows-first" shape in the context of writing to the viewports of a `RealizationSink` derivative is that such a grid is guaranteed to work with "linear write only" realization backends. See important notes about "Cross realization backend compatibility" in `?write_block` in the **S4Arrays** package for more information.

See Also

- `setAutoBlockSize` and `setAutoBlockShape` to control the geometry of automatic blocks.
- `blockApply` and family for convenient block processing of an array-like object.
- `ArrayGrid` in the **S4Arrays** package for the formal representation of grids and viewports.
- The `makeCappedVolumeBox` utility to make *capped volume boxes*.
- `chunkGrid`.
- `read_block` and `write_block` in the **S4Arrays** package.

Examples

```
## -----
## A VERSION OF sum() THAT USES BLOCK PROCESSING
## -----

block_sum <- function(a, grid) {
  sums <- lapply(grid, function(viewport) sum(read_block(a, viewport)))
  sum(unlist(sums))
}
```

```

## On an ordinary matrix:
m <- matrix(runif(600), ncol=12)
m_grid <- defaultAutoGrid(m, block.length=120)
sum1 <- block_sum(m, m_grid)
sum1

## On a DelayedArray object:
library(HDF5Array)
M <- as(m, "HDF5Array")
sum2 <- block_sum(M, m_grid)
sum2

sum3 <- block_sum(M, colAutoGrid(M, block.length=120))
sum3

sum4 <- block_sum(M, rowAutoGrid(M, block.length=80))
sum4

## Sanity checks:
sum0 <- sum(m)
stopifnot(identical(sum1, sum0))
stopifnot(identical(sum2, sum0))
stopifnot(identical(sum3, sum0))
stopifnot(identical(sum4, sum0))

## -----
## defaultAutoGrid()
## -----
grid <- defaultAutoGrid(m, block.length=120)
grid
as.list(grid) # turn the grid into a list of ArrayViewport objects
table(lengths(grid))
stopifnot(maxlength(grid) <= 120)

grid <- defaultAutoGrid(m, block.length=120,
                        block.shape="first-dim-grows-first")
grid
table(lengths(grid))
stopifnot(maxlength(grid) <= 120)

grid <- defaultAutoGrid(m, block.length=120,
                        block.shape="last-dim-grows-first")
grid
table(lengths(grid))
stopifnot(maxlength(grid) <= 120)

defaultAutoGrid(m, block.length=100)
defaultAutoGrid(m, block.length=75)
defaultAutoGrid(m, block.length=25)
defaultAutoGrid(m, block.length=20)
defaultAutoGrid(m, block.length=10)

```



```
## -----
## rowAutoGrid() AND colAutoGrid()
## -----
rowAutoGrid(m, nrow=10) # 5 blocks of 10 rows each
rowAutoGrid(m, nrow=15) # 3 blocks of 15 rows each plus 1 block of 5 rows
colAutoGrid(m, ncol=5)  # 2 blocks of 5 cols each plus 1 block of 2 cols

## See '?RealizationSink' for advanced examples of user-implemented
## block processing using colAutoGrid() and a realization sink.

## -----
## REPLACE DEFAULT AUTOMATIC GRID MAKER WITH USER-DEFINED ONE
## -----
getAutoGridMaker()
setAutoGridMaker(function(x) colAutoGrid(x, ncol=5))
getAutoGridMaker()

blockApply(m, function(block) currentViewport())

## Reset automatic grid maker to factory settings:
setAutoGridMaker()
```

blockApply

blockApply() and family

Description

A family of convenience functions to walk on the blocks of an array-like object and process them.

Usage

Main looping functions:

```
blockApply(x, FUN, ..., grid=NULL, as.sparse=FALSE,
           BPPARAM=getAutoBPPARAM(), verbose=NA)
```

```
blockReduce(FUN, x, init, ..., BREAKIF=NULL, grid=NULL, as.sparse=FALSE,
            verbose=NA)
```

Lower-level looping functions:

```
gridApply(grid, FUN, ..., BPPARAM=getAutoBPPARAM(), verbose=NA)
gridReduce(FUN, grid, init, ..., BREAKIF=NULL, verbose=NA)
```

Retrieve grid context for the current block/viewport:

```
effectiveGrid(envir=parent.frame(2))
currentBlockId(envir=parent.frame(2))
currentViewport(envir=parent.frame(2))
```

Get/set automatic parallel back-end:

```
getAutoBPPARAM()
setAutoBPPARAM(BPPARAM=NULL)

## For testing/debugging callback functions:
set_grid_context(effective_grid, current_block_id, envir=parent.frame(1))
```

Arguments

x	An array-like object, typically a DelayedArray object or derivative.
FUN	For <code>blockApply</code> and <code>blockReduce</code>, FUN is the callback function to apply to each block of data in x. More precisely, FUN will be called on each block of data in x defined by the grid used to walk on x. IMPORTANT: If <code>as.sparse</code> is set to FALSE, all blocks will be passed to FUN as ordinary arrays. If it's set to TRUE, they will be passed as SparseArraySeed objects. If it's set to NA, then <code>is_sparse(x)</code> determines how they will be passed to FUN. For <code>gridApply()</code> and <code>gridReduce()</code>, FUN is the callback function to apply to each <code>**viewport**</code> in grid. Beware that FUN must take at least <code>**two**</code> arguments for <code>blockReduce()</code> and <code>gridReduce()</code>. More precisely: • <code>blockReduce()</code> will perform <code>init <- FUN(block, init, ...)</code> on each block, so FUN must take at least arguments <code>block</code> and <code>init</code>. • <code>gridReduce()</code> will perform <code>init <- FUN(viewport, init, ...)</code> on each viewport, so FUN must take at least arguments <code>viewport</code> and <code>init</code>. In both cases, the exact names of the two arguments doesn't really matter. Also FUN is expected to return a value of the same type as its 2nd argument (<code>init</code>).
...	Additional arguments passed to FUN.
grid	The grid used for the walk, that is, an ArrayGrid object that defines the blocks (or viewports) to walk on. For <code>blockApply()</code> and <code>blockReduce()</code> the supplied grid must be compatible with the geometry of x. If not specified, an automatic grid is used. By default <code>defaultAutoGrid(x)</code> is called to create an automatic grid. The <i>automatic grid maker</i> can be changed with <code>setAutoGridMaker()</code>. See <code>?setAutoGridMaker</code> for more information.
as.sparse	Passed to the internal calls to <code>read_block</code> . See <code>?read_block</code> in the S4Arrays package for more information.
BPPARAM	A NULL, in which case blocks are processed sequentially, or a BiocParallelParam instance (from the BiocParallel package), in which case they are processed in parallel. The specific BiocParallelParam instance determines the parallel back-end to use. See <code>?BiocParallelParam</code> in the BiocParallel package for more information about parallel back-ends.
verbose	Whether block processing progress should be displayed or not. If set to NA (the default), verbosity is controlled by <code>DelayedArray:::get_verbose_block_processing()</code> . Setting <code>verbose</code> to TRUE or FALSE overrides this.

init	The value to pass to the first call to FUN(block, init) (or FUN(viewport, init)) when blockReduce() (or gridReduce()) starts the walk. Note that blockReduce() and gridReduce() always operate sequentially.
BREAKIF	An optional callback function that detects a break condition. Must return TRUE or FALSE. At each iteration blockReduce() (and gridReduce()) will call it on the result of init <- FUN(block, init) (on the result of init <- FUN(viewport, init) for gridReduce()) and exit the walk if BREAKIF(init) returned TRUE.
envir	Do not use (unless you know what you are doing).
effective_grid, current_block_id	See Details below.

Details

effectiveGrid(), currentBlockId(), and currentViewport() return the "grid context" for the block/viewport being currently processed. By "grid context" we mean:

- The *effective grid*, that is, the user-supplied grid or defaultAutoGrid(x) if the user didn't supply any grid.
- The *current block id* (a.k.a. block rank).
- The *current viewport*, that is, the [ArrayViewport](#) object describing the position of the current block w.r.t. the effective grid.

Note that effectiveGrid(), currentBlockId(), and currentViewport() can only be called (with no arguments) from ****within**** the callback functions FUN and/or BREAKIF passed to blockApply() and family.

If you need to be able to test/debug your callback function as a standalone function, set an arbitrary *effective grid* and *current block id* by calling

```
set_grid_context(effective_grid, current_block_id)
```

****right before**** calling the callback function.

Value

For blockApply() and gridApply(), a list with one list element per block/viewport visited.

For blockReduce() and gridReduce(), the result of the last call to FUN.

For effectiveGrid(), the grid ([ArrayGrid](#) object) being effectively used.

For currentBlockId(), the id (a.k.a. rank) of the current block.

For currentViewport(), the viewport ([ArrayViewport](#) object) of the current block.

See Also

- [defaultAutoGrid](#) and family to create automatic grids to use for block processing of array-like objects.
- [ArrayGrid](#) in the **S4Arrays** package for the formal representation of grids and viewports.
- [read_block](#) and [write_block](#) in the **S4Arrays** package.
- [MulticoreParam](#), [SnowParam](#), and [bpparam](#), from the **BiocParallel** package.
- [DelayedArray](#) objects.

Examples

```

m <- matrix(1:60, nrow=10)
m_grid <- defaultAutoGrid(m, block.length=16, block.shape="hypercube")

## -----
## blockApply()
## -----
blockApply(m, identity, grid=m_grid)
blockApply(m, sum, grid=m_grid)

blockApply(m, function(block) {block + currentBlockId()*1e3}, grid=m_grid)
blockApply(m, function(block) currentViewport(), grid=m_grid)
blockApply(m, dim, grid=m_grid)

## The grid does not need to be regularly spaced:
a <- array(runif(8000), dim=c(25, 40, 8))
a_tickmarks <- list(c(7L, 15L, 25L), c(14L, 22L, 40L), c(2L, 8L))
a_grid <- ArbitraryArrayGrid(a_tickmarks)
a_grid
blockApply(a, function(block) sum(log(block + 0.5)), grid=a_grid)

## See block processing in action:
blockApply(m, function(block) sum(log(block + 0.5)), grid=m_grid,
           verbose=TRUE)

## Use parallel evaluation:
library(BiocParallel)
if (.Platform$OS.type != "windows") {
  BPPARAM <- MulticoreParam(workers=4)
} else {
  ## MulticoreParam() is not supported on Windows so we use
  ## SnowParam() on this platform.
  BPPARAM <- SnowParam(4)
}
blockApply(m, function(block) sum(log(block + 0.5)), grid=m_grid,
           BPPARAM=BPPARAM, verbose=TRUE)
## Note that blocks can be visited in any order!

## -----
## blockReduce()
## -----
FUN <- function(block, init) anyNA(block) || init
blockReduce(FUN, m, init=FALSE, grid=m_grid, verbose=TRUE)

m[10, 1] <- NA
blockReduce(FUN, m, init=FALSE, grid=m_grid, verbose=TRUE)

## With early bailout:
blockReduce(FUN, m, init=FALSE, BREAKIF=identity, grid=m_grid,
           verbose=TRUE)

## Note that this is how the anyNA() method for DelayedArray objects is

```

```
## implemented.
```

chunkGrid	<i>chunkGrid</i>
-----------	------------------

Description

chunkGrid and chunkdim are internal generic functions not aimed to be used directly by the user.

Usage

```
chunkGrid(x)  
chunkdim(x)
```

Arguments

x	An array-like object.
---	-----------------------

Details

Coming soon...

Value

chunkGrid returns NULL or an [ArrayGrid](#) object defining a grid on reference array x.

chunkdim returns NULL or the chunk dimensions in an integer vector parallel to `dim(x)`.

See Also

- [defaultAutoGrid](#) and family to create automatic grids to use for block processing of array-like objects.
- [DelayedArray](#) objects.
- [ArrayGrid](#) in the **S4Arrays** package for the formal representation of grids and viewports.

Examples

```
## Coming soon...
```

compat

Functions and classes that have moved to S4Arrays

Description

Some functions and classes that used to be defined in the **DelayedArray** package have been moved to the new **S4Arrays** package in BioC 3.17. The corresponding symbols are still exported by the **DelayedArray** package for backward compatibility with existing code.

WARNING: This is a temporary situation only. Packages that import these symbols from **DelayedArray** must be modified to import them from **S4Arrays** instead.

These symbols are actually documented in the **S4Arrays** package. See:

- `S4Arrays::t.Array`
- `S4Arrays::makeNindexFromArrayViewport`
- `S4Arrays::ArrayGrid`
- `S4Arrays::DummyArrayGrid`
- `S4Arrays::RegularArrayGrid`
- `S4Arrays::ArbitraryArrayGrid`
- `S4Arrays::extract_array`
- `S4Arrays::is_sparse`
- `S4Arrays::read_block`
- `S4Arrays::write_block`

ConstantArray

A DelayedArray subclass that contains a constant value

Description

A [DelayedArray](#) subclass to efficiently mimic an array containing a constant value, without actually creating said array in memory.

Usage

```
## Constructor function:
ConstantArray(dim, value=NA)
```

Arguments

dim	The dimensions (specified as an integer vector) of the ConstantArray object to create.
value	Vector (atomic or list) of length 1, containing the value to fill the matrix.

Details

This class allows us to efficiently create arrays containing a single value. For example, we can create matrices full of NA values, to serve as placeholders for missing assays when combining SummarizedExperiment objects.

Value

A ConstantArray (or ConstantMatrix) object. (Note that ConstantMatrix extends ConstantArray.)

Author(s)

Aaron Lun

See Also

- [DelayedArray](#) objects.
- [DelayedArray-utils](#) for common operations on [DelayedArray](#) objects.
- [RleArray](#) objects for representing in-memory Run Length Encoded array-like datasets.

Examples

```
## This would ordinarily take up 8 TB of memory:
CM <- ConstantArray(c(1e6, 1e6), value=NA_real_)
CM

CM2 <-ConstantArray(c(4, 1e6), value=55)
rbind(CM, CM2)
```

DelayedAbind-class *DelayedAbind objects*

Description

NOTE: This man page is about [DelayedArray](#) internals and is provided for developers and advanced users only.

The DelayedAbind class provides a formal representation of a *delayed* abind() operation. It is a concrete subclass of the [DelayedNaryOp](#) virtual class, which itself is a subclass of the [DelayedOp](#) virtual class:

```
DelayedOp
  ^
  |
DelayedNaryOp
  ^
  |
DelayedAbind
```


Examples

```
## DelayedAbind extends DelayedNaryOp which extends DelayedOp:
extends("DelayedAbind")

## -----
## BASIC EXAMPLE
## -----
m1 <- matrix(101:128, ncol=4)
m2 <- matrix(runif(16), ncol=4)
M1 <- DelayedArray(m1)
M2 <- DelayedArray(m2)
showtree(M1)
showtree(M2)

M3 <- rbind(M1, M2)
showtree(M3)
class(M3@seed)      # a DelayedAbind object

M4 <- cbind(t(M1), M2)
showtree(M4)
class(M4@seed)      # a DelayedAbind object

## -----
## PROPAGATION OF SPARSITY
## -----
## DelayedAbind objects always propagate sparsity (granted that all the
## input arrays are sparse).

sm1 <- sparseMatrix(i=c(1, 1, 7, 7), j=c(1, 4, 1, 4),
                    x=c(11, 14, 71, 74), dims=c(7, 4))
SM1 <- DelayedArray(sm1)
sm2 <- sparseMatrix(i=c(1, 1, 4, 4), j=c(1, 4, 1, 4),
                    x=c(11, 14, 41, 44), dims=c(4, 4))
SM2 <- DelayedArray(sm2)
showtree(SM1)
showtree(SM2)
is_sparse(SM1)      # TRUE
is_sparse(SM2)      # TRUE

SM3 <- rbind(SM1, SM2)
showtree(SM3)
class(SM3@seed)     # a DelayedAbind object
is_sparse(SM3@seed) # TRUE

SM4 <- cbind(SM2, t(SM1))
showtree(SM4)
class(SM4@seed)     # a DelayedAbind object
is_sparse(SM4@seed) # TRUE

M5 <- rbind(SM2, M1) # 2nd input array is not sparse!
showtree(M5)
class(M5@seed)      # a DelayedAbind object
```

```

is_sparse(M5@seed)    # FALSE

## -----
## SANITY CHECKS
## -----
stopifnot(class(M3@seed) == "DelayedAbind")
stopifnot(class(M4@seed) == "DelayedAbind")
stopifnot(class(SM3@seed) == "DelayedAbind")
stopifnot(is_sparse(SM3@seed))
stopifnot(class(SM4@seed) == "DelayedAbind")
stopifnot(is_sparse(SM4@seed))
stopifnot(class(M5@seed) == "DelayedAbind")
stopifnot(!is_sparse(M5@seed))

```

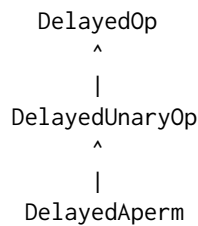
DelayedAperm-class	<i>DelayedAperm objects</i>
--------------------	-----------------------------

Description

NOTE: This man page is about [DelayedArray](#) internals and is provided for developers and advanced users only.

The DelayedAperm class provides a formal representation of a *delayed "extended aperm()" operation*, that is, of a delayed [aperm\(\)](#) that can drop and/or add *ineffective* dimensions. Note that since only *ineffective* dimensions (i.e. dimensions with an extent of 1) can be dropped or added, the length of the output array is guaranteed to be the same as the length of the input array.

DelayedAperm is a concrete subclass of the [DelayedUnaryOp](#) virtual class, which itself is a subclass of the [DelayedOp](#) virtual class:



DelayedAperm objects are used inside a [DelayedArray](#) object to represent the *delayed "extended aperm()" operations* carried by the object. They're never exposed to the end user and are not intended to be manipulated directly.

Usage

```

## S4 method for signature 'DelayedAperm'
is_noop(x)

## S4 method for signature 'DelayedAperm'

```

```
summary(object, ...)
```

```
## ~ ~ ~ Seed contract ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~
```

```
## S4 method for signature 'DelayedAperm'
```

```
dim(x)
```

```
## S4 method for signature 'DelayedAperm'
```

```
dimnames(x)
```

```
## S4 method for signature 'DelayedAperm'
```

```
extract_array(x, index)
```

```
## ~ ~ ~ Propagation of sparsity ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~
```

```
## S4 method for signature 'DelayedAperm'
```

```
is_sparse(x)
```

```
## S4 method for signature 'DelayedAperm'
```

```
OLD_extract_sparse_array(x, index)
```

Arguments

x, object	A DelayedAperm object.
index	See <code>?extract_array</code> in the S4Arrays package for a description of the index argument.
...	Not used.

See Also

- `DelayedOp` objects.
- `showtree` to visualize the nodes and access the leaves in the tree of delayed operations carried by a `DelayedArray` object.
- `extract_array` in the **S4Arrays** package.
- `OLD_extract_sparse_array`.

Examples

```
## DelayedAperm extends DelayedUnaryOp which extends DelayedOp:
extends("DelayedAperm")

## -----
## BASIC EXAMPLES
## -----
a0 <- array(1:20, dim=c(1, 10, 2))
A0 <- DelayedArray(a0)
showtree(A0)

A <- aperm(A0, perm=c(2, 3, 1))
```

```

showtree(A)
class(A@seed)      # a DelayedAperm object

M1 <- drop(A0)
showtree(M1)
class(M1@seed)      # a DelayedAperm object

M2 <- t(M1)
showtree(M2)
class(M2@seed)      # a DelayedAperm object

## -----
## PROPAGATION OF SPARSITY
## -----
## DelayedAperm objects always propagate sparsity.

sa0 <- as(a0, "SparseArraySeed")
SA0 <- DelayedArray(sa0)
showtree(SA0)
is_sparse(SA0)      # TRUE

SA <- aperm(SA0, perm=c(2, 3, 1))
showtree(SA)
class(SA@seed)      # a DelayedAperm object
is_sparse(SA@seed)  # TRUE

## -----
## SANITY CHECKS
## -----
stopifnot(class(A@seed) == "DelayedAperm")
stopifnot(class(M1@seed) == "DelayedAperm")
stopifnot(class(M2@seed) == "DelayedAperm")
stopifnot(class(SA@seed) == "DelayedAperm")
stopifnot(is_sparse(SA@seed))

```

DelayedArray-class *DelayedArray objects*

Description

Wrapping an array-like object (typically an on-disk object) in a DelayedArray object allows one to perform common array operations on it without loading the object in memory. In order to reduce memory usage and optimize performance, operations on the object are either *delayed* or executed using a block processing mechanism.

Usage

```

DelayedArray(seed) # constructor function
type(x)

```

Arguments

seed	An array-like object.
x	Typically a DelayedArray object. More generally type() is expected to work on any array-like object (that is, any object for which dim(x) is not NULL), or any ordinary vector (i.e. atomic or non-atomic).

In-memory versus on-disk realization

To *realize* a DelayedArray object (i.e. to trigger execution of the delayed operations carried by the object and return the result as an ordinary array), call `as.array` on it. However this realizes the full object at once *in memory* which could require too much memory if the object is big. A big DelayedArray object is preferably realized *on disk* e.g. by calling `writeHDF5Array` on it (this function is defined in the **HDF5Array** package) or coercing it to an `HDF5Array` object with `as(x, "HDF5Array")`. Other on-disk backends can be supported. This uses a block processing strategy so that the full object is not realized at once in memory. Instead the object is processed block by block i.e. the blocks are realized in memory and written to disk one at a time. See `?writeHDF5Array` in the **HDF5Array** package for more information about this.

Accessors

DelayedArray objects support the same set of getters as ordinary arrays i.e. `dim()`, `length()`, and `dimnames()`. In addition, they support `type()`, `nseed()`, `seed()`, and `path()`.

`type()` is the DelayedArray equivalent of `typeof()` (or `storage.mode()`) for ordinary arrays and vectors. Note that, for convenience and consistency, `type()` also supports ordinary arrays and vectors. It should also support any array-like object, that is, any object `x` for which `dim(x)` is not NULL.

`dimnames()`, `seed()`, and `path()` also work as setters.

Subsetting

A DelayedArray object can be subsetted with `[]` like an ordinary array, but with the following differences:

- *N-dimensional single bracket subsetting* (i.e. subsetting of the form `x[i_1, i_2, ..., i_n]` with one (possibly missing) subscript per dimension) returns a DelayedArray object where the subsetting is actually *delayed*. So it's a very light operation. One notable exception is when `drop=TRUE` and the result has only one dimension, in which case it is *realized* as an ordinary vector (atomic or list). Note that NAs in the subscripts are not supported.
- *1D-style single bracket subsetting* (i.e. subsetting of the form `x[i]`) only works if the subscript `i` is a numeric or logical vector, or a logical array-like object with the same dimensions as `x`, or a numeric matrix with one column per dimension in `x`. When `i` is a numeric vector, all the indices in it must be ≥ 1 and $\leq \text{length}(x)$. NAs in the subscripts are not supported. This is NOT a delayed operation (block processing is triggered) i.e. the result is *realized* as an ordinary vector (atomic or list). One exception is when `x` has only one dimension and `drop` is set to `FALSE`, in which case the subsetting is *delayed*.

Subsetting with `[[` is supported but only the 1D-style form of it at the moment, that is, subsetting of the form `x[[i]]` where `i` is a *single* numeric value ≥ 1 and $\leq \text{length}(x)$. It is equivalent to `x[i][[1]]`.

Subassignment to a DelayedArray object with `[<-` is also supported like with an ordinary array, but with the following restrictions:

- *N-dimensional subassignment* (i.e. subassignment of the form `x[i_1, i_2, ..., i_n] <- value` with one (possibly missing) subscript per dimension) only accepts a replacement value (a.k.a. right value) that is an array-like object (e.g. ordinary array, dgCMatix object, DelayedArray object, etc...) or an ordinary vector (atomic or list) of length 1.
- *1D-style subassignment* (a.k.a. 1D-style subassignment, that is, subassignment of the form `x[i] <- value`) only works if the subscript `i` is a logical DelayedArray object of the same dimensions as `x` and if the replacement value is an ordinary vector (atomic or list) of length 1.
- *Filling with a vector*, that is, subassignment of the form `x[] <- v` where `v` is an ordinary vector (atomic or list), is only supported if the length of the vector is a divisor of `nrow(x)`.

These 3 forms of subassignment are implemented as *delayed* operations so are very light.

Single value replacement (`x[[...]] <- value`) is not supported yet.

See Also

- [showtree](#) for [DelayedArray](#) accessors `nseed`, `seed`, and `path`.
- [realize](#) for realizing a DelayedArray object in memory or on disk.
- [blockApply](#) and family for convenient block processing of an array-like object.
- [DelayedArray-utils](#) for common operations on DelayedArray objects.
- [DelayedMatrix-utils](#) for common operations on DelayedMatrix objects.
- [DelayedArray-stats](#) for statistical functions on DelayedArray objects.
- [DelayedMatrix-stats](#) for DelayedMatrix row/col summarization.
- [ConstantArray](#) objects for mimicking an array containing a constant value, without actually creating said array in memory.
- [RleArray](#) objects for representing in-memory Run Length Encoded array-like datasets.
- [HDF5Array](#) objects in the **HDF5Array** package.
- [DataFrame](#) objects in the **S4Vectors** package.
- [array](#) objects in base R.

Examples

```
## -----
## A. WRAP AN ORDINARY ARRAY IN A DelayedArray OBJECT
## -----
a <- array(runif(1500000), dim=c(10000, 30, 5))
A <- DelayedArray(a)
A
## The seed of a DelayedArray object is **always** treated as a
## "read-only" object so will never be modified by the operations
## we perform on A:
stopifnot(identical(a, seed(A)))
type(A)
```

```

## N-dimensional single bracket subsetting:
m <- a[11:20 , 5, -3] # an ordinary matrix
M <- A[11:20 , 5, -3] # a DelayedMatrix object
stopifnot(identical(m, as.array(M)))

## 1D-style single bracket subsetting:
A[11:20]
A[A <= 1e-5]
stopifnot(identical(a[a <= 1e-5], A[A <= 1e-5]))

## Subassignment:
A[A < 0.2] <- NA
a[a < 0.2] <- NA
stopifnot(identical(a, as.array(A)))

A[2:5, 1:2, ] <- array(1:40, c(4, 2, 5))
a[2:5, 1:2, ] <- array(1:40, c(4, 2, 5))
stopifnot(identical(a, as.array(A)))

## Other operations:
crazy <- function(x) (5 * x[ , , 1] ^ 3 + 1L) * log(x[, , 2])
b <- crazy(a)
head(b)

B <- crazy(A) # very fast! (all operations are delayed)
B

cs <- colSums(b)
CS <- colSums(B)
stopifnot(identical(cs, CS))

## -----
## B. WRAP A DataFrame OBJECT IN A DelayedArray OBJECT
## -----
## Generate random coverage and score along an imaginary chromosome:
cov <- Rle(sample(20, 5000, replace=TRUE), sample(6, 5000, replace=TRUE))
score <- Rle(sample(100, nrun(cov), replace=TRUE), runLength(cov))

DF <- DataFrame(cov, score)
A2 <- DelayedArray(DF)
A2
seed(A2) # 'DF'

## Coercion of a DelayedMatrix object to DataFrame produces a DataFrame
## object with Rle columns:
as(A2, "DataFrame")
stopifnot(identical(DF, as(A2, "DataFrame")))

t(A2) # transposition is delayed so is very fast and memory-efficient
colSums(A2)

## -----
## C. AN HDF5Array OBJECT IS A (PARTICULAR KIND OF) DelayedArray OBJECT

```

```

## -----
library(HDF5Array)
A3 <- as(a, "HDF5Array") # write 'a' to an HDF5 file
A3
is(A3, "DelayedArray")  # TRUE
seed(A3)                 # an HDF5ArraySeed object

B3 <- crazy(A3)          # very fast! (all operations are delayed)
B3                       # not an HDF5Array object anymore because
                        # now it carries delayed operations

CS3 <- colSums(B3)
stopifnot(identical(cs, CS3))

## -----
## D. PERFORM THE DELAYED OPERATIONS
## -----
as(B3, "HDF5Array")      # "realize" 'B3' on disk

## If this is just an intermediate result, you can either keep going
## with B3 or replace it with its "realized" version:
B3 <- as(B3, "HDF5Array") # no more delayed operations on new 'B3'
seed(B3)
path(B3)

## For convenience, realize() can be used instead of explicit coercion.
## The current "automatic realization backend" controls where
## realization happens e.g. in memory if set to NULL or in an HDF5
## file if set to "HDF5Array":
D <- cbind(B3, exp(B3))
D
setAutoRealizationBackend("HDF5Array")
D <- realize(D)
D
## See '?setAutoRealizationBackend' for more information about
## "realization backends".

## -----
## E. MODIFY THE PATH OF A DelayedArray OBJECT
## -----
## This can be useful if the file containing the array data is on a
## shared partition but the exact path to the partition depends on the
## machine from which the data is being accessed.
## For example:

## Not run:
library(HDF5Array)
A <- HDF5Array("/path/to/lab_data/my_precious_data.h5")
path(A)

## Operate on A...
## Now A carries delayed operations.
## Make sure path(A) still works:
path(A)

```



```

## Save A:
save(A, file="A.rda")

## A.rda should be small (it doesn't contain the array data).
## Send it to a co-worker that has access to my_precious_data.h5.

## Co-worker loads it:
load("A.rda")
path(A)

## A is broken because path(A) is incorrect for co-worker:
A # error!

## Co-worker fixes the path (in this case this is better done using the
## dirname() setter rather than the path() setter):
dirname(A) <- "E:/other/path/to/lab_data"

## A "works" again:
A

## End(Not run)

## -----
## F. WRAP A SPARSE MATRIX IN A DelayedArray OBJECT
## -----
## Not run:
M <- 75000L
N <- 1800L
p <- sparseMatrix(sample(M, 9000000, replace=TRUE),
                  sample(N, 9000000, replace=TRUE),
                  x=runif(9000000), dims=c(M, N))
P <- DelayedArray(p)
P
p2 <- as(P, "sparseMatrix")
stopifnot(identical(p, p2))

## The following is based on the following post by Murat Tasan on the
## R-help mailing list:
## https://stat.ethz.ch/pipermail/r-help/2017-May/446702.html

## As pointed out by Murat, the straight-forward row normalization
## directly on sparse matrix 'p' would consume too much memory:
row_normalized_p <- p / rowSums(p^2) # consumes too much memory
## because the rowSums() result is being recycled (appropriately) into a
## *dense* matrix with dimensions equal to dim(p).

## Murat came up with the following solution that is very fast and
## memory-efficient:
row_normalized_p1 <- Diagonal(x=1/sqrt(Matrix::rowSums(p^2)))

## With a DelayedArray object, the straight-forward approach uses a
## block processing strategy behind the scene so it doesn't consume

```

```
## too much memory.

## First, let's see block processing in action:
DelayedArray::set_verbose_block_processing(TRUE)
## and check the automatic block size:
getAutoBlockSize()

row_normalized_P <- P / sqrt(DelayedArray::rowSums(P^2))

## Increasing the block size increases the speed but also memory usage:
setAutoBlockSize(2e8)
row_normalized_P2 <- P / sqrt(DelayedArray::rowSums(P^2))
stopifnot(all.equal(row_normalized_P, row_normalized_P2))

## Back to sparse representation:
DelayedArray::set_verbose_block_processing(FALSE)
row_normalized_p2 <- as(row_normalized_P, "sparseMatrix")
stopifnot(all.equal(row_normalized_p1, row_normalized_p2))

setAutoBlockSize() # reset automatic block size to factory settings

## End(Not run)
```

DelayedArray-stats *Statistical functions on DelayedArray objects*

Description

Statistical functions on [DelayedArray](#) objects.

All these functions are implemented as delayed operations.

Usage

```
## --- The Normal Distribution ----- ##

## S4 method for signature 'DelayedArray'
dnorm(x, mean=0, sd=1, log=FALSE)
## S4 method for signature 'DelayedArray'
pnorm(q, mean=0, sd=1, lower.tail=TRUE, log.p=FALSE)
## S4 method for signature 'DelayedArray'
qnorm(p, mean=0, sd=1, lower.tail=TRUE, log.p=FALSE)

## --- The Binomial Distribution --- ##

## S4 method for signature 'DelayedArray'
dbinom(x, size, prob, log=FALSE)
## S4 method for signature 'DelayedArray'
pbinom(q, size, prob, lower.tail=TRUE, log.p=FALSE)
```

```
## S4 method for signature 'DelayedArray'
qbinom(p, size, prob, lower.tail=TRUE, log.p=FALSE)

## --- The Poisson Distribution ---- ##

## S4 method for signature 'DelayedArray'
dpois(x, lambda, log=FALSE)
## S4 method for signature 'DelayedArray'
ppois(q, lambda, lower.tail=TRUE, log.p=FALSE)
## S4 method for signature 'DelayedArray'
qpois(p, lambda, lower.tail=TRUE, log.p=FALSE)

## --- The Logistic Distribution --- ##

## S4 method for signature 'DelayedArray'
dlogis(x, location=0, scale=1, log=FALSE)
## S4 method for signature 'DelayedArray'
plogis(q, location=0, scale=1, lower.tail=TRUE, log.p=FALSE)
## S4 method for signature 'DelayedArray'
qlogis(p, location=0, scale=1, lower.tail=TRUE, log.p=FALSE)
```

Arguments

`x`, `q`, `p` A [DelayedArray](#) object.
`mean`, `sd`, `log`, `lower.tail`, `log.p`, `size`, `prob`, `lambda`, `location`, `scale`
 See `?stats::dnorm`, `?stats::dbinom`, `?stats::dpois`, and `?stats::dlogis`,
 for a description of these arguments.

See Also

- [dnorm](#), [dbinom](#), [dpois](#), and [dlogis](#) in the **stats** package for the corresponding operations on ordinary arrays or matrices.
- [DelayedMatrix-stats](#) for [DelayedMatrix](#) row/col summarization.
- [DelayedArray](#) objects.
- [HDF5Array](#) objects in the **HDF5Array** package.
- [array](#) objects in base R.

Examples

```
a <- array(4 * runif(1500000), dim=c(10000, 30, 5))
A <- DelayedArray(a)
A

A2 <- dnorm(A + 1)[ , , -3] # very fast! (operations are delayed)
A2

a2 <- as.array(A2)          # "realize" 'A2' in memory (as an ordinary
                             # array)
```

```

DelayedArray(a2) == A2      # DelayedArray object of type "logical"
stopifnot(all(DelayedArray(a2) == A2))

library(HDF5Array)
A3 <- as(A2, "HDF5Array")   # "realize" 'A2' on disk (as an HDF5Array
                             # object)

A3 == A2                    # DelayedArray object of type "logical"
stopifnot(all(A3 == A2))

## See '?DelayedArray' for general information about DelayedArray objects
## and their "realization".

```

DelayedArray-utils *Common operations on DelayedArray objects*

Description

Common operations on [DelayedArray](#) objects.

Details

The operations currently supported on [DelayedArray](#) objects are:

Delayed operations:

- `rbind` and `cbind`
- all the members of the [Ops](#), [Math](#), and [Math2](#) groups
- `!`
- `is.na`, `is.finite`, `is.infinite`, `is.nan`
- `type<-`
- `lengths`
- `nchar`, `tolower`, `toupper`, `grepl`, `sub`, `gsub`
- `pmax2` and `pmin2`
- [sweep](#)
- [scale](#) (when the supplied center and scale are not TRUE)
- statistical functions like `dnorm`, `dbinom`, `dpois`, and `dlogis` (for the Normal, Binomial, Poisson, and Logistic distribution, respectively) and related functions (documented in [DelayedArray-stats](#))

Block-processed operations:

- `anyNA`, `which`
- `unique`, `table`
- all the members of the [Summary](#) group

- `mean`
- `apply`

Mix delayed and block-processed operations:

- `scale` (when the supplied center and/or scale are TRUE)

See Also

- `cbind` in the **base** package for `rbind/cbind`'ing ordinary arrays.
- `arbind` and `acbind` in this package (**DelayedArray**) for binding ordinary arrays of arbitrary dimensions along their rows or columns.
- `is.na`, `!`, `table`, `mean`, `apply`, and `%*%` in the **base** package for the corresponding operations on ordinary arrays or matrices.
- `DelayedMatrix-utils` for common operations on `DelayedMatrix` objects.
- `DelayedArray-stats` for statistical functions on `DelayedArray` objects.
- `DelayedMatrix-stats` for `DelayedMatrix` row/col summarization.
- `DelayedArray` objects.
- `HDF5Array` objects in the **HDF5Array** package.
- `S4groupGeneric` in the **methods** package for the members of the `Ops`, `Math`, and `Math2` groups.
- `sweep` and `scale` in the **base** package.

Examples

```
## -----
## BIND DelayedArray OBJECTS
## -----
## DelayedArray objects can be bound along their 1st (rows) or 2nd
## (columns) dimension with rbind() or cbind(). These operations are
## equivalent to arbind() and acbind(), respectively, and are all
## delayed.

## On 2D objects:
library(HDF5Array)
toy_h5 <- system.file("extdata", "toy.h5", package="HDF5Array")
h5ls(toy_h5)

M1 <- HDF5Array(toy_h5, "M1")
M2 <- HDF5Array(toy_h5, "M2")

M12 <- rbind(M1, t(M2))          # delayed
M12
colMeans(M12)                   # block-processed

## On objects with more than 2 dimensions:
example(arbind) # to create arrays a1, a2, a3
```

```

A1 <- DelayedArray(a1)
A2 <- DelayedArray(a2)
A3 <- DelayedArray(a3)
A123 <- rbind(A1, A2, A3)          # delayed
A123

## On 1D objects:
v1 <- array(11:15, 5, dimnames=list(LETTERS[1:5]))
v2 <- array(letters[1:3])
V1 <- DelayedArray(v1)
V2 <- DelayedArray(v2)
V12 <- rbind(V1, V2)
V12

## Not run: cbind(V1, V2) # Error! (the objects to cbind() must have at least 2
# dimensions)

## End(Not run)

## Note that base::rbind() and base::cbind() do something completely
## different on ordinary arrays that are not matrices. They treat them
## as if they were vectors:
rbind(a1, a2, a3)
cbind(a1, a2, a3)
rbind(v1, v2)
cbind(v1, v2)

## Also note that DelayedArray objects of arbitrary dimensions can be
## stored inside a DataFrame object as long as they all have the same
## first dimension (nrow()):
DF <- DataFrame(M=I(tail(M1, n=5)), A=I(A3), V=I(V1))
DF[-3, ]
DF2 <- rbind(DF, DF)
DF2$V

## Sanity checks:
m1 <- as.matrix(M1)
m2 <- as.matrix(M2)
stopifnot(identical(rbind(m1, t(m2)), as.matrix(M12)))
stopifnot(identical(arbind(a1, a2, a3), as.array(A123)))
stopifnot(identical(arbind(v1, v2), as.array(V12)))
stopifnot(identical(rbind(DF$M, DF$M), DF2$M))
stopifnot(identical(rbind(DF$A, DF$A), DF2$A))
stopifnot(identical(rbind(DF$V, DF$V), DF2$V))

## -----
## MORE OPERATIONS
## -----

M1 >= 0.5 & M1 < 0.75          # delayed
log(M1)                        # delayed
pmax2(M2, 0)                   # delayed

```

```

type(M2) <- "integer"          # delayed
M2

## table() is block-processed:
a4 <- array(sample(50L, 2000000L, replace=TRUE), c(200, 4, 2500))
A4 <- as(a4, "HDF5Array")
table(A4)
a5 <- array(sample(20L, 2000000L, replace=TRUE), c(200, 4, 2500))
A5 <- as(a5, "HDF5Array")
table(A5)

A4 - 2 * A5                    # delayed
table(A4 - 2 * A5)            # block-processed

## range() is block-processed:
range(A4 - 2 * A5)
range(M1)

cmeans <- colMeans(M2)        # block-processed
sweep(M2, 2, cmeans)          # delayed
scale(M2)                     # delayed & block-processed
scale(M2, center=FALSE, scale=10) # delayed

```

DelayedMatrix-stats *DelayedMatrix* row/col summarization

Description

Only a small number of row/col summarization methods are provided by the **DelayedArray** package.

See the **DelayedMatrixStats** package for an extensive set of row/col summarization methods.

Usage

```

## S4 method for signature 'DelayedMatrix'
rowSums(x, na.rm=FALSE, dims=1)
## S4 method for signature 'DelayedMatrix'
colSums(x, na.rm=FALSE, dims=1)

## S4 method for signature 'DelayedMatrix'
rowMeans(x, na.rm=FALSE, dims=1)
## S4 method for signature 'DelayedMatrix'
colMeans(x, na.rm=FALSE, dims=1)

## S4 method for signature 'DelayedMatrix'
rowMins(x, rows=NULL, cols=NULL, na.rm=FALSE)
## S4 method for signature 'DelayedMatrix'
colMins(x, rows=NULL, cols=NULL, na.rm=FALSE)

```

```
## S4 method for signature 'DelayedMatrix'
rowMaxs(x, rows=NULL, cols=NULL, na.rm=FALSE)
## S4 method for signature 'DelayedMatrix'
colMaxs(x, rows=NULL, cols=NULL, na.rm=FALSE)

## S4 method for signature 'DelayedMatrix'
rowRanges(x, rows=NULL, cols=NULL, na.rm=FALSE)
## S4 method for signature 'DelayedMatrix'
colRanges(x, rows=NULL, cols=NULL, na.rm=FALSE)
```

Arguments

<code>x</code>	A DelayedMatrix object.
<code>na.rm</code>	Should missing values (including NaN) be omitted from the calculations?
<code>dims, rows, cols</code>	These arguments are not supported. Don't use them.

Details

All these operations are block-processed.

See Also

- The **DelayedMatrixStats** package for more row/col summarization methods for [DelayedMatrix](#) objects.
- [rowSums](#) in the **base** package and [rowMaxs](#) in the **matrixStats** package for row/col summarization of an ordinary matrix.
- [DelayedMatrix-utils](#) for other common operations on [DelayedMatrix](#) objects.
- [DelayedMatrix](#) objects.
- [matrix](#) objects in base R.

Examples

```
library(HDF5Array)
toy_h5 <- system.file("extdata", "toy.h5", package="HDF5Array")
h5ls(toy_h5)

M1 <- HDF5Array(toy_h5, "M1")
M2 <- HDF5Array(toy_h5, "M2")

M12 <- rbind(M1, t(M2))      # delayed

## All these operations are block-processed.

rowSums(M12)
colSums(M12)

rowMeans(M12)
```



```

colMeans(M12)

rmins <- rowMins(M12)
cmins <- colMins(M12)

rmaxs <- rowMaxs(M12)
cmaxs <- colMaxs(M12)

rranges <- rowRanges(M12)
cranges <- colRanges(M12)

## Sanity checks:
m12 <- rbind(as.matrix(M1), t(as.matrix(M2)))
stopifnot(identical(rowSums(M12), rowSums(m12)))
stopifnot(identical(colSums(M12), colSums(m12)))
stopifnot(identical(rowMeans(M12), rowMeans(m12)))
stopifnot(identical(colMeans(M12), colMeans(m12)))
stopifnot(identical(rmins, rowMins(m12)))
stopifnot(identical(cmins, colMins(m12)))
stopifnot(identical(rmaxs, rowMaxs(m12)))
stopifnot(identical(cmaxs, colMaxs(m12)))
stopifnot(identical(rranges, cbind(rmins, rmaxs, deparse.level=0)))
stopifnot(identical(cranges, cbind(cmins, cmaxs, deparse.level=0)))

```

DelayedMatrix-utils *Common operations on DelayedMatrix objects*

Description

Common operations on [DelayedMatrix](#) objects.

Details

In addition to the operations supported on [DelayedArray](#) objects, [DelayedMatrix](#) objects support the following operations:

Delayed operations:

- `t`

Block-processed operations:

- `rowsum` and `colsum`
- matrix multiplication (`%*%`) of an ordinary matrix by a [DelayedMatrix](#) object
- matrix row/col summarization (see `?DelayedMatrix-stats`)

See Also

- `rowsum` in the **base** package for computing column sums across rows of an ordinary matrix for each level of a grouping variable.
- `DelayedArray-utils` for common operations on `DelayedArray` objects.
- `DelayedArray-stats` for statistical functions on `DelayedArray` objects.
- `DelayedMatrix-stats` for `DelayedMatrix` row/col summarization.
- `setAutoRealizationBackend` for how to set a *automatic realization backend*.
- `writeHDF5Array` in the **HDF5Array** package for writing an array-like object to an HDF5 file and other low-level utilities to control the location of automatically created HDF5 datasets.
- `DelayedArray` objects.
- `HDF5Array` objects in the **HDF5Array** package.
- `array` objects in base R.

Examples

```
## -----
## rowsum() / colsum()
## -----
library(HDF5Array)
set.seed(123)
m0 <- matrix(runif(14400000), ncol=2250,
              dimnames=list(NULL, sprintf("C%04d", 1:2250)))
M0 <- writeHDF5Array(m0, chunkdim=c(200, 250))
dimnames(M0) <- dimnames(m0)

## --- rowsum() ---

group <- sample(90, nrow(M0), replace=TRUE) # define groups of rows
rs <- rowsum(M0, group)
rs[1:5, 1:8]
rs2 <- rowsum(M0, group, reorder=FALSE)
rs2[1:5, 1:8]

## Let's see block processing in action:
DelayedArray::set_verbose_block_processing(TRUE)
setAutoBlockSize(2e6)
rs3 <- rowsum(M0, group)
setAutoBlockSize()
DelayedArray::set_verbose_block_processing(FALSE)

## Sanity checks:
stopifnot(all.equal(rowsum(m0, group), rs))
stopifnot(all.equal(rowsum(m0, group, reorder=FALSE), rs2))
stopifnot(all.equal(rs, rs3))

## --- colsum() ---

group <- sample(30, ncol(M0), replace=TRUE) # define groups of cols
cs <- colsum(M0, group)
```

```

cs[1:5, 1:7]
cs2 <- colsum(M0, group, reorder=FALSE)
cs2[1:5, 1:7]

## Sanity checks:
stopifnot(all.equal(colsum(m0, group), cs))
stopifnot(all.equal(cs, t(rowsum(t(m0), group))))
stopifnot(all.equal(cs, t(rowsum(t(M0), group))))
stopifnot(all.equal(colsum(m0, group, reorder=FALSE), cs2))
stopifnot(all.equal(cs2, t(rowsum(t(m0), group, reorder=FALSE))))
stopifnot(all.equal(cs2, t(rowsum(t(M0), group, reorder=FALSE))))

## -----
## MATRIX MULTIPLICATION
## -----
library(HDF5Array)
toy_h5 <- system.file("extdata", "toy.h5", package="HDF5Array")
h5ls(toy_h5)
M1 <- HDF5Array(toy_h5, "M1")

## Matrix multiplication is not delayed: the output matrix is realized
## block by block. The current "automatic realization backend" controls
## where realization happens e.g. in memory as an ordinary matrix if not
## (i.e. set to NULL) or in an HDF5 file if set to "HDF5Array".
## See '?setAutoRealizationBackend' for more information about
## "realization backends".
## The output matrix is returned as a DelayedMatrix object with no delayed
## operations on it. The exact class of the object depends on the backend
## e.g. it will be HDF5Matrix with "HDF5Array" backend.

m <- matrix(runif(50000), ncol=nrow(M1))

## Set backend to NULL for in-memory realization:
setAutoRealizationBackend()
P1 <- m %*% M1
P1

## Set backend to HDF5Array for realization in HDF5 file:
setAutoRealizationBackend("HDF5Array")

## With the HDF5Array backend, the output matrix will be written to an
## automatic location on disk:
getHDF5DumpFile() # HDF5 file where the output matrix will be written
lsHDF5DumpFile()

P2 <- m %*% M1
P2

lsHDF5DumpFile()

## Use setHDF5DumpFile() and setHDF5DumpName() from the HDF5Array package
## to control the location of automatically created HDF5 datasets.

```

DelayedNaryIsoOp-class
DelayedNaryIsoOp objects

NOTE: This man page is about [DelayedArray](#) internals and is provided for developers and advanced users only.

```

DelayedOp
  ^
  |
DelayedNaryOp
  ^
  |
DelayedNaryIsoOp

```

Usage

[illegible]

```
is_sparse(x)

## S4 method for signature 'DelayedNaryIsoOp'
OLD_extract_sparse_array(x, index)
```

Arguments

x, object	A DelayedNaryIsoOp object.
index	See ?extract_array in the S4Arrays package for a description of the index argument.
...	Not used.

See Also

- [DelayedOp](#) objects.
- [showtree](#) to visualize the nodes and access the leaves in the tree of delayed operations carried by a [DelayedArray](#) object.
- [extract_array](#) in the **S4Arrays** package.
- [OLD_extract_sparse_array](#).

Examples

```
## DelayedNaryIsoOp extends DelayedNaryOp which extends DelayedOp:
extends("DelayedNaryIsoOp")

## -----
## BASIC EXAMPLE
## -----
m1 <- matrix(101:130, ncol=5)
m2 <- matrix(runif(30), ncol=5)
M1 <- DelayedArray(m1)
M2 <- DelayedArray(m2)
showtree(M1)
showtree(M2)

M <- M1 / M2
showtree(M)
class(M@seed)      # a DelayedNaryIsoOp object

## -----
## PROPAGATION OF SPARSITY
## -----
sm1 <- sparseMatrix(i=c(1, 6), j=c(1, 4), x=c(11, 64), dims=6:5)
SM1 <- DelayedArray(sm1)
sm2 <- sparseMatrix(i=c(2, 6), j=c(1, 5), x=c(21, 65), dims=6:5)
SM2 <- DelayedArray(sm2)
showtree(SM1)
showtree(SM2)
is_sparse(SM1)      # TRUE
is_sparse(SM2)      # TRUE
```

```

SM3 <- SM1 - SM2
showtree(SM3)
class(SM3@seed)      # a DelayedNaryIsoOp object
is_sparse(SM3@seed)   # TRUE

M4 <- SM1 / SM2
showtree(M4)
class(M4@seed)        # a DelayedNaryIsoOp object
is_sparse(M4@seed)    # FALSE

## -----
## SANITY CHECKS
## -----
stopifnot(class(M@seed) == "DelayedNaryIsoOp")
stopifnot(class(SM3@seed) == "DelayedNaryIsoOp")
stopifnot(is_sparse(SM3@seed))
stopifnot(class(M4@seed) == "DelayedNaryIsoOp")
stopifnot(!is_sparse(M4@seed))

```

DelayedOp-class

DelayedOp objects

Description

NOTE: This man page is about [DelayedArray](#) internals and is provided for developers and advanced users only.

In a [DelayedArray](#) object, the delayed operations are stored as a tree where the leaves are operands and the nodes are the operations. Each node in the tree is a DelayedOp derivative representing a particular delayed operation.

DelayedOp is a virtual class with 8 concrete subclasses. Each subclass provides a formal representation for a particular kind of delayed operation.

Usage

```
is_noop(x)
```

Arguments

x A DelayedSubset, DelayedAperm, or DelayedSetDimnames object.

Details

8 types of nodes are currently supported. Each type is a DelayedOp subclass:

Node type	Represented operation

DelayedOp (VIRTUAL)	

* DelayedUnaryOp (VIRTUAL)		
o DelayedSubset		Multi-dimensional single bracket subsetting.
o DelayedAperm		Extended aperm() (can drop and/or add ineffective dimensions).
o DelayedUnaryIsoOp (VIRTUAL)		Unary op that preserves the geometry.
- DelayedUnaryIsoOpStack		Simple ops stacked together.
- DelayedUnaryIsoOpWithArgs		One op with vector-like arguments along the dimensions of the input.
- DelayedSubassign		Multi-dimensional single bracket subassignment.
- DelayedSetDimnames		Set/replace the dimnames.

* DelayedNaryOp (VIRTUAL)		
o DelayedNaryIsoOp		N-ary op that preserves the geometry.
o DelayedAbind		abind()

All the nodes are array-like objects that must comply with the *seed contract* i.e. they must support `dim()`, `dimnames()`, and `extract_array()`. See [?extract_array](#) in the **S4Arrays** package for more information about the *seed contract*.

`is_noop()` can only be called on a `DelayedSubset`, `DelayedAperm`, or `DelayedSetDimnames` object at the moment, and will return `TRUE` if the object represents a no-op.

Note

The `DelayedOp` virtual class and its 8 concrete subclasses are used inside a `DelayedArray` object to represent delayed operations carried by the object. They're never exposed to the end user and are not intended to be manipulated directly.

See Also

- `DelayedOp` concrete subclasses: [DelayedSubset](#), [DelayedAperm](#), [DelayedUnaryIsoOpStack](#), [DelayedUnaryIsoOpWithArgs](#), [DelayedSubassign](#), [DelayedSetDimnames](#), [DelayedNaryIsoOp](#), and [DelayedAbind](#).
- [DelayedArray](#) objects.
- [showtree](#) to visualize the nodes and access the leaves in the tree of delayed operations carried by a [DelayedArray](#) object.
- [simplify](#) to simplify the tree of delayed operations carried by a [DelayedArray](#) object.
- [extract_array](#) in the **S4Arrays** package.


```
## DelayedSetDimnames objects inherit the default
## is_sparse() and OLD_extract_sparse_array() methods defined
## for DelayedUnaryIsoOp derivatives.
```

Arguments

```
x, object      A DelayedSetDimnames object.
...           Not used.
```

See Also

- [DelayedOp](#) objects.
- [showtree](#) to visualize the nodes and access the leaves in the tree of delayed operations carried by a [DelayedArray](#) object.

Examples

```
## DelayedSetDimnames extends DelayedUnaryIsoOp, which extends
## DelayedUnaryOp, which extends DelayedOp:
extends("DelayedSetDimnames")

## -----
## BASIC EXAMPLE
## -----
m0 <- matrix(1:30, ncol=5, dimnames=list(letters[1:6], NULL))
M2 <- M1 <- M0 <- DelayedArray(m0)
showtree(M0)

dimnames(M1) <- list(NULL, LETTERS[1:5])
showtree(M1)
class(M1@seed)      # a DelayedSetDimnames object

colnames(M2) <- LETTERS[1:5]
showtree(M2)
class(M2@seed)      # a DelayedSetDimnames object

## -----
## PROPAGATION OF SPARSITY
## -----
## DelayedSetDimnames objects always propagate sparsity.

sm0 <- sparseMatrix(i=c(1, 4), j=c(1, 3), x=c(11, 43), dims=4:3)
SM <- SM0 <- DelayedArray(sm0)
showtree(SM0)
is_sparse(SM0)      # TRUE

dimnames(SM) <- list(letters[1:4], LETTERS[1:3])
showtree(SM)
class(SM@seed)      # a DelayedSetDimnames object
is_sparse(SM@seed)  # TRUE
```



```

showtree(M2)
class(M2@seed)      # a DelayedSubassign object

## -----
## PROPAGATION OF SPARSITY
## -----

## DelayedSubassign objects don't propagate sparsity at the moment, that
## is, is_sparse() always returns FALSE on them.

## -----
## SANITY CHECKS
## -----
stopifnot(class(M1@seed) == "DelayedSubassign")
stopifnot(class(M2@seed) == "DelayedSubassign")

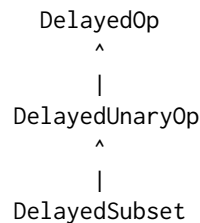
```

DelayedSubset-class *DelayedSubset objects*

Description

NOTE: This man page is about [DelayedArray](#) internals and is provided for developers and advanced users only.

The DelayedSubset class provides a formal representation of a *delayed multi-dimensional single bracket subsetting operation*. It is a concrete subclass of the [DelayedUnaryOp](#) virtual class, which itself is a subclass of the [DelayedOp](#) virtual class:



DelayedSubset objects are used inside a [DelayedArray](#) object to represent the *delayed multi-dimensional single bracket subsetting operations* carried by the object. They're never exposed to the end user and are not intended to be manipulated directly.

Usage

```

## S4 method for signature 'DelayedSubset'
is_noop(x)

## S4 method for signature 'DelayedSubset'
summary(object, ...)

```



```
## -----
## PROPAGATION OF SPARSITY
## -----
sm0 <- sparseMatrix(i=c(1, 4), j=c(1, 3), x=c(11, 43), dims=4:3)
SM0 <- DelayedArray(sm0)
showtree(SM0)
is_sparse(SM0)      # TRUE

SM1 <- SM0[-1, 3:2, drop=FALSE]
showtree(SM1)
class(SM1@seed)      # a DelayedSubset object
is_sparse(SM1@seed)  # TRUE

## Duplicated indices break structural sparsity.
M2 <- SM0[-1, c(3:2, 2), drop=FALSE]
showtree(M2)
class(M2@seed)      # a DelayedSubset object
is_sparse(M2@seed)  # FALSE

## -----
## SANITY CHECKS
## -----
stopifnot(class(A@seed) == "DelayedSubset")
stopifnot(class(SM1@seed) == "DelayedSubset")
stopifnot(is_sparse(SM1@seed))
stopifnot(class(M2@seed) == "DelayedSubset")
stopifnot(!is_sparse(M2@seed))
```

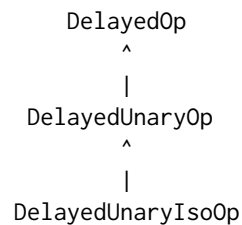
DelayedUnaryIsoOpStack-class

DelayedUnaryIsoOpStack objects

Description

NOTE: This man page is about [DelayedArray](#) internals and is provided for developers and advanced users only.

The DelayedUnaryIsoOpStack class provides a formal representation of a *stack of delayed unary isometric operations*, that is, of a group of delayed unary isometric operations stacked (a.k.a. piped) together. It is a concrete subclass of the [DelayedUnaryIsoOp](#) virtual class, which itself is a subclass of the [DelayedUnaryOp](#) virtual class, which itself is a subclass of the [DelayedOp](#) virtual class:



- `a` is the input array.
- The output (out) is an array of same dimensions as `a`.

In addition, each operation (OP) in the pipe must satisfy the property that each value in the output array must be determined ***solely*** by the corresponding value in the input array. In other words:

```
a |> OP |> `[`(i_1, i_2, ..., i_n) # i.e. OP(a)[i_1, i_2, ..., i_n]
```

must be equal to:

```
a |> `[`(i_1, i_2, ..., i_n) |> OP # i.e. OP(a[i_1, i_2, ..., i_n])
```

for any valid multidimensional index (`i_1, i_2, ..., i_n`).

We refer to this property as the *locality principle*.

Concrete examples:

1. Things like `is.na()`, `is.finite()`, logical negation (`!`), `nchar()`, `tolower()`.
2. Most functions in the [Math](#) and [Math2](#) groups e.g. `log()`, `sqrt()`, `abs()`, `ceiling()`, `round()`, etc... Notable exceptions are the `cum*()` functions (`cummin()`, `cummax()`, `cumsum()`, and `cumprod()`): they don't satisfy the *locality principle*.
3. Operations in the [Ops](#) group when one operand is an array and the other a scalar e.g. `a + 10`, `2 ^ a`, `a <= 0.5`, etc...

See Also

- [DelayedOp](#) objects.
- [showtree](#) to visualize the nodes and access the leaves in the tree of delayed operations carried by a [DelayedArray](#) object.
- [extract_array](#) in the [S4Arrays](#) package.
- [OLD_extract_sparse_array](#).

Examples

```
## DelayedUnaryIsoOpStack extends DelayedUnaryIsoOp, which extends
## DelayedUnaryOp, which extends DelayedOp:
extends("DelayedUnaryIsoOpStack")

## -----
## BASIC EXAMPLE
## -----
m0 <- matrix(runif(12), ncol=3)
M0 <- DelayedArray(m0)
showtree(M0)

M <- log(1 + M0) / 10
showtree(M)
class(M@seed)      # a DelayedUnaryIsoOpStack object
```



```

## -----
## PROPAGATION OF SPARSITY
## -----
sm0 <- sparseMatrix(i=c(1, 4), j=c(1, 3), x=c(11, 43), dims=4:3)
SM0 <- DelayedArray(sm0)
showtree(SM0)
is_sparse(SM0)      # TRUE

M1 <- SM0 - 11
showtree(M1)
class(M1@seed)      # a DelayedUnaryIsoOpStack object
is_sparse(M1@seed)  # FALSE

SM2 <- 10 * SM0
showtree(SM2)
class(SM2@seed)     # a DelayedUnaryIsoOpStack object
is_sparse(SM2@seed) # TRUE

M3 <- SM0 / 0
showtree(M3)
class(M3@seed)      # a DelayedUnaryIsoOpStack object
is_sparse(M3@seed)  # FALSE

SM4 <- log(1 + SM0) / 10
showtree(SM4)
class(SM4@seed)     # a DelayedUnaryIsoOpStack object
is_sparse(SM4@seed) # TRUE

SM5 <- 2 ^ SM0 - 1
showtree(SM5)
class(SM5@seed)     # a DelayedUnaryIsoOpStack object
is_sparse(SM5@seed) # TRUE

## -----
## SANITY CHECKS
## -----
stopifnot(class(M@seed) == "DelayedUnaryIsoOpStack")
stopifnot(class(M1@seed) == "DelayedUnaryIsoOpStack")
stopifnot(!is_sparse(M1@seed))
stopifnot(class(SM2@seed) == "DelayedUnaryIsoOpStack")
stopifnot(is_sparse(SM2@seed))
stopifnot(class(M3@seed) == "DelayedUnaryIsoOpStack")
stopifnot(!is_sparse(M3@seed))
stopifnot(class(SM4@seed) == "DelayedUnaryIsoOpStack")
stopifnot(is_sparse(SM4@seed))
stopifnot(class(SM5@seed) == "DelayedUnaryIsoOpStack")
stopifnot(is_sparse(SM5@seed))

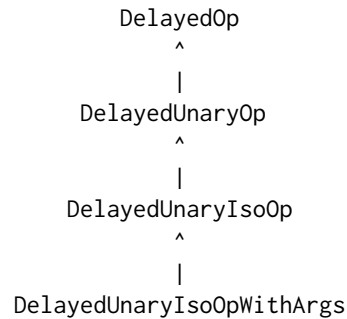
```

DelayedUnaryIsoOpWithArgs-class

*DelayedUnaryIsoOpWithArgs objects***Description**

NOTE: This man page is about [DelayedArray](#) internals and is provided for developers and advanced users only.

The DelayedUnaryIsoOpWithArgs class provides a formal representation of a *delayed unary isometric operation with vector-like arguments going along the dimensions of the input array*. It is a concrete subclass of the [DelayedUnaryIsoOp](#) virtual class, which itself is a subclass of the [DelayedUnaryOp](#) virtual class, which itself is a subclass of the [DelayedOp](#) virtual class:



DelayedUnaryIsoOpWithArgs objects are used inside a [DelayedArray](#) object to represent the *delayed unary isometric operations with vector-like arguments going along the dimensions of the input array* carried by the object. They're never exposed to the end user and are not intended to be manipulated directly.

Usage

```

## S4 method for signature 'DelayedUnaryIsoOpWithArgs'
summary(object, ...)

## ~ ~ ~ Seed contract ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~

## DelayedUnaryIsoOpWithArgs objects inherit the default dim()
## and dimnames() methods defined for DelayedUnaryIsoOp
## derivatives, but overwrite their extract_array() method.

## S4 method for signature 'DelayedUnaryIsoOpWithArgs'
extract_array(x, index)

## ~ ~ ~ Propagation of sparsity ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~

## S4 method for signature 'DelayedUnaryIsoOpWithArgs'
is_sparse(x)
  
```

```
## S4 method for signature 'DelayedUnaryIsoOpWithArgs'
OLD_extract_sparse_array(x, index)
```

Arguments

x, object	A DelayedUnaryIsoOpWithArgs object.
index	See ?extract_array in the S4Arrays package for a description of the index argument.
...	Not used.

Details

A DelayedUnaryIsoOpWithArgs object is used to represent the delayed version of an operation of the form:

```
out <- OP(L1, L2, ..., a, R1, R2, ...)
```

where:

- OP is an isometric array transformation i.e. a transformation that returns an array with the same dimensions as the input array.
- a is the input array.
- L1, L2, etc... are the left arguments.
- R1, R2, etc... are the right arguments.
- The output (out) is an array of same dimensions as a.

Some of the arguments (left or right) can go along the dimensions of the input array. For example if a is a 12 x 150 x 5 array, argument L2 is considered to go along the 3rd dimension if its length is 5 and if the result of:

```
OP(L1, L2[k], ..., a[, , k, drop=FALSE], R1, R2, ...)
```

is the same as `out[, , k, drop=FALSE]` for any index k.

More generally speaking, if, say, arguments L2, L3, R1, and R2 go along the 3rd, 1st, 2nd, and 1st dimensions, respectively, then each value in the output array (`a[i, j, k]`) must be determined solely by the corresponding values in the input array (`a[i, j, k]`) and arguments (`L2[k]`, `L3[i]`, `R1[j]`, `R2[i]`). In other words, `out[i, j, k]` must be equal to:

```
OP(L1, L2[k], L3[i], ..., a[i, j, k], R1[j], R2[i], ...)
```

for any $1 \leq i \leq 12$, $1 \leq j \leq 150$, and $1 \leq k \leq 5$.

We refer to this property as the *locality principle*.

Concrete examples:

1. Addition (or any operation in the [Ops](#) group) of an array `a` and an atomic vector `v` of length `dim(a)[[1]]`:
 - ``+`(a, v)`: OP is ``+``, right argument goes along the 1st dimension.
 - ``<=`(a, v)`: OP is ``<=``, right argument goes along the 1st dimension.
 - ``&`(v, a)`: OP is ``&``, left argument goes along the 1st dimension.
2. `scale(x, center=v1, scale=v2)`: OP is `scale`, right arguments center and scale go along the 2nd dimension.

Note that if OP has no argument that goes along a dimension of the input array, then the delayed operation is better represented with a [DelayedUnaryIsoOpStack](#) object.

See Also

- [DelayedOp](#) objects.
- [showtree](#) to visualize the nodes and access the leaves in the tree of delayed operations carried by a [DelayedArray](#) object.
- [extract_array](#) in the **S4Arrays** package.
- [OLD_extract_sparse_array](#).

Examples

```
## DelayedUnaryIsoOpWithArgs extends DelayedUnaryIsoOp, which extends
## DelayedUnaryOp, which extends DelayedOp:
extends("DelayedUnaryIsoOpWithArgs")

## -----
## BASIC EXAMPLE
## -----
m0 <- matrix(runif(12), ncol=3)
M0 <- DelayedArray(m0)
showtree(M0)

M <- M0 + 101:104
showtree(M)
class(M@seed)      # a DelayedUnaryIsoOpWithArgs object

## -----
## PROPAGATION OF SPARSITY
## -----
sm0 <- sparseMatrix(i=c(1, 4), j=c(1, 3), x=c(11, 43), dims=4:3)
SM0 <- DelayedArray(sm0)
showtree(SM0)
is_sparse(SM0)      # TRUE

M1 <- SM0 + 101:104
showtree(M1)
class(M1@seed)      # a DelayedUnaryIsoOpWithArgs object
is_sparse(M1@seed)   # FALSE

SM2 <- SM0 * 101:104
```

```

showtree(SM2)
class(SM2@seed)      # a DelayedUnaryIsoOpWithArgs object
is_sparse(SM2@seed)  # TRUE

SM3 <- SM0 * c(101:103, 0)
showtree(SM3)
class(SM3@seed)      # a DelayedUnaryIsoOpWithArgs object
is_sparse(SM3@seed)  # TRUE

M4 <- SM0 * c(101:103, NA)
showtree(M4)
class(M4@seed)       # a DelayedUnaryIsoOpWithArgs object
is_sparse(M4@seed)   # FALSE

M5 <- SM0 * c(101:103, Inf)
showtree(M5)
class(M5@seed)       # a DelayedUnaryIsoOpWithArgs object
is_sparse(M5@seed)   # FALSE

SM6 <- SM0 / 101:104
showtree(SM6)
class(SM6@seed)      # a DelayedUnaryIsoOpWithArgs object
is_sparse(SM6@seed)  # TRUE

M7 <- SM0 / c(101:103, 0)
showtree(M7)
class(M7@seed)       # a DelayedUnaryIsoOpWithArgs object
is_sparse(M7@seed)   # FALSE

M8 <- SM0 / c(101:103, NA)
showtree(M8)
class(M8@seed)       # a DelayedUnaryIsoOpWithArgs object
is_sparse(M8@seed)   # FALSE

SM9 <- SM0 / c(101:103, Inf)
showtree(SM9)
class(SM9@seed)      # a DelayedUnaryIsoOpWithArgs object
is_sparse(SM9@seed)  # TRUE

M10 <- 101:104 / SM0
showtree(M10)
class(M10@seed)      # a DelayedUnaryIsoOpWithArgs object
is_sparse(M10@seed)  # FALSE

## -----
## ADVANCED EXAMPLE
## -----
## Not ready yet!
#op <- DelayedArray::new_DelayedUnaryIsoOpWithArgs(m0,
#          scale,
#          Rargs=list(center=c(1, 0, 100), scale=c(10, 1, 1)),
#          Ralong=c(2, 2))

```


Arguments

maxvol	The maximum volume of the box to return.
maxdim	The dimensions of the constraining box.
shape	The shape of the box to return.
refdim	The dimensions of the reference array of the grid to return.
viewport_len	The maximum length of the elements (a.k.a. viewports) of the grid to return.
viewport_shape	The shape of the elements (a.k.a. viewports) of the grid to return.

Details

makeCappedVolumeBox returns the dimensions of a box that satisfies the following constraints:

1. The volume of the box is as close as possible to (but no bigger than) maxvol.
2. The box fits in the *constraining box* i.e. in the box whose dimensions are specified via maxdim.
3. The box has a non-zero volume if the *constraining box* has a non-zero volume.
4. The shape of the box is as close as possible to the requested shape.

The supported shapes are:

- hypercube: The box should be as close as possible to an *hypercube* (a.k.a. *n-cube*), that is, the ratio between its biggest and smallest dimensions should be as close as possible to 1.
- scale: The box should have the same proportions as the *constraining box*.
- first-dim-grows-first: The box will be grown along its 1st dimension first, then along its 2nd dimension, etc...
- last-dim-grows-first: Like first-dim-grows-first but starting along the last dimension.

See Also

- [defaultAutoGrid](#) and family to create automatic grids to use for block processing of array-like objects.
- [ArrayGrid](#) in the **S4Arrays** package for the formal representation of grids and viewports.

Examples

```
## -----
## makeCappedVolumeBox()
## -----

maxdim <- c(50, 12) # dimensions of the "constraining box"

## "hypercube" shape:
makeCappedVolumeBox(40, maxdim)
makeCappedVolumeBox(120, maxdim)
makeCappedVolumeBox(125, maxdim)
makeCappedVolumeBox(200, maxdim)
```

```

## "scale" shape:
makeCappedVolumeBox(40, maxdim, shape="scale")
makeCappedVolumeBox(160, maxdim, shape="scale")

## "first-dim-grows-first" and "last-dim-grows-first" shapes:
makeCappedVolumeBox(120, maxdim, shape="first-dim-grows-first")
makeCappedVolumeBox(149, maxdim, shape="first-dim-grows-first")
makeCappedVolumeBox(150, maxdim, shape="first-dim-grows-first")

makeCappedVolumeBox(40, maxdim, shape="last-dim-grows-first")
makeCappedVolumeBox(59, maxdim, shape="last-dim-grows-first")
makeCappedVolumeBox(60, maxdim, shape="last-dim-grows-first")

## -----
## makeRegularArrayGridOfCappedLengthViewports()
## -----

grid1a <- makeRegularArrayGridOfCappedLengthViewports(maxdim, 40)
grid1a
as.list(grid1a) # turn the grid into a list of ArrayViewport objects
table(lengths(grid1a))
stopifnot(maxlength(grid1a) <= 40) # sanity check

grid1b <- makeRegularArrayGridOfCappedLengthViewports(maxdim, 40,
                                                       "first-dim-grows-first")
grid1b
as.list(grid1b) # turn the grid into a list of ArrayViewport objects
table(lengths(grid1b))
stopifnot(maxlength(grid1b) <= 40) # sanity check

grid2a <- makeRegularArrayGridOfCappedLengthViewports(maxdim, 120)
grid2a
as.list(grid2a) # turn the grid into a list of ArrayViewport objects
table(lengths(grid2a))
stopifnot(maxlength(grid2a) <= 120) # sanity check

grid2b <- makeRegularArrayGridOfCappedLengthViewports(maxdim, 120,
                                                       "first-dim-grows-first")
grid2b
as.list(grid2b) # turn the grid into a list of ArrayViewport objects
table(lengths(grid2b))
stopifnot(maxlength(grid2b) <= 120) # sanity check

grid3a <- makeRegularArrayGridOfCappedLengthViewports(maxdim, 200)
grid3a
as.list(grid3a) # turn the grid into a list of ArrayViewport objects
table(lengths(grid3a))
stopifnot(maxlength(grid3a) <= 200) # sanity check

grid3b <- makeRegularArrayGridOfCappedLengthViewports(maxdim, 200,
                                                       "first-dim-grows-first")
grid3b
as.list(grid3b) # turn the grid into a list of ArrayViewport objects

```



```
table(lengths(grid3b))
stopifnot(maxlength(grid3b) <= 200) # sanity check
```

read_sparse_block	<i>read_sparse_block()</i>
-------------------	----------------------------

Description

An internal generic not meant to be called directly by the end user.

RealizationSink	<i>RealizationSink objects</i>
-----------------	--------------------------------

Description

Use a RealizationSink object in combination with [write_block\(\)](#) to write blocks of array data to disk.

RealizationSink is a virtual class with various concrete subclasses supporting writing data to specific formats.

sinkApply() is a convenience function for walking on a RealizationSink object, typically for the purpose of filling it with blocks of data.

Note that [write_block\(\)](#) is typically used inside the callback function passed to sinkApply().

Usage

```
## Walk on a RealizationSink derivative:
sinkApply(sink, FUN, ..., grid=NULL, verbose=NA)

## Backend-agnostic RealizationSink constructor:
AutoRealizationSink(dim, dimnames=NULL, type="double", as.sparse=FALSE)

## Get/set the "automatic realization backend":
getAutoRealizationBackend()
setAutoRealizationBackend(BACKEND=NULL)
supportedRealizationBackends()
```

Arguments

sink	A **writable** array-like object, typically a RealizationSink derivative. Some important notes: • DelayedArray objects are NEVER writable, even when they don't carry delayed operations (e.g. HDF5Array objects from the HDF5Array package), and even when they don't carry delayed operations **and** have all their data in memory (e.g. RleArray objects). In other words, there are NO exceptions.
------	---

- RealizationSink is a **virtual** class so sink will always be a RealizationSink **derivative**, that is, an object that belongs to a **concrete** subclass of the RealizationSink class (e.g. an [HDF5RealizationSink](#) object from the **HDF5Array** package).
- RealizationSink derivatives are considered array-like objects i.e. they have dimensions and possibly dimnames.

Although [write_block\(\)](#) and [sinkApply\(\)](#) will typically be used on a RealizationSink derivative, they can also be used on an ordinary array or other writable in-memory array-like objects like **dgCMatrix** objects from the **Matrix** package.

FUN	The callback function to apply to each viewport of the grid used to walk on sink. sinkApply() will perform <code>sink <- FUN(sink, viewport, ...)</code> on each viewport, so FUN must take at least two arguments, typically sink and viewport (but the exact names can differ).
	The function is expected to return its 1st argument (sink) possibly modified (e.g. when FUN contains a call to write_block() , which is typically the case).
...	Additional arguments passed to FUN.
grid	The grid used for the walk, that is, an ArrayGrid object that defines the viewports to walk on. It must be compatible with the geometry of sink. If not specified, an automatic grid is created by calling defaultSinkAutoGrid(sink) , and used. See ?defaultSinkAutoGrid for more information.
verbose	Whether block processing progress should be displayed or not. If set to NA (the default), verbosity is controlled by <code>DelayedArray::get_verbose_block_processing()</code> . Setting verbose to TRUE or FALSE overrides this.
dim	The dimensions (specified as an integer vector) of the RealizationSink derivative to create.
dimnames	The dimnames (specified as a list of character vectors or NULLs) of the RealizationSink derivative to create.
type	The type of the data that will be written to the RealizationSink derivative to create.
as.sparse	Whether the data should be written as sparse or not to the RealizationSink derivative to create. Not all <i>realization backends</i> support this.
BACKEND	NULL (the default), or a single string specifying the name of a realization backend e.g. "HDF5Array" or "RleArray" etc...

Details

***** The RealizationSink API *****

The DelayedArray package provides a simple API for writing blocks of array data to disk (or to memory): the "RealizationSink API". This API allows the developer to write code that is agnostic about the particular on-disk (or in-memory) format being used to store the data.

Here is how to use it:

1. Create a realization sink.
2. Write blocks of array data to the realization sink with one or several calls to [write_block\(\)](#).
3. Close the realization sink with [close\(\)](#).

4. Coerce the realization sink to `DelayedArray`.

A realization sink is formally represented by a `RealizationSink` derivative. Note that `RealizationSink` is a virtual class with various concrete subclasses like `HDF5RealizationSink` from the **HDF5Array** package, or `RleRealizationSink`. Each subclass implements the "RealizationSink API" for a specific realization backend.

To create a realization sink, use the specific constructor function. This function should be named as the class itself e.g. `HDF5RealizationSink()`.

To create a realization sink in a backend-agnostic way, use `AutoRealizationSink()`. It will create a `RealizationSink` derivative for the current *automatic realization backend* (see below).

Once writing to the realization sink is completed, the `RealizationSink` derivative must be closed (with `close(sink)`), then coerced to `DelayedArray` (with `as(sink, "DelayedArray")`). What specific `DelayedArray` derivative this coercion will return depends on the specific class of the `RealizationSink` derivative. For example, if `sink` is an `HDF5RealizationSink` object from the **HDF5Array** package, then `as(sink, "DelayedArray")` will return an `HDF5Array` instance (the `HDF5Array` class is a `DelayedArray` subclass).

*** The *automatic realization backend* ***

The *automatic realization backend* is a user-controlled global setting that indicates what specific `RealizationSink` derivative `AutoRealizationSink()` should return. In the context of block processing of a `DelayedArray` object, this controls where/how realization happens e.g. as an ordinary array if not set (i.e. set to `NULL`), or as an `HDF5Array` object if set to `"HDF5Array"`, or as an `RleArray` object if set to `"RleArray"`, etc...

Use `getAutoRealizationBackend()` or `setAutoRealizationBackend()` to get or set the *automatic realization backend*.

Use `supportedRealizationBackends()` to get the list of realization backends that are currently supported.

*** Cross realization backend compatibility ***

Two important things to keep in mind for developers aiming at writing code that is compatible across realization backends:

- Realization backends don't necessarily support concurrent writing.
More precisely: Even though it is safe to assume that any `DelayedArray` object will support concurrent `read_block()` calls, it is not so safe to assume that any `RealizationSink` derivative will support concurrent calls to `write_block()`. For example, at the moment, `HDF5RealizationSink` objects do not support concurrent writing.
This means that in order to remain compatible across realization backends, code that contains calls to `write_block()` should NOT be parallelized.
- Some realization backends are "linear write only", that is, they don't support *random write access*, only *linear write access*.
Such backends will provide a realization sink where the blocks of data must be written in linear order (i.e. by ascending rank). Furthermore, the geometry of the blocks must also be compatible with *linear write access*, that is, they must have a "first-dim-grows-first" shape. Concretely this means that the grid used to walk on the realization sink must be created with something like:

```
colAutoGrid(sink)
```

for a two-dimensional sink, or with something like:

```
defaultSinkAutoGrid(sink)
```

for a sink with an arbitrary number of dimensions.

See [?defaultSinkAutoGrid](#) for more information.

For obvious reasons, "linear write only" realization backends do not support concurrent writing.

Value

For `sinkApply()`, its 1st argument (`sink`) possibly modified (e.g. when callback function `FUN` contains a call to [write_block\(\)](#), which is typically the case).

For `AutoRealizationSink()`, a `RealizationSink` derivative with the class associated with the current *automatic realization backend*.

For `getAutoRealizationBackend`, `NULL` (no backend set yet) or a single string specifying the name of the *automatic realization backend* currently in use.

For `supportedRealizationBackends`, a data frame with 1 row per supported realization backend.

See Also

- [read_block](#) and [write_block](#) in the **S4Arrays** package.
- [ArrayGrid](#) in the **S4Arrays** package for the formal representation of grids and viewports.
- [defaultSinkAutoGrid](#) to create an automatic grid on a `RealizationSink` derivative.
- [SparseArraySeed](#) objects.
- [blockApply](#) and family for convenient block processing of an array-like object.
- [HDF5RealizationSink](#) objects in the **HDF5Array** package.
- [HDF5-dump-management](#) in the **HDF5Array** package to control the location and physical properties of automatically created HDF5 datasets.
- [RleArray](#) objects.
- [DelayedArray](#) objects.
- [array](#) objects in base R.

Examples

```
## -----
## USING THE "RealizationSink API": EXAMPLE 1
## -----

## -- STEP 1 --
## Create a realization sink. Note that instead of creating a
## realization sink by calling a backend-specific sink constructor
## (e.g. HDF5Array::HDF5RealizationSink), we set the "automatic
## realization backend" to "HDF5Array" and use backend-agnostic
## constructor AutoRealizationSink():
setAutoRealizationBackend("HDF5Array")
sink <- AutoRealizationSink(c(35L, 50L, 8L))
```

```

dim(sink)

## -- STEP 2 --
## Define the grid of viewports to walk on. Here we define a grid made
## of very small viewports on 'sink'. Note that, for real-world use cases,
## block processing will typically use grids made of much bigger
## viewports, usually obtained with defaultSinkAutoGrid().
## Also please note that this grid would not be compatible with "linear
## write only" realization backends. See "Cross realization backend
## compatibility" above in this man page for more information.
sink_grid <- RegularArrayGrid(dim(sink), spacings=c(20, 20, 4))

## -- STEP 3 --
## Walk on the grid, and, for each viewport, write random data to it.
for (bid in seq_along(sink_grid)) {
  viewport <- sink_grid[[bid]]
  block <- array(runif(length(viewport)), dim=dim(viewport))
  sink <- write_block(sink, viewport, block)
}

## -- An alternative to STEP 3 --
FUN <- function(sink, viewport) {
  block <- array(runif(length(viewport)), dim=dim(viewport))
  write_block(sink, viewport, block)
}
sink <- sinkApply(sink, FUN, grid=sink_grid, verbose=TRUE)

## -- STEP 4 --
## Close the sink and turn it into a DelayedArray object:
close(sink)
A <- as(sink, "DelayedArray")
A

setAutoRealizationBackend() # unset automatic realization backend

## -----
## USING THE "RealizationSink API": EXAMPLE 2
## -----

## Say we have a 3D array and want to collapse its 3rd dimension by
## summing the array elements that are stacked vertically, that is, we
## want to compute the matrix M obtained by doing sum(A[i, j, ]) for all
## valid i and j. This is very easy to do with an ordinary array:
collapse_3rd_dim <- function(a) apply(a, MARGIN=1:2, sum)

## or, in a slightly more efficient way:
collapse_3rd_dim <- function(a) {
  m <- matrix(0, nrow=nrow(a), ncol=ncol(a))
  for (z in seq_len(dim(a)[[3]]))
    m <- m + a[ , , z]
  m
}

```

```

## With a toy 3D array:
a <- array(runif(8000), dim=c(25, 40, 8))
dim(collapse_3rd_dim(a))
stopifnot(identical(sum(a), sum(collapse_3rd_dim(a)))) # sanity check

## Now say that A is so big that even M wouldn't fit in memory. This is
## a situation where we'd want to compute M block by block:

## -- STEP 1 --
## Create the 2D realization sink:
setAutoRealizationBackend("HDF5Array")
sink <- AutoRealizationSink(dim(a)[1:2])
dim(sink)

## -- STEP 2 --
## Define two grids: one for 'sink' and one for 'a'. Since we're going
## to walk on the two grids simultaneously, read a block from 'a' and
## write it to 'sink', we need to make sure that we define grids that
## are "aligned". More precisely, the two grids must have the same number
## of viewports, and the viewports in one must correspond to the viewports
## in the other one:
sink_grid <- colAutoGrid(sink, ncol=10)
a_spacings <- c(dim(sink_grid[[1L]]), dim(a)[[3]])
a_grid <- RegularArrayGrid(dim(a), spacings=a_spacings)
dims(sink_grid) # dimensions of the individual viewports
dims(a_grid)    # dimensions of the individual viewports

## Let's check that our two grids are actually "aligned":
stopifnot(identical(length(sink_grid), length(a_grid)))
stopifnot(identical(dims(sink_grid), dims(a_grid)[ , 1:2, drop=FALSE]))

## -- STEP 3 --
## Walk on the two grids simultaneously:
for (bid in seq_along(sink_grid)) {
  ## Read block from 'a'.
  a_viewport <- a_grid[[bid]]
  block <- read_block(a, a_viewport)
  ## Collapse it.
  block <- collapse_3rd_dim(block)
  ## Write the collapsed block to 'sink'.
  sink_viewport <- sink_grid[[bid]]
  sink <- write_block(sink, sink_viewport, block)
}

## -- An alternative to STEP 3 --
FUN <- function(sink, sink_viewport) {
  ## Read block from 'a'.
  bid <- currentBlockId()
  a_viewport <- a_grid[[bid]]
  block <- read_block(a, a_viewport)
  ## Collapse it.
  block <- collapse_3rd_dim(block)
  ## Write the collapsed block to 'sink'.

```

```

    write_block(sink, sink_viewport, block)
}
sink <- sinkApply(sink, FUN, grid=sink_grid, verbose=TRUE)

## -- STEP 4 --
## Close the sink and turn it into a DelayedArray object:
close(sink)
M <- as(sink, "DelayedArray")
M

## Sanity check:
stopifnot(identical(collapse_3rd_dim(a), as.array(M)))

setAutoRealizationBackend() # unset automatic realization backend

## -----
## USING THE "RealizationSink API": AN ADVANCED EXAMPLE
## -----

## Say we have 2 matrices with the same number of columns. Each column
## represents a biological sample:
library(HDF5Array)
R <- as(matrix(runif(75000), ncol=1000), "HDF5Array") # 75 rows
G <- as(matrix(runif(250000), ncol=1000), "HDF5Array") # 250 rows

## Say we want to compute the matrix U obtained by applying the same
## binary functions FUN() to all samples i.e. U is defined as:
##
##   U[ , j] <- FUN(R[ , j], G[ , j]) for 1 <= j <= 1000
##
## Note that FUN() should return a vector of constant length, say 200,
## so U will be a 200x1000 matrix. A naive implementation would be:
##
##   pFUN <- function(r, g) {
##     stopifnot(ncol(r) == ncol(g)) # sanity check
##     sapply(seq_len(ncol(r)), function(j) FUN(r[ , j], g[ , j]))
##   }
##
## But because U is going to be too big to fit in memory, we can't
## just do pFUN(R, G). So we want to compute U block by block and
## write the blocks to disk as we go. The blocks will be made of full
## columns. Also since we need to walk on 2 matrices at the same time
## (R and G), we can't use blockApply() or blockReduce() so we'll use
## a "for" loop.

## Before we get to the "for" loop, we need 4 things:

## 1. Two grids of blocks, one on R and one on G. The blocks in the
##    two grids must contain the same number of columns. We arbitrarily
##    choose to use blocks of 150 columns:
R_grid <- colAutoGrid(R, ncol=150)
G_grid <- colAutoGrid(G, ncol=150)

```

```

## 2. The function pFUN(). It will take 2 blocks as input, 1 from R
##    and 1 from G, apply FUN() to all the samples in the blocks,
##    and return a matrix with one columns per sample:
pFUN <- function(r, g) {
  stopifnot(ncol(r) == ncol(g)) # sanity check
  ## Return a matrix with 200 rows with random values. Completely
  ## artificial sorry. A realistic example would actually need to
  ## apply the same binary function to r[,j] and g[,j] for
  ## 1 <= j <= ncol(r).
  matrix(runif(200 * ncol(r)), nrow=200)
}

## 3. A RealizationSink derivative where to write the matrices returned
##    by pFUN() as we go:
setAutoRealizationBackend("HDF5Array")
U_sink <- AutoRealizationSink(c(200L, 1000L))

## 4. Finally, we create a grid on U_sink with viewports that contain
##    the same number of columns as the corresponding blocks in R and G:
U_grid <- colAutoGrid(U_sink, ncol=150)

## Note that the three grids should have the same number of viewports:
stopifnot(length(U_grid) == length(R_grid))
stopifnot(length(U_grid) == length(G_grid))

## 5. Now we can proceed. We use a "for" loop to walk on R and G
##    simultaneously, block by block, apply pFUN(), and write the
##    output of pFUN() to U_sink:
for (bid in seq_along(U_grid)) {
  R_block <- read_block(R, R_grid[[bid]])
  G_block <- read_block(G, G_grid[[bid]])
  U_block <- pFUN(R_block, G_block)
  U_sink <- write_block(U_sink, U_grid[[bid]], U_block)
}

## An alternative to the "for" loop is to use sinkApply():
FUN <- function(U_sink, U_viewport) {
  bid <- currentBlockId()
  R_block <- read_block(R, R_grid[[bid]])
  G_block <- read_block(G, G_grid[[bid]])
  U_block <- pFUN(R_block, G_block)
  write_block(U_sink, U_viewport, U_block)
}
U_sink <- sinkApply(U_sink, FUN, grid=U_grid, verbose=TRUE)

close(U_sink)
U <- as(U_sink, "DelayedArray")
U

setAutoRealizationBackend() # unset automatic realization backend

## -----
## VERY BASIC (BUT ALSO VERY ARTIFICIAL) USAGE OF THE

```



```

## read_block()/write_block() COMBO
## -----

##### On an ordinary matrix #####
m1 <- matrix(1:30, ncol=5)

## Define a viewport on 'm1':
block1_dim <- c(4, 3)
viewport1 <- ArrayViewport(dim(m1), IRanges(c(3, 2), width=block1_dim))

## Read/transform/write:
block1 <- read_block(m1, viewport1)
write_block(m1, viewport1, block1 + 1000L)

## Define another viewport on 'm1':
viewport1b <- ArrayViewport(dim(m1), IRanges(c(1, 3), width=block1_dim))

## Read/transform/write:
write_block(m1, viewport1b, block1 + 1000L)

## No-op:
m <- write_block(m1, viewport1, read_block(m1, viewport1))
stopifnot(identical(m1, m))

##### On a 3D array #####
a3 <- array(1:60, 5:3)

## Define a viewport on 'a3':
block3_dim <- c(2, 4, 1)
viewport3 <- ArrayViewport(dim(a3), IRanges(c(1, 1, 3), width=block3_dim))

## Read/transform/write:
block3 <- read_block(a3, viewport3)
write_block(a3, viewport3, block3 + 1000L)

## Define another viewport on 'a3':
viewport3b <- ArrayViewport(dim(a3), IRanges(c(3, 1, 3), width=block3_dim))

## Read/transform/write:
write_block(a3, viewport3b, block3 + 1000L)

## No-op:
a <- write_block(a3, viewport3, read_block(a3, viewport3))
stopifnot(identical(a3, a))

## -----
## LESS BASIC (BUT STILL VERY ARTIFICIAL) USAGE OF THE
## read_block()/write_block() COMBO
## -----

grid1 <- RegularArrayGrid(dim(m1), spacings=c(3L, 2L))
grid1
length(grid1) # number of blocks defined by the grid

```

```

read_block(m1, grid1[[3L]]) # read 3rd block
read_block(m1, grid1[[1L, 3L]])

## Walk on the grid, column by column:
m1a <- m1
for (bid in seq_along(grid1)) {
  viewport <- grid1[[bid]]
  block <- read_block(m1a, viewport)
  block <- bid * 1000L + block
  m1a <- write_block(m1a, viewport, block)
}
m1a

## Walk on the grid, row by row:
m1b <- m1
for (i in seq_len(dim(grid1)[[1]])) {
  for (j in seq_len(dim(grid1)[[2]])) {
    viewport <- grid1[[i, j]]
    block <- read_block(m1b, viewport)
    block <- (i * 10L + j) * 1000L + block
    m1b <- write_block(m1b, viewport, block)
  }
}
m1b

## -----
## supportedRealizationBackends() AND FAMILY
## -----

getAutoRealizationBackend() # no backend set yet

supportedRealizationBackends()
setAutoRealizationBackend("HDF5Array")
getAutoRealizationBackend() # backend is set to "HDF5Array"
supportedRealizationBackends()

getHDF5DumpChunkLength()
setHDF5DumpChunkLength(500L)
getHDF5DumpChunkShape()

sink <- AutoRealizationSink(c(120L, 50L))
class(sink) # HDF5-specific realization sink
dim(sink)
chunkdim(sink)

grid <- defaultSinkAutoGrid(sink, block.length=600)
for (bid in seq_along(grid)) {
  viewport <- grid[[bid]]
  block <- 101 * bid + runif(length(viewport))
  dim(block) <- dim(viewport)
  sink <- write_block(sink, viewport, block)
}

```

```
close(sink)
A <- as(sink, "DelayedArray")
A

setAutoRealizationBackend() # unset automatic realization backend
```

realize	<i>Realize a DelayedArray object</i>
---------	--------------------------------------

Description

Realize a [DelayedArray](#) object in memory or on disk.

Usage

```
realize(x, ...)

## S4 method for signature 'ANY'
realize(x, BACKEND=getAutoRealizationBackend())
```

Arguments

x	The array-like object to realize.
...	Additional arguments passed to methods.
BACKEND	A single string specifying the name of the <i>realization backend</i> . Use the current <i>automatic realization backend</i> by default i.e. the backend returned by getAutoRealizationBackend() .

Value

A [DelayedArray](#) object. More precisely, it returns `DelayedArray(as.array(x))` when the backend is set to NULL (the default). Otherwise it returns an instance of the class associated with the specified backend (which should extend [DelayedArray](#)).

See Also

- [getAutoRealizationBackend](#) and [setAutoRealizationBackend](#) for getting and setting the current *automatic realization backend*.
- [DelayedArray](#) objects.
- [RleArray](#) objects.
- [HDF5Array](#) objects in the **HDF5Array** package.
- [array](#) objects in base R.

Examples

```
library(HDF5Array)
toy_h5 <- system.file("extdata", "toy.h5", package="HDF5Array")
h5ls(toy_h5)
M1 <- HDF5Array(toy_h5, "M1")
M2 <- HDF5Array(toy_h5, "M2")
M3 <- rbind(log(M1), t(M2))

supportedRealizationBackends()
getAutoRealizationBackend() # backend is set to NULL
realize(M3) # realization as ordinary array

setAutoRealizationBackend("RleArray")
getAutoRealizationBackend() # backend is set to "RleArray"
realize(M3) # realization as RleArray object

setAutoRealizationBackend("HDF5Array")
getAutoRealizationBackend() # backend is set to "HDF5Array"
realize(M3) # realization in HDF5 file
```

RleArray-class

RleArray objects

Description

The RleArray class is a [DelayedArray](#) subclass for representing an in-memory Run Length Encoded array-like dataset.

All the operations available for [DelayedArray](#) objects work on RleArray objects.

Usage

```
## Constructor function:
RleArray(data, dim, dimnames, chunksize=NULL)
```

Arguments

data	An Rle object, or an ordinary list of Rle objects, or an RleList object, or a DataFrame object where all the columns are Rle objects. More generally speaking, data can be any list-like object where all the list elements are Rle objects.
dim	The dimensions of the object to be created, that is, an integer vector of length one or more giving the maximal indices in each dimension.
dimnames	The <i>dimnames</i> of the object to be created. Must be NULL or a list of length the number of dimensions. Each list element must be either NULL or a character vector along the corresponding dimension.
chunksize	Experimental. Don't use!

Value

An RleArray (or RleMatrix) object. (Note that RleMatrix extends RleArray.)

See Also

- [Rle](#) and [DataFrame](#) objects in the **S4Vectors** package and [RleList](#) objects in the **IRanges** package.
- [DelayedArray](#) objects.
- [DelayedArray-utils](#) for common operations on [DelayedArray](#) objects.
- [realize](#) for realizing a DelayedArray object in memory or on disk.
- [ConstantArray](#) objects for mimicking an array containing a constant value, without actually creating said array in memory.
- [HDF5Array](#) objects in the **HDF5Array** package.
- The [RleArraySeed](#) helper class.

Examples

```
## -----
## A. BASIC EXAMPLE
## -----

data <- Rle(sample(6L, 500000, replace=TRUE), 8)
a <- array(data, dim=c(50, 20, 4000)) # array() expands the Rle object
                                     # internally with as.vector()

A <- RleArray(data, dim=c(50, 20, 4000)) # Rle object is NOT expanded
A

object.size(a)
object.size(A)

stopifnot(identical(a, as.array(A)))

as(A, "Rle") # deconstruction

toto <- function(x) (5 * x[, , 1] ^ 3 + 1L) * log(x[, , 2])
m1 <- toto(a)
head(m1)

M1 <- toto(A) # very fast! (operations are delayed)
M1

stopifnot(identical(m1, as.array(M1)))

cs <- colSums(m1)
CS <- colSums(M1)
stopifnot(identical(cs, CS))

## Coercing a DelayedMatrix object to DataFrame produces a DataFrame
```

```

## object with Rle columns:
as(M1, "DataFrame")

## -----
## B. MAKING AN RleArray OBJECT FROM A LIST-LIKE OBJECT OF Rle OBJECTS
## -----

## From a DataFrame object:
DF <- DataFrame(A=Rle(sample(3L, 100, replace=TRUE)),
                B=Rle(sample(3L, 100, replace=TRUE)),
                C=Rle(sample(3L, 100, replace=TRUE) - 0.5),
                row.names=sprintf("ID%03d", 1:100))

M2 <- RleArray(DF)
M2

A3 <- RleArray(DF, dim=c(25, 6, 2))
A3

M4 <- RleArray(DF, dim=c(25, 12), dimnames=list(LETTERS[1:25], NULL))
M4

## From an ordinary list:
## If all the supplied Rle objects have the same length and if the 'dim'
## argument is not specified, then the RleArray() constructor returns an
## RleMatrix object with 1 column per Rle object. If the 'dimnames'
## argument is not specified, then the names on the list are propagated
## as the colnames of the returned object.
data <- as.list(DF)
M2b <- RleArray(data)
A3b <- RleArray(data, dim=c(25, 6, 2))
M4b <- RleArray(data, dim=c(25, 12), dimnames=list(LETTERS[1:25], NULL))

data2 <- list(Rle(sample(3L, 9, replace=TRUE)) * 11L,
              Rle(sample(3L, 15, replace=TRUE)))
## Not run:
  RleArray(data2) # error! (cannot infer the dim)

## End(Not run)
RleArray(data2, dim=c(4, 6))

## From an RleList object:
data <- RleList(data)
M2c <- RleArray(data)
A3c <- RleArray(data, dim=c(25, 6, 2))
M4c <- RleArray(data, dim=c(25, 12), dimnames=list(LETTERS[1:25], NULL))

data2 <- RleList(data2)
## Not run:
  RleArray(data2) # error! (cannot infer the dim)

## End(Not run)
RleArray(data2, dim=4:2)

```

```

## Sanity checks:
data0 <- as.vector(unlist(DF, use.names=FALSE))
m2 <- matrix(data0, ncol=3, dimnames=dimnames(M2))
stopifnot(identical(m2, as.matrix(M2)))
rownames(m2) <- NULL
stopifnot(identical(m2, as.matrix(M2b)))
stopifnot(identical(m2, as.matrix(M2c)))
a3 <- array(data0, dim=c(25, 6, 2))
stopifnot(identical(a3, as.array(A3)))
stopifnot(identical(a3, as.array(A3b)))
stopifnot(identical(a3, as.array(A3c)))
m4 <- matrix(data0, ncol=12, dimnames=dimnames(M4))
stopifnot(identical(m4, as.matrix(M4)))
stopifnot(identical(m4, as.matrix(M4b)))
stopifnot(identical(m4, as.matrix(M4c)))

## -----
## C. COERCING FROM RleList OR DataFrame TO RleMatrix
## -----

## Coercing an RleList object to RleMatrix only works if all the list
## elements in the former have the same length.
x <- RleList(A=Rle(sample(3L, 20, replace=TRUE)),
             B=Rle(sample(3L, 20, replace=TRUE)))
M <- as(x, "RleMatrix")
stopifnot(identical(x, as(M, "RleList")))

x <- DataFrame(A=x[[1]], B=x[[2]], row.names=letters[1:20])
M <- as(x, "RleMatrix")
stopifnot(identical(x, as(M, "DataFrame")))

## -----
## D. CONSTRUCTING A LARGE RleArray OBJECT
## -----

## The RleArray() constructor does not accept a "long" Rle object (i.e.
## an object of length > .Machine$integer.max) at the moment:
## Not run:
RleArray(Rle(5, 3e9), dim=c(3, 1e9)) # error!

## End(Not run)

## The workaround is to supply a list of Rle objects instead:

toy_Rle <- function() {
  run_lens <- c(sample(4), sample(rep(c(1:19, 40) * 3, 6e4)), sample(4))
  run_vals <- sample(700, length(run_lens), replace=TRUE) / 5
  Rle(run_vals, run_lens)
}
rle_list <- lapply(1:80, function(j) toy_Rle()) # takes about 20 sec.

## Cumulative length of all the Rle objects is > .Machine$integer.max:

```

```

sum(lengths(rle_list)) # 3.31e+09

## Feed 'rle_list' to the RleArray() constructor:
dim <- c(14395, 320, 719)
A <- RleArray(rle_list, dim)
A

## Because all the Rle objects in 'rle_list' have the same length, we
## can call RleArray() on it without specifying the 'dim' argument. This
## returns an RleMatrix object where each column corresponds to an Rle
## object in 'rle_list':
M <- RleArray(rle_list)
M
stopifnot(identical(as(rle_list, "RleList"), as(M, "RleList")))

## -----
## E. CHANGING THE TYPE OF AN RleArray OBJECT FROM "double" TO "integer"
## -----

## An RleArray object is an in-memory object so it can be useful to
## reduce its memory footprint. For an object of type "double" this can
## be done by changing its type to "integer" (integers are half the size
## of doubles in memory). Of course this only makes sense if this results
## in a loss of precision that is acceptable.
## On an ordinary array (or matrix) 'a', this is simply a matter of
## doing 'storage.mode(a) <- "integer"'. However, with a DelayedArray
## object, things are a little bit different. Let's do this on a subset
## of the RleMatrix object 'M' created in the previous section.

M1 <- as(M[1:6e5, ], "RleMatrix")
rm(M)

## First of all, it's important to be aware that object.size() (from
## package utils) is NOT reliable on RleArray objects! This is because
## the data in an RleArray object is stored in an environment and
## object.size() stubbornly refuses to take the content of an environment
## into account when computing its size:
object.size(list2env(list(aa=1:10))) # 56 bytes
object.size(list2env(list(aa=1:1e6))) # always 56 bytes!

## So we'll use obj_size() instead (from package lobstr):
library(lobstr)
obj_size(list2env(list(aa=1:10))) # 264 B
obj_size(list2env(list(aa=1:1e6))) # 4 MB
obj_size(list2env(list(aa=as.double(1:1e6)))) # 8 MB

obj_size(M1) # 16.7 MB

type(M1) <- "integer" # Delayed!
M1
# Note the class: it's no longer RleMatrix!
# (That's because the object now carries delayed
# operations.)

```



```
## Because changing the type is a delayed operation, the memory footprint
## of the object has not changed yet (remember that the original data in
## a DelayedArray object is stored in its "seed" and its seed is never
## modified **in-place**, that is, no operation on the object will ever
## modify its seed):
obj_size(M1) # Still the same (well, a very tiny more, because the
              # object is now carrying one more delayed operation,
              # the `type<-` operation)

## To effectively reduce the memory footprint of the object, a new object
## needs to be created. This is achieved simply by **realizing** M1 as a
## (new) RleArray object. Note that this realization will use block
## processing:

DelayedArray:::set_verbose_block_processing(TRUE) # See block processing
                                                    # in action.
getAutoBlockSize()      # Automatic block size (100 Mb by default).
setAutoBlockSize(20e6)  # Set automatic block size to 20 Mb.

M2 <- as(M1, "RleArray")
DelayedArray:::set_verbose_block_processing(FALSE)
setAutoBlockSize()      # Reset automatic block size to factory settings.

M2
obj_size(M2) # 6.91 MB (Less than half the original size! This is
              # because RleArray objects use some internal tricks to
              # reduce memory footprint even more when the data in
              # their seed is of type "integer".)

## Finally note that the 2-step approach described here (i.e.
## type(A) <- "integer" followed by realization) is generic and works
## on any kind of DelayedArray object or derivative. In particular,
## after doing 'type(A) <- "integer"', 'A' can be realized as anything
## as long as the realization backend is supported (e.g. could be
## 'as(A, "HDF5Array")' or 'as(A, "TENxMatrix")') and realization will
## always use block processing so the array data will never be fully
## loaded in memory.
```

RleArraySeed-class	<i>RleArraySeed objects</i>
--------------------	-----------------------------

Description

RleArraySeed is a low-level helper class for representing an in-memory Run Length Encoded array-like dataset. RleArraySeed objects are not intended to be used directly. Most end users should create and manipulate [RleArray](#) objects instead. See [?RleArray](#) for more information.

Details

No operation can be performed directly on an `RleArraySeed` object. It first needs to be wrapped in a `DelayedArray` object. The result of this wrapping is an `RleArray` object (an `RleArray` object is just an `RleArraySeed` object wrapped in a `DelayedArray` object).

See Also

- `RleArray` objects.
- `Rle` objects in the `S4Vectors` package.

showtree

Visualize and access the leaves of a tree of delayed operations

Description

`showtree` can be used to visualize the tree of delayed operations carried by a `DelayedArray` object. Use `nseed`, `seed`, or `path` to access the number of seeds, the seed, or the seed path of a `DelayedArray` object, respectively.

Use `seedApply` to apply a function to the seeds of a `DelayedArray` object.

Usage

```
showtree(x, show.node.dim=TRUE)

nseed(x)          # seed counter
seed(x)           # seed getter and setter
path(object, ...) # path getter and setter

seedApply(x, FUN, ...)
```

Arguments

<code>x, object</code>	Typically a <code>DelayedArray</code> object but can also be a <code>DelayedOp</code> object or a list where each element is a <code>DelayedArray</code> or <code>DelayedOp</code> object.
<code>show.node.dim</code>	TRUE or FALSE. If TRUE (the default), the nodes dimensions and data type are displayed.
<code>FUN</code>	The function to be applied to each leaf in <code>x</code> .
<code>...</code>	Optional arguments to <code>FUN</code> for <code>seedApply()</code> . Additional arguments passed to methods for <code>path()</code> .

Value

The number of seeds contained in `x` for `nseed`.
 The seed contained in `x` for `seed`.
 The path of the seed contained in `object` for `path`.
 A list of length `nseed(x)` for `seedApply`.

See Also

- [simplify](#) to simplify the tree of delayed operations carried by a [DelayedArray](#) object.
- [DelayedOp](#) objects.
- [DelayedArray](#) objects.

Examples

```
## -----
## showtree(), nseed(), and seed()
## -----
m1 <- matrix(runif(150), nrow=15, ncol=10)
M1 <- DelayedArray(m1)
showtree(M1)
seed(M1)

M2 <- log(t(M1[5:1, c(TRUE, FALSE)] + 10))[-1, ]
showtree(M2)

## In the above example, the tree is linear i.e. all the operations
## are represented by unary nodes. The simplest way to know if a
## tree is linear is by counting its leaves with nseed():
nseed(M2) # only 1 leaf means the tree is linear
seed(M2)

dimnames(M1) <- list(letters[1:15], LETTERS[1:10])
showtree(M1)

m2 <- matrix(1:20, nrow=10)
Y <- cbind(t(M1[, 10:1]), DelayedArray(m2), M1[6:15, "A", drop=FALSE])
showtree(Y)
showtree(Y, show.node.dim=FALSE)
nseed(Y) # the tree is not linear

Z <- t(Y[10:1, ])[1:15, ] + 0.4 * M1
showtree(Z)
nseed(Z) # the tree is not linear

## -----
## seedApply()
## -----
seedApply(Y, class)
seedApply(Y, dim)
```

Description

NOTE: The tools documented in this man page are primarily intended for developers or advanced users curious about the internals of the **DelayedArray** package. End users typically don't need them for their regular use of **DelayedArray** objects.

In a **DelayedArray** object, the delayed operations are stored as a tree of **DelayedOp** objects. See **?DelayedOp** for more information about this tree.

`simplify` can be used to simplify the tree of delayed operations in a **DelayedArray** object.

`isPristine` can be used to know whether a **DelayedArray** object is *pristine* or not. A **DelayedArray** object is considered *pristine* when it carries no delayed operation. Note that an object that carries delayed operations that do nothing (e.g. $A + 0$) is not considered *pristine*.

`contentIsPristine` can be used to know whether the delayed operations in a **DelayedArray** object *touch* its array elements or not.

`netSubsetAndAperm` returns an object that represents the *net subsetting* and *net dimension rearrangement* of all the delayed operations in a **DelayedArray** object.

Usage

```
simplify(x, incremental=FALSE)

isPristine(x, ignore.dimnames=FALSE)
contentIsPristine(x)
netSubsetAndAperm(x, as.DelayedOp=FALSE)
```

Arguments

<code>x</code>	Typically a DelayedArray object but can also be a DelayedOp object (except for <code>isPristine</code>).
<code>incremental</code>	For internal use.
<code>ignore.dimnames</code>	TRUE or FALSE. When TRUE, the object is considered <i>pristine</i> even if its dimnames have been modified and no longer match the dimnames of its seed (in which case the object carries a single delayed operations of type DelayedSetDimnames).
<code>as.DelayedOp</code>	TRUE or FALSE. Controls the form of the returned object. See details below.

Details

`netSubsetAndAperm` is only supported on a **DelayedArray** object `x` with a single seed i.e. `nseed(x) == 1`.

The mapping between the array elements of `x` and the array elements of its seed is affected by the following delayed operations carried by `x`: `[], drop()`, and `aperm()`. `x` can carry any number of each of these operations in any order but their net result can always be described by a *net subsetting* followed by a *net dimension rearrangement*.

`netSubsetAndAperm(x)` returns an object that represents the *net subsetting* and *net dimension rearrangement*. The `as.DelayedOp` argument controls in what form this object should be returned:

- If `as.DelayedOp` is FALSE (the default), the returned object is a list of subscripts that describes the *net subsetting*. The list contains one subscript per dimension in the seed. Each subscript can be either a vector of positive integers or a NULL. A NULL indicates a *missing subscript*. In addition, if `x` carries delayed operations that rearrange its dimensions (i.e. operations that drop and/or permute some of the original dimensions), the *net dimension rearrangement* is described in a `dimmap` attribute added to the list. This attribute is an integer vector parallel to `dim(x)` that reports how the dimensions of `x` are mapped to the dimensions of its seed.
- If `as.DelayedOp` is TRUE, the returned object is a linear tree with 2 [DelayedOp](#) nodes and a leaf node. The leaf node is the seed of `x`. Walking the tree from the seed, the 2 [DelayedOp](#) nodes are of type [DelayedSubset](#) and [DelayedAperm](#), in that order (this reflects the order in which the operations apply). More precisely, the returned object is a [DelayedAperm](#) object with one child (the [DelayedSubset](#) object), and one grandchild (the seed of `x`). The [DelayedSubset](#) and [DelayedAperm](#) nodes represent the *net subsetting* and *net dimension rearrangement*, respectively. Either or both of them can be a no-op.

Note that the returned object describes how the array elements of `x` map to their corresponding array element in `seed(x)`.

Value

The simplified object for `simplify`.

TRUE or FALSE for `contentIsPristine`.

An ordinary list (possibly with the `dimmap` attribute on it) for `netSubsetAndAperm`. Unless `as.DelayedOp` is set to TRUE, in which case a [DelayedAperm](#) object is returned (see Details section above for more information).

See Also

- [showtree](#) to visualize and access the leaves of a tree of delayed operations carried by a [DelayedArray](#) object.
- [DelayedOp](#) objects.
- [DelayedArray](#) objects.

Examples

```
## -----
## Simplification of the tree of delayed operations
## -----
m1 <- matrix(runif(150), nrow=15, ncol=10)
M1 <- DelayedArray(m1)
showtree(M1)

## By default, the tree of delayed operations carried by a DelayedArray
## object gets simplified each time a delayed operation is added to it.
## This can be disabled via a global option:
options(DelayedArray.simplify=FALSE)
M2 <- log(t(M1[5:1, c(TRUE, FALSE)] + 10))[-1, ]
showtree(M2) # linear tree
```

```

## Note that as part of the simplification process, some operations
## can be reordered:
options(DelayedArray.simplify=TRUE)
M2 <- log(t(M1[5:1, c(TRUE, FALSE)] + 10))[-1, ]
showtree(M2) # linear tree

options(DelayedArray.simplify=FALSE)

dimnames(M1) <- list(letters[1:15], LETTERS[1:10])
showtree(M1) # linear tree

m2 <- matrix(1:20, nrow=10)
Y <- cbind(t(M1[, 10:1]), DelayedArray(m2), M1[6:15, "A", drop=FALSE])
showtree(Y) # non-linear tree

Z <- t(Y[10:1, ])[1:15, ] + 0.4 * M1
showtree(Z) # non-linear tree

Z@seed@seeds
Z@seed@seeds[[2]]@seed # reaching to M1
Z@seed@seeds[[1]]@seed@seed@seed@seed@seed # reaching to Y

## -----
## isPristine()
## -----
m <- matrix(1:20, ncol=4, dimnames=list(letters[1:5], NULL))
M <- DelayedArray(m)

isPristine(M) # TRUE
isPristine(log(M)) # FALSE
isPristine(M + 0) # FALSE
isPristine(t(M)) # FALSE
isPristine(t(t(M))) # TRUE
isPristine(cbind(M, M)) # FALSE
isPristine(cbind(M)) # TRUE

dimnames(M) <- NULL
isPristine(M) # FALSE
isPristine(M, ignore.dimnames=TRUE) # TRUE
isPristine(t(t(M)), ignore.dimnames=TRUE) # TRUE
isPristine(cbind(M, M), ignore.dimnames=TRUE) # FALSE

## -----
## contentIsPristine()
## -----
a <- array(1:40, c(4, 5, 2))
A <- DelayedArray(a)

stopifnot(contentIsPristine(A))
stopifnot(contentIsPristine(A[1, , ]))
stopifnot(contentIsPristine(t(A[1, , ])))
stopifnot(contentIsPristine(cbind(A[1, , ], A[2, , ])))
dimnames(A) <- list(LETTERS[1:4], letters[1:5], NULL)

```

```

stopifnot(contentIsPristine(A))

contentIsPristine(log(A))      # FALSE
contentIsPristine(A - 11:14)  # FALSE
contentIsPristine(A * A)      # FALSE

## -----
## netSubsetAndAperm()
## -----

a <- array(1:40, c(4, 5, 2))
M <- aperm(DelayedArray(a)[ , -1, ] / 100)[ , , 3] + 99:98
M
showtree(M)

netSubsetAndAperm(M) # 1st dimension was dropped, 2nd and 3rd
                     # dimension were permuted (transposition)

op2 <- netSubsetAndAperm(M, as.DelayedOp=TRUE)
op2      # 2 nested delayed operations
op1 <- op2@seed
class(op1)      # DelayedSubset
class(op2)      # DelayedAperm
op1@index
op2@perm

DelayedArray(op2) # same as M from a [, drop(), and aperm() point of
                  # view but the individual array elements are now
                  # reset to their original values i.e. to the values
                  # they have in the seed
stopifnot(contentIsPristine(DelayedArray(op2)))

## A simple function that returns TRUE if a DelayedArray object carries
## no "net subsetting" and no "net dimension rearrangement":
is_aligned_with_seed <- function(x)
{
  if (nseed(x) != 1L)
    return(FALSE)
  op2 <- netSubsetAndAperm(x, as.DelayedOp=TRUE)
  op1 <- op2@seed
  is_noop(op1) && is_noop(op2)
}

M <- DelayedArray(a[ , , 1])
is_aligned_with_seed(log(M + 11:14) > 3)      # TRUE
is_aligned_with_seed(M[4:1, ])                # FALSE
is_aligned_with_seed(M[4:1, ][4:1, ])        # TRUE
is_aligned_with_seed(t(M))                   # FALSE
is_aligned_with_seed(t(t(M)))                 # TRUE
is_aligned_with_seed(t(0.5 * t(M[4:1, ])[ , 4:1])) # TRUE

options(DelayedArray.simplify=TRUE)

```

SparseArraySeed-class *SparseArraySeed objects*

Description

SparseArraySeed objects are used internally to support block processing of array-like objects.

Usage

```
## Constructor function:
SparseArraySeed(dim, nzindex=NULL, nzdata=NULL, dimnames=NULL, check=TRUE)

## Getters (in addition to dim(), length(), and dimnames()):
nzindex(x)
nzdata(x)
sparsity(x)

## Two low-level utilities:
dense2sparse(x)
sparse2dense(sas)
```

Arguments

dim	The dimensions (specified as an integer vector) of the SparseArraySeed object to create.
nzindex	A matrix containing the array indices of the nonzero data. This must be an integer matrix like one returned by <code>base::arrayInd</code> , that is, with <code>length(dim)</code> columns and where each row is an n-uplet representing an <i>array index</i> .
nzdata	A vector (atomic or list) of length <code>nrow(nzindex)</code> containing the nonzero data.
dimnames	The <i>dimnames</i> of the object to be created. Must be NULL or a list of length the number of dimensions. Each list element must be either NULL or a character vector along the corresponding dimension.
check	Should the object be validated upon construction?
x	A SparseArraySeed object for the <code>nzindex</code> , <code>nzdata</code> , and <code>sparsity</code> getters. An array-like object for <code>dense2sparse</code> .
sas	A SparseArraySeed object.

Value

- For `SparseArraySeed()`: A SparseArraySeed instance.
- For `nzindex()`: The matrix containing the array indices of the nonzero data.
- For `nzdata()`: The vector of nonzero data.

- For `sparsity()`: The number of zero-valued elements in the implicit array divided by the total number of array elements (a.k.a. the length of the array).
- For `dense2sparse()`: A `SparseArraySeed` instance.
- For `sparse2dense()`: An ordinary array.

See Also

- [SparseArraySeed-utils](#) for native operations on `SparseArraySeed` objects.
- S4 classes [dgCMatrix](#), [dgRMatrix](#), and [lsparseMatrix](#), defined in the **Matrix** package, for the de facto standard of sparse matrix representations in R.
- The [read_block](#) function in the **S4Arrays** package.
- [blockApply](#) and family for convenient block processing of an array-like object.
- [extract_array](#) in the **S4Arrays** package.
- [DelayedArray](#) objects.
- [arrayInd](#) in the **base** package.
- [array](#) objects in base R.

Examples

```
## -----
## EXAMPLE 1
## -----
dim1 <- 5:3
nzindex1 <- Lindex2Mindex(sample(60, 8), 5:3)
nzdata1 <- 11.11 * seq_len(nrow(nzindex1))
sas1 <- SparseArraySeed(dim1, nzindex1, nzdata1)

dim(sas1)      # the dimensions of the implicit array
length(sas1)   # the length of the implicit array
nzindex(sas1)
nzdata(sas1)
type(sas1)
sparsity(sas1)

sparse2dense(sas1)
as.array(sas1) # same as sparse2dense(sas1)

## Not run:
as.matrix(sas1) # error!

## End(Not run)
## -----
## EXAMPLE 2
## -----
m2 <- matrix(c(5:-2, rep.int(c(0L, 99L), 11)), ncol=6)
sas2 <- dense2sparse(m2)
class(sas2)
dim(sas2)
length(sas2)
```

```

nzindex(sas2)
nzdata(sas2)
type(sas2)
sparsity(sas2)

stopifnot(identical(sparse2dense(sas2), m2))

as.matrix(sas2) # same as sparse2dense(sas2)

t(sas2)
stopifnot(identical(as.matrix(t(sas2)), t(as.matrix(sas2))))

## -----
## COERCION FROM/TO dg[C|R]Matrix OR lg[C|R]Matrix OBJECTS
## -----
## dg[C|R]Matrix and lg[C|R]Matrix objects are defined in the Matrix
## package.

## dgCMatrix/dgRMatrix:

M2C <- as(sas2, "dgCMatrix")
stopifnot(identical(M2C, as(m2, "dgCMatrix")))

sas2C <- as(M2C, "SparseArraySeed")
## 'sas2C' is the same as 'sas2' except that 'nzdata(sas2C)' has
## type "double" instead of "integer":
stopifnot(all.equal(sas2, sas2C))
typeof(nzdata(sas2C)) # double
typeof(nzdata(sas2))  # integer

M2R <- as(sas2, "dgRMatrix")
stopifnot(identical(M2R, as(m2, "dgRMatrix")))
sas2R <- as(M2R, "SparseArraySeed")
stopifnot(all.equal(as.matrix(sas2), as.matrix(sas2R)))

## lgCMatrix/lgRMatrix:

m3 <- m2 == 99 # logical matrix
sas3 <- dense2sparse(m3)
class(sas3)
type(sas3)

M3C <- as(sas3, "lgCMatrix")
stopifnot(identical(M3C, as(m3, "lgCMatrix")))
sas3C <- as(M3C, "SparseArraySeed")
identical(as.matrix(sas3), as.matrix(sas3C))

M3R <- as(sas3, "lgRMatrix")
#stopifnot(identical(M3R, as(m3, "lgRMatrix")))
sas3R <- as(M3R, "SparseArraySeed")
identical(as.matrix(sas3), as.matrix(sas3R))

## -----

```

```
## SEED CONTRACT
## -----
## SparseArraySeed objects comply with the "seed contract".
## In particular they support extract_array():
extract_array(sas1, list(c(5, 3:2, 5), NULL, 3))

## See '?extract_array' in the S4Arrays package for more information
## about the "seed contract".

## This means that they can be wrapped in a DelayedArray object:
A1 <- DelayedArray(sas1)
A1

## A big very sparse DelayedMatrix object:
nzindex4 <- cbind(sample(25000, 600000, replace=TRUE),
                  sample(195000, 600000, replace=TRUE))
nzdata4 <- runif(600000)
sas4 <- SparseArraySeed(c(25000, 195000), nzindex4, nzdata4)
sparsity(sas4)

M4 <- DelayedArray(sas4)
M4
colSums(M4[, 1:20])
```

SparseArraySeed-utils *Operate natively on SparseArraySeed objects*

Description

Some utilities to operate natively on [SparseArraySeed](#) objects. Mostly for internal use by the **DelayedArray** package e.g. they support block processed methods for sparse [DelayedArray](#) objects like `sum()`, `mean()`, `which()`, etc...

Usage

```
## S4 method for signature 'SparseArraySeed'
is.na(x)

## S4 method for signature 'SparseArraySeed'
is.infinite(x)

## S4 method for signature 'SparseArraySeed'
is.nan(x)

## S4 method for signature 'SparseArraySeed'
tolower(x)

## S4 method for signature 'SparseArraySeed'
toupper(x)
```

```

## S4 method for signature 'SparseArraySeed'
nchar(x, type="chars", allowNA=FALSE, keepNA=NA)

## S4 method for signature 'SparseArraySeed'
anyNA(x, recursive=FALSE)

## S4 method for signature 'SparseArraySeed'
which(x, arr.ind=FALSE, useNames=TRUE)

## <-<-<-< "Summary" group generic <-<-<-<

## S4 method for signature 'SparseArraySeed'
max(x, ..., na.rm=FALSE)

## S4 method for signature 'SparseArraySeed'
min(x, ..., na.rm=FALSE)

## S4 method for signature 'SparseArraySeed'
range(x, ..., finite=FALSE, na.rm=FALSE)

## S4 method for signature 'SparseArraySeed'
sum(x, ..., na.rm=FALSE)

## S4 method for signature 'SparseArraySeed'
prod(x, ..., na.rm=FALSE)

## S4 method for signature 'SparseArraySeed'
any(x, ..., na.rm=FALSE)

## S4 method for signature 'SparseArraySeed'
all(x, ..., na.rm=FALSE)

## <-<-<-< others <-<-<-<

## S4 method for signature 'SparseArraySeed'
mean(x, na.rm=FALSE)

```

Arguments

<code>x</code>	A SparseArraySeed object.
<code>type</code> , <code>allowNA</code> , <code>keepNA</code>	See ?base:: nchar for a description of these arguments.
<code>recursive</code>	See ?base:: anyNA for a description of this argument.
<code>arr.ind</code>	See ?base:: which for a description of this argument.
<code>useNames</code>	Ignored.

...	Unsupported.
na.rm	TRUE or FALSE (the default). Should NA's and NaN's be removed?
finite	TRUE or FALSE (the default). Should non-finite values be removed?

Value

See corresponding functions in the **base** package.

See Also

- [SparseArraySeed](#) objects.

Examples

```
## Create a SparseArraySeed object:
dim1 <- 5:3
nzindex1 <- Lindex2Mindex(sample(60, 14), 5:3)
sas1 <- SparseArraySeed(dim1, nzindex1, nzdata=sample(0:13))

## Apply native operations:
sum(sas1)
range(sas1)
mean(sas1)

## Sanity checks:
stopifnot(identical(sum(as.array(sas1)), sum(sas1)))
stopifnot(identical(range(as.array(sas1)), range(sas1)))
stopifnot(identical(mean(as.array(sas1)), mean(sas1)))
```

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