

Package ‘Streamer’

October 17, 2024

Type Package

Title Enabling stream processing of large files

Version 1.50.0

Author Martin Morgan, Nishant Gopalakrishnan

Maintainer Martin Morgan <martin.morgan@roswellpark.org>

Description Large data files can be difficult to work with in R, where data generally resides in memory. This package encourages a style of programming where data is 'streamed' from disk into R via a 'producer' and through a series of 'consumers' that, typically reduce the original data to a manageable size. The package provides useful Producer and Consumer stream components for operations such as data input, sampling, indexing, and transformation; see package?Streamer for details.

License Artistic-2.0

LazyLoad yes

Imports methods, graph, RBGL, parallel, BiocGenerics

Suggests RUnit, Rsamtools (>= 1.5.53), GenomicAlignments, Rgraphviz

biocViews Infrastructure, DataImport

Collate AllGenerics.R Streamer-class.R Producer-class.R
Consumer-class.R Stream-class.R ConnectionProducer-classes.R
RawInput-class.R Seq-class.R Downsample-class.R
FunctionProducerConsumer-classes.R ParallelParam-classes.R
Team-class.R Team-methods.R Reducer-class.R DAGParam-class.R
DAGParam-methods.R DAGTeam-class.R Utility-classes.R
lapply-methods.R stream-methods.R plot-methods.R zzz.R

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

git_branch RELEASE_3_19

git_last_commit 680f4d6

git_last_commit_date 2024-04-30

Repository Bioconductor 3.19

Date/Publication 2024-10-16

Contents

Streamer-package	2
ConnectionProducer	3
Consumer	5
DAGTeam	6
Downsample	7
Function*	8
ParallelParam	9
Producer	11
RawInput	12
Reducer	14
reset	15
Seq	16
status	17
Stream	18
Team	20
Utility	21
yield	22
Index	23

Streamer-package	<i>Package to enable stream (iterative) processing of large data</i>
------------------	--

Description

Large data files can be difficult to work with in R, where data generally resides in memory. This package encourages a style of programming where data is 'streamed' from disk into R through a series of components that, typically, reduce the original data to a manageable size. The package provides useful [Producer](#) and [Consumer](#) components for operations such as data input, sampling, indexing, and transformation.

Details

The central paradigm in this package is a [Stream](#) composed of a [Producer](#) and zero or more [Consumer](#) components. The [Producer](#) is responsible for input of data, e.g., from the file system. A [Consumer](#) accepts data from a [Producer](#) and performs transformations on it. The [Stream](#) function is used to assemble a [Producer](#) and zero or more [Consumer](#) components into a single string.

The [yield](#) function can be applied to a stream to generate one 'chunk' of data. The definition of chunk depends on the stream and its components. A common paradigm repeatedly invokes [yield](#) on a stream, retrieving chunks of the stream for further processing.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Producer](#), [Consumer](#) are the main types of stream components. Use [Stream](#) to connect components, and [yield](#) to iterate a stream.

Examples

```
## About this package
packageDescription("Streamer")

## Existing stream components
getClass("Producer") # Producer classes
getClass("Consumer") # Consumer classes

## An example
fl <- system.file("extdata", "s_1_sequence.txt", package="Streamer")
b <- RawInput(fl, 100L, reader=rawReaderFactory(1e4))
s <- Stream(RawToChar(), Rev(), b)
s
head(yield(s)) # First chunk
close(b)

b <- RawInput(fl, 5000L, verbose=TRUE)
d <- Downsample(sampledSize=50)
s <- Stream(RawToChar(), d, b)
s
s[[2]]

## Processing the first ten chunks of the file
i <- 1
while (10 >= i && 0L != length(chunk <- yield(s)))
{
  cat("chunk", i, "length", length(chunk), "\n")
  i <- i + 1
}
close(b)
```

ConnectionProducer *Producer classes to read file connections*

Description

ConnectionProducer classes include ScanProducer, ReadLinesProducer, and ReadTableProducer, providing Streamer interfaces to scan, readLines, and read.table.

Usage

```
ScanProducer(file, ..., fileArgs=list(), scanArgs=list(...))
ReadLinesProducer(con, ..., conArgs=list(), readLinesArgs=list(...))
ReadTableProducer(file, ..., fileArgs=list(), readTableArgs=list(...))
```

```
## S3 method for class 'ConnectionProducer'
close(con, ...)
```

Arguments

file, con The file or connection to be used for input. See [connections](#).
 ... Additional arguments, e.g., nlines, to scan, readLines, etc.
 fileArgs, conArgs Arguments, e.g., mode, encoding, to be used when invoking [reset\(\)](#).
 scanArgs, readLinesArgs, readTableArgs Arguments to scan, readLines, etc., when reading a file or connection; provide this argument when ... contains arguments (especially verbose=TRUE) to be used by the class.

Methods

See [Producer Methods](#).

Internal Class Fields and Methods

Internal fields of this class are described with, e.g., `getRefClass("ReadLinesProducer")$fields`.

Internal methods of this class are described with `getRefClass("ReadLinesProducer")$methods()` and `getRefClass("ReadLinesProducer")$help()`.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Streamer-package](#), [Producer-class](#), [Streamer-class](#).

Examples

```
f1 <- system.file(package="Rsamtools", "extdata", "ex1.sam")

p <- ReadLinesProducer(f1, n = 1000) # read 1000 lines at a time
while (length(y <- yield(p)))
  print(length(y))
close(p)

p <- ReadTableProducer(f1, quote="", fill=TRUE, nrows=1000)
while (length(y <- yield(p)))
  print(dim(y))

reset(p)
dim(yield(p))

## connections opened 'under the hood' are closed, with warnings
```

```
rm(p); gc()

## avoid warnings by managing connections
p <- ScanProducer(file(f1, "r"), verbose=TRUE,
                  scanArgs=list(what=character()))
length(yield(p))
close(p)
rm(p); gc()
```

Consumer

Class defining methods for all consumers

Description

A virtual base class representing components that can consume data from a Producer, and yield data to the user or other Consumer instances. A Consumer typically transforms records from one form to another. Producer and Consumer instances are associated with each other through the [Stream](#) function.

Methods

Methods defined on this class include:

Stream Construct a stream from one Producer and one or more Consumer. See ?Stream.

Internal Class Fields and Methods

Internal fields of this class are described with, e.g., `getRefClass("Consumer")$fields`.

Internal methods of this class are described with `getRefClass("Consumer")$methods()` and `getRefClass("Consumer")$`

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Streamer-package](#), [Streamer-class](#), [Producer-class](#), [Stream-class](#).

Examples

```
showClass("Consumer")
```

DAGTeam

*Consumer classes for directed acyclic graph evaluation***Description**

A [Consumer](#) to route incoming tasks through nodes connected as a directed acyclic graph.

Usage

```
DAGParam(x, ...)
```

```
DAGTeam(..., dagParam = DAGParam(), teamParam = MulticoreParam(1L))
```

```
## S3 method for class 'DAGTeam'
plot(x, y, ...)
```

Arguments

x	A matrix or data.frame with columns 'From', 'To', or a graphNEL object (from the graph package) describing a directed acyclic graph.
...	For DAGTeam, named FunctionConsumer instances, one for each node in the graph. The FunctionConsumer corresponding to the first node in the graph must accept one argument; remaining FunctionConsumer instances must have as input arguments the names of the nodes from which the inputs derive, as in the example below. For DAGParam when x is a data.frame or matrix, data.frame columns W, V or additional arguments W, V as described in ftM2graphNEL .
dagParam	A DAGParam instance, with all nodes referenced in the graph represented by FunctionConsumer instances in ...
teamParam	A ParallelParam instance, such as generated by MulticoreParam(). Currently ignored (all calculations are performed on a single thread).
y	Unused.

Constructors

Use DAGParam and DAGTeam to construct instances of these classes, with ParallelParam instances created by, e.g., MulticoreParam.

Methods

See [Consumer](#) Methods.

Internal Class Fields and Methods

Internal fields of this class are described with, e.g., `getRefClass("MulticoreTeam")$fields`.
Internal methods of this class are described with `getRefClass("MulticoreTeam")$methods()` and `getRefClass("MulticoreTeam")$help()`.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Team](#) applies a single function across multiple threads..

Examples

```
df <- data.frame(From = c("A", "A", "B", "C"),
                 To   = c("B", "C", "D", "D"),
                 stringsAsFactors=FALSE)
dagParam <- DAGParam(df)
dteam <- DAGTeam(A=FunctionConsumer(function(y) y),
                B=FunctionConsumer(function(A) -A),
                C=FunctionConsumer(function(A) 1 / A),
                D=FunctionConsumer(function(B, C) B + C),
                dagParam=dagParam)

plot(dteam)

strm <- Stream(Seq(to=10), dteam)
sapply(strm, c)
reset(strm)
```

Downsample

Consumer class to down-sample data

Description

A [Consumer](#)-class to select records with fixed probability, returning a yield of fixed size. Successive calls to `yield` result in sampling of subsequent records in the stream, until the stream is exhausted.

Usage

```
Downsample(probability=0.1, sampledSize=1e6, ...)
```

Arguments

<code>probability</code>	A <code>numeric(1)</code> between 0, 1 indicating the probability with which a record should be retained.
<code>...</code>	Additional arguments, passed to the <code>\$new</code> method of the underlying reference class. Currently unused.
<code>sampledSize</code>	A <code>integer(1)</code> indicating the number of records to return.

Methods

See [Consumer](#) Methods.

Internal Class Fields and Methods

Internal fields of this class are described with, e.g., `getRefClass("Downsample")$fields`.

Internal methods of this class are described with `getRefClass("Downsample")$methods()` and `getRefClass("Downsample")$help()`.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Stream](#)

Examples

```
showClass("Downsample")
```

Function*

Classes for user-defined Producers and Consumers

Description

The `FunctionProducer` and `FunctionConsumer` classes provide an easy way to quickly create `Producer` and `Consumer` instances from user-provided functions.

Usage

```
FunctionProducer(FUN, RESET, ..., state=NULL)
FunctionConsumer(FUN, RESET, ..., state=NULL)
```

Arguments

<code>FUN</code>	User defined function to yield successive records in the stream. The <code>FunctionProducer</code> function must return an object of length 0 (e.g., <code>logical(0)</code>) when the stream is complete.
<code>RESET</code>	An optional function of one argument ('state') to reset the stream to its original state. If missing, the stream cannot be reset.
<code>...</code>	Arguments passed to the <code>Producer</code> -class or <code>Consumer</code> -class constructors.
<code>state</code>	Any information, made available to <code>RESET</code> .

Constructors

Use `FunctionProducer` or `FunctionConsumer` to construct instances of this class.

Methods

See [Producer](#) and [Consumer](#) Methods.

Internal Class Fields and Methods

Internal fields of this class are described with, e.g., `getRefClass("FunctionProducer")$fields`.

Internal methods of this class are described with `getRefClass("FunctionProducer")$methods()` and `getRefClass("FunctionProducer")$help()`.

Author(s)

Nishant Gopalakrishnan ngopalak@fhcrc.org

See Also

[Stream](#)

Examples

```
## A ProducerFunction
producerFun <- function()
  ## produce the mean of 10 random uniform numbers
  ## stop when the mean is greater than 0.8
  {
    x <- mean(runif(10))
    if (x > .8) numeric(0) else x
  }
randomSampleMeans <- FunctionProducer(producerFun)
result <- sapply(randomSampleMeans, c)
length(result)
head(result)

## A FunctionConsumer:
consumerFun <- function(y)
  ## transform input by -10 log10
  {
    -10 * log10(y)
  }

neg10log10 <- FunctionConsumer(consumerFun)

strm <- Stream(randomSampleMeans, neg10log10)
result <- sapply(strm, c)
length(result)
head(result)
```

Description

Configure and register parallel calculations, e.g., for [Team](#) evaluation.

Usage

```
MulticoreParam(size = getOption("mc.cores", 2L),
  mc.set.seed = TRUE, ...)
register(param)
```

Arguments

size	The number of members in the parallel cluster.
mc.set.seed	logical(1); see ?mcparallel on unix platforms.
param	A ParallelParam instance, such as generated by MulticoreParam().
...	Additional arguments, e.g., verbose, passed to the Streamer class.

Constructors

Use MulticoreParam to construct instances of this class.

Methods

register Invoked with an argument `param` stores the `param` for use in subsequent parallel computation. Use `NULL` to clear the register. The function returns, invisibly, the previously registered parameter instance, if any.

Internal Class Fields and Methods

Internal fields of this class are described with, e.g., `getRefClass("MulticoreParam")$fields`.

Internal methods of this class are described with `getRefClass("MulticoreParam")$methods()` and `getRefClass("MulticoreParam")$help()`.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Team](#) to apply one function in parallel, [DAGTeam](#) to evaluate functions whose dependencies are represented as directed acyclic graphs.

Examples

```
if (.Platform$OS.type != "windows") {
  oparam <- register()      ## previous setting
  param <- MulticoreParam() ## default multicore settings
  register(param)         ## register for future use, e.g., Team
  register(oparam)       ## reset original
}
```

 Producer

Class defining methods for all Producers

Description

A virtual class representing components that can read data from connections, and yield records to the user or a Consumer instance. A Producer represents a source of data, responsible for parsing a file or other data source into records to be passed to Consumer classes. Producer and Consumer instances are associated with each other through the [Stream](#) function.

Usage

```
## S4 method for signature 'Producer'
lapply(X, FUN, ...)

## S4 method for signature 'Producer'
sapply(X, FUN, ..., simplify=TRUE, USE.NAMES=TRUE)
```

Arguments

X	An instance of class Producer
FUN	A function to be applied to each successful yield() of X.
...	Additional arguments to FUN.
simplify	See ?base::sapply.
USE.NAMES	See ?base::sapply but note that names do not usually make sense for instances of class Producer.

Methods

Methods defined on this class include:

Stream Construct a stream from one Producer and one or more Consumer. See ?Stream.

yield Yield a single result (e.g., data.frame) from the Producer.

reset Reset, if possible, the Producer.

lapply, sapply Apply FUN to each result applied to yield(), simplifying (using simplify2array) if possible for sapply. Partial results on error can be recovered using [tryCatch](#), as illustrated below. Infinite producers will of course exhaust memory.

Internal Class Fields and Methods

Internal fields of this class are described with, e.g., `getRefClass("Producer")$fields`.

Internal methods of this class are described with `getRefClass("Producer")$methods()` and `getRefClass("Producer")$`

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Streamer-package](#), [Consumer-class](#), [Streamer-class](#).

Examples

```
showClass("Producer")
showMethods(class="Producer", where="package:Streamer")

sapply(Seq(to=47, yieldSize=7), function(elt) {
  c(n = length(elt), xbar = mean(elt))
})

## recover partial results
fun = function(i) if (i == 5) stop("oops, i == 5") else i
res <- tryCatch(sapply(Seq(to=10), fun), error=function(err) {
  warning(conditionMessage(err),
    "\n only partial results available")
  simplify2array(err$partialResult)
})
res
```

RawInput

Class "RawInput"

Description

A [Producer](#)-class to interpret files as raw (binary) data. Users interact with this class through the constructor [RawInput](#) and methods [yield](#), [reset](#), and [Stream](#).

This class requires two helper functions; the ‘factory’ methods defined on this page can be used to supply these. `rawReaderFactory` creates a ‘reader’, whose responsibility it is to accept a connection and return a vector of predefined type, e.g., `raw`. `rawParserFactory` creates a ‘parser’, responsible for parsing a buffer and vector of the same type as produced by the reader into records. The final record may be incomplete (e.g., because reader does not return complete records), and regardless of completion status is the content of `buf` on the subsequent invocation of `parser`. `length(buf)` or `length(bin)` may be 0, as when the first or final record is parsed.

Usage

```
RawInput(con, yieldSize = 1e+06, reader = rawReaderFactory(),
  parser = rawParserFactory(), ...)
rawReaderFactory(blockSize = 1e+06, what)
rawParserFactory(separator = charToRaw("\n"), trim = separator)
```

Arguments

<code>con</code>	A character string or connection (opened as "rb" mode) from which raw input will be retrieved.
<code>yieldSize</code>	The number of records the input parser is to yield.

reader	A function of one argument (con, an open connection positioned at the start of the file, or at the position the con was in at the end of the previous invocation of the reader function) that returns a vector of type raw.
parser	A function of two arguments (buf, bin), parsing the raw vector c(buf, bin) into records.
...	Additional arguments, passed to the \$new method of this class. Currently ignored.
blockSize	The number of bytes to read at one time.
what	The type of data to read, as the argument to readBin .
separator	A raw vector indicating the unique sequence of bytes by which record starts are to be recognized. The parser supplied here includes the record separator at the start of each record.
trim	A raw vector that is a prefix of separator, and that is to be removed from the record during parsing.

Fields

- con: Object of class connection. An R [connection](#) opened in “rb” mode from which data will be read.
- blockSize: Object of class integer. Size (e.g., number of raw bytes) input during each [yield](#).
- reader: Object of class function. A function used to input blockSize elements. See [rawReaderFactory](#).
- parser: Object of class function. A function used to parse raw input into records, e.g., breaking a raw vector on new lines ‘\n’. See [rawParserFactory](#)
- .buffer: Object of class raw. Contains read but not parsed raw stream data.
- .records: Object of class list. Parsed but not yet yield-ed records.
- .parsedRecords: Object of class integer. Total number of records parsed by the Producer.

Class-Based Methods

- reset(): Remove buffer and current records, reset record counter, re-open con.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Stream](#)

Examples

```
f1 <- system.file("extdata", "s_1_sequence.txt", package="Streamer")
b <- RawInput(f1, 100L, reader=rawReaderFactory(1e4))
length(value <- yield(b))
head(value)
close(b)
```

 Reducer

Consumer class to combine successive records

Description

A [Consumer](#)-class to reduce N successive records into a single yield.

Usage

```
Reducer(FUN, init, ..., yieldNth = NA_integer_)
```

Arguments

<code>FUN</code>	A function of two arguments, where the first argument is the result of the previous reduction (or <code>init</code> , if specified, for the first record) and the second argument is the current record.
<code>init</code>	An optional initial value to initiate the reduction. When present, <code>init</code> is used to initial each yield.
<code>...</code>	Additional arguments, passed to the <code>\$new</code> method of the underlying reference class. Currently unused.
<code>yieldNth</code>	A positive integer indicating how many upstream yields are combined before the Reducer yields. A value of <code>NA_integer_</code> indicates reduction of all records in the input stream.

Methods

See [Consumer](#) Methods.

Internal Class Fields and Methods

Internal fields of this class are described with, e.g., `getRefClass("Reducer")$fields`.

Internal methods of this class are described with `getRefClass("Reducer")$methods()` and `getRefClass("Reducer")$he`.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Stream](#)

Examples

```

s <- Stream(Seq(to=10), Reducer("+"))
yield(s)    ## sum(1:10), i.e., Reduce over the entire stream
s <- Stream(Seq(to=10), Reducer("+", yieldNth=5))
yield(s)    ## sum(1:5)
yield(s)    ## sum(6:10)
s <- Stream(Seq(to=10), Reducer("+", init=10, yieldNth=5))
sapply(s, c) ## 10 + c(sum(1:5), sum(6:10))
if (.Platform$OS.type != "windows") {
  s <- Stream(Seq(to=10),
              Team(function(i) { Sys.sleep(1); i },
                    param=MulticoreParam(10L)),
              Reducer("+"))
  system.time(y <- yield(s))
  y
}

```

reset

*Function to reset a Stream, Producer, or Consumer***Description**

reset on a stream invokes the reset method of all components of the stream; on a component, it invokes the reset method of the component and all inputs to the component.

Usage

```
reset(x, ...)
```

Arguments

x A Stream, Producer, or Consumer object.
... Additional arguments, currently unused.

Value

A reference to x, the stream or component on which reset was invoked.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Stream](#), [Producer](#), [Consumer](#).

Examples

```
## see example(Stream)
```

Seq

*Producer class to generate (numeric) sequences***Description**

A [Producer](#)-class to generate a sequence (possibly long) of numbers.

Usage

```
Seq(from = 1L, to=.Machine$integer.max, by = 1L, yieldSize=1L,
    ...)
```

Arguments

from	A starting value of any type (e.g., integer, numeric supported by <code>base::seq</code>).
to	An ending value, typically of the same type as <code>from</code> .
by	A value, typically of the same class as <code>from</code> , indicating the increment between successive numbers in the sequence. <code>by = 0</code> can create an infinite stream.
yieldSize	A <code>integer(1)</code> indicating the length of the output sequence each time <code>yield()</code> is invoked.
...	Additional arguments passed to Producer .

Constructors

Use `Seq` to construct instances of this class.

Methods

See [Producer](#) Methods.

Internal Class Fields and Methods

Internal fields of this class are described with `getRefClass("Seq")$fields`.

Internal methods of this class are described with `getRefClass("Seq")$methods()` and `getRefClass("Seq")$help()`.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Stream](#)

Examples

```
s <- Seq(1, 10, yieldSize=5)
while(length(y <- yield(s)))
  print(y)
```

status	<i>Function to report current status of a stream</i>
--------	--

Description

status invoked on a stream yields the current status of the stream, as reported by the status methods of each component.

Usage

```
status(x, ...)
```

```
## S4 method for signature 'Streamer'  
status(x, ...)
```

Arguments

x	A Stream, Producer, or Consumer object.
...	Additional arguments, currently unused.

Value

A component-specific summary the current status

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Stream](#), [Producer](#), [Consumer](#).

Examples

```
## see example(Stream)
```

Stream

Class to represent a Producer and zero or more Consumers

Description

An ordered collection of Consumer and Producer components combined into a single entity. Applying a method such as `yield` to `Stream` invokes `yield` on the terminal Consumer component of the stream, yielding one batch from the stream. The result of `yield` is defined by the Producer and Consumer components of the stream.

Usage

```
Stream(x, ..., verbose=FALSE)

## S4 method for signature 'Stream'
length(x)

## S4 method for signature 'Stream,numeric'
x[[i, j, ...]]

## S4 method for signature 'Stream'
lapply(X, FUN, ...)

## S4 method for signature 'Stream'
sapply(X, FUN, ..., simplify=TRUE, USE.NAMES=TRUE)
```

Arguments

<code>x, X</code>	For <code>Stream</code> , <code>x</code> is a Producer instance. For other functions, an instance of class <code>Stream</code> .
<code>FUN</code>	A function to be applied to each successful <code>yield()</code> of <code>X</code> .
<code>i, j</code>	Numeric index of the <code>i</code> th stream element (<code>j</code> is ignored by this method).
<code>...</code>	For <code>Stream</code> , zero or more Consumer instances. For <code>lapply</code> , <code>sapply</code> , additional arguments to <code>FUN</code> .
<code>simplify</code>	See <code>?base::sapply</code> .
<code>USE.NAMES</code>	See <code>?base::sapply</code> but note that names do not usually make sense for instances of class <code>Producer</code> .
<code>verbose</code>	A <code>logical(1)</code> indicating whether status information should be reported.

Constructors

Arguments to `Stream` must consist of a single Producer and zero or more Consumer components.

When invoked with the Producer as the first argument, `Stream(P, C1, C2)` produces a stream in which the data is read by `P`, then processed by `C1`, then processed by `C2`.

When invoked with the Consumer as the first argument, the . . . must include a Producer as the *last* argument. `Stream(C1, C2, P)` produces a stream in which the data is read by P, then processed by C2, then processed by C1.

Methods

Methods defined on this class include:

length The number of components in this stream.

[[The *i*th component (including inputs) of this stream.

yield Yield a single result (e.g., `data.frame`) from the stream.

reset Reset, if possible, each component of the stream.

lapply, sapply Apply FUN to each result applied to `yield()`, simplifying (using `simplify2array`) if possible for `sapply`. Partial results on error can be recovered using `tryCatch`, as illustrated on the help page [Producer](#).

Internal Class Fields and Methods

Internal fields of this class are described with, e.g., `getRefClass("FunctionProducer")$fields`.

Internal methods of this class are described with `getRefClass("FunctionProducer")$methods()` and `getRefClass("FunctionProducer")$help()`.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Streamer-package](#), [Consumer-class](#), [Producer-class](#).

Examples

```
f1 <- system.file("extdata", "s_1_sequence.txt", package="Streamer")
b <- RawInput(f1, 100L, reader=rawReaderFactory(1e4))
s <- Stream(b, Rev(), RawToChar())
s
yield(s)
reset(s)
while (length(yield(s))) cat("tick\n")
close(b)

strm <- Stream(Seq(to=10), FunctionConsumer(function(y) 1/y))
sapply(strm, c)
```

Team

Consumer classes for parallel evaluation

Description

A [Consumer](#) to divide incoming tasks amongst processes for parallel evaluation; not supported on Windows.

Usage

```
Team(FUN, ..., param)
```

Arguments

<code>FUN</code>	A function of one argument (the input to this consumer), to be applied to each element of the stream. The return value of the function is the value yield'ed.
<code>...</code>	Additional arguments (e.g., <code>verbose</code> , passed to the Consumer constructor.
<code>param</code>	If provided, a <code>ParallelParam</code> instance, such as generated by <code>MulticoreParam()</code> .

Constructors

Use `Team` to construct instances of this class.

When `param` is missing, `Team` consults the registry (see [register](#)) for a parallel parameter class. If none is found and `.Platform$OS.type == "unix"`, a default `MulticoreParam` instance is used. An error is signaled on other operating systems (i.e., Windows)

Methods

See [Consumer](#) Methods.

Internal Class Fields and Methods

Internal fields of this class are described with, e.g., `getRefClass("MulticoreTeam")$fields`.

Internal methods of this class are described with `getRefClass("MulticoreTeam")$methods()` and `getRefClass("MulticoreTeam")$help()`.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[ParallelParam](#) for configuring parallel environments. [DAGTeam](#) apply functions organized as a directed acyclic graph.

Examples

```
if (.Platform$OS.type != "windows") {
  param <- MulticoreParam(size=5)
  team <- Team(function(x) { Sys.sleep(1); mean(x) }, param=param)
  s <- Stream(Seq(to=50, yieldSize=5), team)
  system.time({while(length(y <- yield(s)))
    print(y)
  }) ## about 2 seconds
}
```

Utility

Consumer classes with simple functionality, e.g., RawToChar, Rev

Description

Utility is a virtual class containing components to create light weight Consumer classes.
RawToChar is a class to convert raw (binary) records to char, applying rawToChar to each record.
Rev reverses the order of current task.

Usage

```
RawToChar(...)  
Rev(...)
```

Arguments

... Arguments passed to the [Consumer](#)-class.

Construction

Use constructors [RawToChar](#), [Rev](#).

Methods

See [Consumer](#) Methods.

Internal Class Fields and Methods

Internal fields of this class are described with, e.g., `getRefClass("Utility")$fields`.

Internal methods of this class are described with `getRefClass("Utility")$methods()` and `getRefClass("Utility")$he`.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Streamer-package](#), [Consumer-class](#), [Streamer-class](#).

Examples

```
showClass("Utility")
```

yield

Function to yield one task from a Stream or Producer

Description

yield invoked on a stream yields one chunk of data or, if the stream is complete, a length zero element of the data. Successive invocations of yield produce successive chunks of data.

Usage

```
yield(x, ...)
```

Arguments

x	A Stream, Producer, or Consumer object.
...	Additional arguments, currently unused.

Value

A chunk of data, with the specific notion of chunk defined by the final component of the stream.

Author(s)

Martin Morgan mtmorgan@fhcrc.org

See Also

[Stream](#), [Producer](#), [Consumer](#).

Examples

```
## see example(Stream)
```

Index

* classes

- ConnectionProducer, 3
- DAGTeam, 6
- Downsample, 7
- Function*, 8
- ParallelParam, 9
- Producer, 11
- RawInput, 12
- Reducer, 14
- Seq, 16
- Stream, 18
- Team, 20
- Utility, 21

* manip

- status, 17
- yield, 22

* methods

- reset, 15

* package

- Streamer-package, 2

[[, Stream, numeric-method (Stream), 18

close.ConnectionProducer
(ConnectionProducer), 3

connection, 13

ConnectionProducer, 3

ConnectionProducer-class
(ConnectionProducer), 3

ConnectionProducer-classes
(ConnectionProducer), 3

connections, 4

Consumer, 2, 3, 5, 6–8, 12, 14, 15, 17, 19–22

Consumer-class (Consumer), 5

DAGParam (DAGTeam), 6

DAGParam, data.frame-method (DAGTeam), 6

DAGParam, graphNEL-method (DAGTeam), 6

DAGParam, matrix-method (DAGTeam), 6

DAGParam, missing-method (DAGTeam), 6

DAGParam-class (DAGTeam), 6

DAGTeam, 6, 10, 20

DAGTeam-class (DAGTeam), 6

Downsample, 7

Downsample-class (Downsample), 7

ftM2graphNEL, 6

Function*, 8

FunctionConsumer, 6

FunctionConsumer (Function*), 8

FunctionConsumer-class (Function*), 8

FunctionProducer (Function*), 8

FunctionProducer-class (Function*), 8

FunctionProducerConsumer-classes
(Function*), 8

lapply, Producer-method (Producer), 11

lapply, Stream-method (Stream), 18

length, Stream-method (Stream), 18

MulticoreParam, 20

MulticoreParam (ParallelParam), 9

MulticoreParam-class (ParallelParam), 9

MulticoreTeam-class (Team), 20

ParallelParam, 9, 20

ParallelParam-class (ParallelParam), 9

ParallelRegister-class (ParallelParam), 9

plot.DAGParam (DAGTeam), 6

plot.DAGTeam (DAGTeam), 6

Producer, 2–5, 8, 11, 12, 15–17, 19, 22

Producer-class (Producer), 11

RawInput, 12, 12

RawInput-class (RawInput), 12

rawParserFactory, 13

rawParserFactory (RawInput), 12

rawReaderFactory, 13

rawReaderFactory (RawInput), 12

RawToChar (Utility), 21

RawToChar-class (Utility), 21

- readBin, [13](#)
- ReadLinesProducer (ConnectionProducer), [3](#)
- ReadLinesProducer-class (ConnectionProducer), [3](#)
- ReadTableProducer (ConnectionProducer), [3](#)
- ReadTableProducer-class (ConnectionProducer), [3](#)
- Reducer, [14](#)
- Reducer-class (Reducer), [14](#)
- register, [20](#)
- register (ParallelParam), [9](#)
- reset, [4](#), [12](#), [15](#)
- reset, Streamer-method (reset), [15](#)
- reset-methods (reset), [15](#)
- Rev (Utility), [21](#)
- Rev-class (Utility), [21](#)

- sapply, Producer-method (Producer), [11](#)
- sapply, Stream-method (Stream), [18](#)
- ScanProducer (ConnectionProducer), [3](#)
- ScanProducer-class (ConnectionProducer), [3](#)
- Seq, [16](#)
- Seq-class (Seq), [16](#)
- show, Consumer-method (Consumer), [5](#)
- status, [17](#)
- status, Streamer-method (status), [17](#)
- status-methods (status), [17](#)
- Stream, [2](#), [3](#), [5](#), [8](#), [9](#), [11–17](#), [18](#), [22](#)
- Stream, Consumer-method (Stream), [18](#)
- Stream, Producer-method (Stream), [18](#)
- Stream-class (Stream), [18](#)
- Stream-methods (Stream), [18](#)
- Streamer, [4](#), [5](#), [12](#), [21](#)
- Streamer (Streamer-package), [2](#)
- Streamer-class (Streamer-package), [2](#)
- Streamer-package, [2](#)

- Team, [7](#), [9](#), [10](#), [20](#)
- Team, missing-method (Team), [20](#)
- Team, MulticoreParam-method (Team), [20](#)
- Team-class (Team), [20](#)
- tryCatch, [11](#), [19](#)

- Utility, [21](#)
- Utility-class (Utility), [21](#)
- Utility-classes (Utility), [21](#)

- yield, [2](#), [3](#), [12](#), [13](#), [22](#)
- yield, Streamer-method (yield), [22](#)
- yield-methods (yield), [22](#)