

Package ‘DelayedDataFrame’

February 1, 2026

Title Delayed operation on DataFrame using standard DataFrame metaphor

Version 1.27.0

Description Based on the standard DataFrame metaphor, we are trying to implement the feature of delayed operation on the DelayedDataFrame, with a slot of lazyIndex, which saves the mapping indexes for each column of DelayedDataFrame. Methods like show, validity check, [/[subsetting, rbind/cbind are implemented for DelayedDataFrame to be operated around lazyIndex. The listData slot stays untouched until a realization call e.g., DataFrame constructor OR as.list() is invoked.

biocViews Infrastructure, DataRepresentation

Depends R (>= 3.6), S4Vectors (>= 0.23.19), DelayedArray (>= 0.7.5)

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Encoding UTF-8

URL <https://github.com/Bioconductor/DelayedDataFrame>

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

Imports methods, stats, BiocGenerics

RoxygenNote 7.0.2

Suggests testthat, knitr, rmarkdown, BiocStyle, SeqArray, GDSArray

Collate LazyIndex-class.R DelayedDataFrame-class.R
DelayedDataFrame-method.R

VignetteBuilder knitr

git_url <https://git.bioconductor.org/packages/DelayedDataFrame>

git_branch devel

git_last_commit 7dcd028

git_last_commit_date 2025-10-29

Repository Bioconductor 3.23

Date/Publication 2026-02-01

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as.list,DelayedDataFrame-method
DelayedDataFrame related methods.

Description

as.list, rbind would incur realization of the lazyIndex slot in DelayedDataFrame object.
 cbind for DelayedDataFrame inherits the lazyIndex's if inputs have any DelayedDataFrame objects. Otherwise, return a new DelayedDataFrame with NULL lazyIndexes.

Usage

```
## S4 method for signature 'DelayedDataFrame'
as.list(x, use.names = TRUE)

## S4 method for signature 'DelayedDataFrame'
names(x)

## S4 method for signature 'DelayedDataFrame'
cbind(..., deparse.level = 1)

## S4 method for signature 'DelayedDataFrame'
bindROWS(
  x,
  objects = list(),
  use.names = TRUE,
  ignore.mcols = FALSE,
  check = TRUE
)

## S4 method for signature 'DelayedDataFrame,ANY'
extractROWS(x, i)
```

```

## S4 method for signature 'DelayedDataFrame'
extractCOLS(x, i)

## S4 method for signature 'DelayedDataFrame'
replaceCOLS(x, i, value)

## S4 method for signature 'DelayedDataFrame'
mergeROWS(x, i, value)

## S4 method for signature 'DelayedDataFrame,ANY,ANY,ANY'
x[i, j, ..., drop = TRUE]

```

Arguments

x	as.list,DelayedDataFrame: a DelayedDataFrame object. OR, [,DelayedDataFrame: DelayedDataFrame object to be subsetted.
use.names	as.list,DelayedDataFrame: whether to use the colnames of DelayedDataFrame as the names for the returned list. OR, bindROWS,DelayedDataFrame: whether to use rownames of the input arguments. Default is TRUE.
...	cbind,DelayedDataFrame: One or more vector-like or matrix-like objects. These can be given as named arguments. OR, [,DelayedDataFrame: other arguments to pass.
deparse.level	See ‘?base::cbind’ for a description of this argument.
objects	the DelayedDataFrame objects to be passed into bindROWS.
ignore.mcols	Logical. This argument is ignored for bindROWS,DelayedDataFrame.
check	Logical. This argument is ignored for bindROWS,DelayedDataFrame.
i	row subscript
value	the new values in the i, j subscripts of DelayedDataFrame object.
j	col subscript
drop	if drop with reduced dimension, default is TRUE.

Value

colnames of DelayedDataFrame

DelayedDataFrame

DelayedDataFrame-class

Description

The DelayedDataFrame class extends the DataFrame class and supports the storage of any type of object (with ‘length’ and ‘[’ methods) as columns.

the `lazyIndex` slot getter and setter for DelayedDataFrame object.

the coercion method between DataFrame and DelayedDataFrame objects.

Usage

```
DelayedDataFrame(..., row.names = NULL, check.names = TRUE)

## S4 method for signature 'DelayedDataFrame'
lazyIndex(x)

.from_DataFrame_to_DelayedDataFrame(from)

.from_DelayedDataFrame_to_DFrame(from, to = "DFrame", strict = TRUE)

lazyIndex(x) <- value

## S4 replacement method for signature 'DelayedDataFrame'
lazyIndex(x) <- value
```

Arguments

...	the arguments to pass into construction of a new DelayedDataFrame.
row.names	the rownames for the newly constructed DelayedDataFrame object.
check.names	logical. If 'TRUE' then the names of the variables in the DelayedDataFrame are checked to ensure that they are syntactically valid variable names and are not duplicated. If necessary they are adjusted (by 'make.names') so that they are.
x	the DelayedDataFrame object.
from	the object to be converted.
to	the class of object to be returned by coercion.
strict	Logical. Whether to force return a DataFrame.
value	the new value of lazyIndex slot for DelayedDataFrame object.

Details

The DelayedDataFrame inherits from DataFrame and behaves very similarly in terms of construction, subsetting, splitting, combining, etc. The most notable exception is that The additional slot of `lazyIndex`, enables `DelayedArray` (with different back-ends) columns to share indexes when possible.

Please be very careful to use this replace method for `lazyIndex` slot. Because it only replace the `lazyIndex` slot, but not necessarily the `nrow` and `rownames` slots. If you want to have synchronized subsetting for all slots, the `[` method should be used.

Value

`lazyIndex<-`: the DelayedDataFrame object with new value of `lazyIndex` slot.

Examples

```
DDF <- DelayedDataFrame(letters, LETTERS)
DDF1 <- DDF[1:10,]
DDF1
```

```
lazyIndex(DDF1)
as(DDF1, "DataFrame")
```

LazyIndex-class*The LazyIndex class and methods.***Description**

The `LazyIndex` class is designed to carry mapping indexes for `DelayedDataFrame` columns. So that some operations (e.g., subsetting) on `DelayedDataFrame` are delayed until a realization call is incurred. (e.g., `as.list()`, `DataFrame()`, ...)

`LazyIndex` constructor.

the subsetting method for `LazyIndex` object.

Usage

```
LazyIndex(listData = list(), index = integer())

## S4 method for signature 'LazyIndex'
cbind(..., deparse.level = 1)

## S4 method for signature 'LazyIndex,ANY,ANY,ANY'
x[i, j, ..., drop = TRUE]
```

Arguments

<code>listData</code>	the list data for all mapping indexes that are used in corresponding <code>DelayedDataFrame</code> object.
<code>index</code>	the position of mapping indexes in <code>listData</code> for each column of the corresponding <code>DelayedDataFrame</code> object.
<code>...</code>	<code>LazyIndex</code> objects.
<code>deparse.level</code>	See <code>?base::cbind</code> for a description of this argument.
<code>x</code>	<code>LazyIndex</code> object.
<code>i</code>	row subscript for <code>LazyIndex</code> , which will subset the <code>listData</code> slot.
<code>j</code>	column subscript for <code>LazyIndex</code> , which will subset the <code>index</code> slot.
<code>drop</code>	Logical. Whether to drop the dimension if any of the dimensions has length 1. Default is TRUE.

Details

the `cbind`, `LazyIndex` method is defined to bind the `LazyIndex`es column-wise when `cbind`, `DelayedDataFrame` function is called.

Value

a `LazyIndex` object.

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