## Package 'Streamer'

December 20, 2024

Type Package

**Title** Enabling stream processing of large files

Version 1.52.0

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\_20

git\_last\_commit ac06971

git\_last\_commit\_date 2024-10-29

Repository Bioconductor 3.20

**Date/Publication** 2024-12-19

2 Streamer-package

## **Contents**

	Streamer-package	1
	ConnectionProducer	1
	Consumer	
	DAGTeam	
	Downsample	1
	Function*	8
	ParallelParam	9
	Producer	1(
	RawInput	12
	Reducer	13
	reset	14
	Seq	1.
	status	1
	Stream	ľ
	Team	18
	Utility	2(
	yield	2
Index		2
Stre	mer-package Package to enable stream (iterative) processing of large data	

## **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.

ConnectionProducer 3

#### **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")</pre>
b <- RawInput(fl, 100L, reader=rawReaderFactory(1e4))</pre>
s <- Stream(RawToChar(), Rev(), b)</pre>
                 # First chunk
head(yield(s))
close(b)
b <- RawInput(fl, 5000L, verbose=TRUE)</pre>
d <- Downsample(sampledSize=50)</pre>
s <- Stream(RawToChar(), d, b)</pre>
s
s[[2]]
## Processing the first ten chunks of the file
i <- 1
while (10 \geq i && 0L != length(chunk \leq 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.
```

4 ConnectionProducer

```
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 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.

```
fl <- system.file(package="Rsamtools", "extdata", "ex1.sam")</pre>
p <- ReadLinesProducer(fl, n = 1000) # read 1000 lines at a time
while (length(y <- yield(p)))</pre>
    print(length(y))
close(p)
p <- ReadTableProducer(fl, quote="", fill=TRUE, nrows=1000)</pre>
while (length(y <- yield(p)))</pre>
    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 <- \mbox{ScanProducer(file(fl, "r"), verbose=TRUE,} \\
                   scanArgs=list(what=character()))
length(yield(p))
close(p)
rm(p); gc()
```

Consumer 5

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 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, ...)
```

6 DAGTeam

## 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).
у	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..

Downsample 7

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

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**

probability A numeric(1) between 0, 1 indicating the probability with which a record should be retained.

... Additional arguments, passed to the \$new method of the underlying reference class. Currently unused.

sampledSize A integer(1) 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

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

8 Function\*

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**

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

#### **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 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

ParallelParam 9

#### **Examples**

```
## A ProducerFunction
producerFun <- function()</pre>
    ## produce the mean of 10 random uniform numbers
    ## stop when the mean is greater than 0.8
{
    x <- mean(runif(10))</pre>
    if (x > .8) numeric(0) else x
randomSampleMeans <- FunctionProducer(producerFun)</pre>
result <- sapply(randomSampleMeans, c)</pre>
length(result)
head(result)
## A FunctionConsumer:
consumerFun <- function(y)</pre>
    ## transform input by -10 log10
{
    -10 * log10(y)
}
neg10log10 <- FunctionConsumer(consumerFun)</pre>
strm <- Stream(randomSampleMeans, neg10log10)</pre>
result <- sapply(strm, c)</pre>
length(result)
head(result)
```

ParallelParam

Classes to configure parallel evaluation

#### **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.

10 Producer

#### 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**

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

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
}</pre>
```

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

```
## $4 method for signature 'Producer'
lapply(X, FUN, ...)
## $4 method for signature 'Producer'
sapply(X, FUN, ..., simplify=TRUE, USE.NAMES=TRUE)
```

Producer 11

#### **Arguments**

Χ	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 are described with, e.g., getRefClass("Producer")\$fields.

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

#### Author(s)

Martin Morgan mtmorgan@fhcrc.org

#### See Also

```
Streamer-package, Consumer-class, Streamer-class.
```

12 RawInput

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

con	A character string or connection (opened as "rb" mode) from which raw input will be retrieved.
yieldSize	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.

Reducer 13

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**

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

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**

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

14 reset

## Methods

See Consumer Methods.

#### **Internal Class Fields and Methods**

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

Internal methods of this class are described with getRefClass("Reducer") methods() and getRefClass("Reducer")

#### Author(s)

Martin Morgan mtmorgan@fhcrc.org

#### See Also

Stream

## **Examples**

```
s <- Stream(Seq(to=10), Reducer("+"))</pre>
             ## sum(1:10), i.e., Reduce over the entire stream
s <- Stream(Seq(to=10), Reducer("+", yieldNth=5))</pre>
             ## sum(1:5)
yield(s)
yield(s)
             ## sum(6:10)
s <- Stream(Seq(to=10), Reducer("+", init=10, yieldNth=5))</pre>
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))</pre>
}
```

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.

Seq 15

#### 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

## Arguments

from	A starting value of any type (e.g., integer, numeric supported by base::seq.
to	An ending value, typically of the same type as from.
by	A value, typically of the same class as from, indicating the increment between successive numbers in the sequence. by $= 0$ can create an infinite stream.
yieldSize	A integer(1) indicating the length of the output sequence each time yield() 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 are described with getRefClass("Seq")\$fields.

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

16 status

#### Author(s)

Martin Morgan mtmorgan@fhcrc.org

#### See Also

Stream

## **Examples**

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

status

Function to report current status of a stream

## 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.
```

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

Stream 17

#### **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**

x, X	For Stream, $\boldsymbol{x}$ is a Producer instance. For other functions, an instance of class Stream.
FUN	A function to be applied to each successful yield() of X.
i, j	Numeric index of the ith stream element (j is ignored by this method).
	For Stream, zero or more Consumer instances. For lapply, sapply, 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.
verbose	A logical(1) 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.

Team Team

#### Methods

Methods defined on this class include:

**length** The number of components in this stream.

[[ The ith 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 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**

```
fl <- system.file("extdata", "s_1_sequence.txt", package="Streamer")
b <- RawInput(fl, 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)</pre>
```

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)
```

Team 19

## **Arguments**

FUN	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.
	Additional arguments (e.g., verbose, passed to the Consumer constructor.
param	If provided, a ParallelParam instance, such as generated by MulticoreParam().

## 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 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.

```
if (.Platform$0S.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
}</pre>
```

20 Utility

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 are described with, e.g., getRefClass("Utility")\$fields.

Internal methods of this class are described with getRefClass("Utility")\$methods() and getRefClass("Utility")

## Author(s)

Martin Morgan mtmorgan@fhcrc.org

#### See Also

```
Streamer-package, Consumer-class, Streamer-class.
```

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

yield 21

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.
```

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

# Index

* classes	ftM2graphNEL, 6
ConnectionProducer, 3	Function*, 8
DAGTeam, 5	FunctionConsumer, $6$
Downsample, 7	<pre>FunctionConsumer (Function*), 8</pre>
Function*, 8	<pre>FunctionConsumer-class(Function*), 8</pre>
ParallelParam, 9	FunctionProducer (Function*), 8
Producer, 10	<pre>FunctionProducer-class(Function*), 8</pre>
RawInput, 12	FunctionProducerConsumer-classes
Reducer, 13	(Function*), 8
Seq, 15	
Stream, 17	lapply, Producer-method (Producer), 10
Team, 18	lapply, Stream-method (Stream), 17
Utility, 20	length, Stream-method (Stream), 17
* manip	MulticoreParam, 19
status, 16	MulticoreParam (ParallelParam), 9
yield, 21	MulticoreParam-class (ParallelParam), 9
* methods	MulticoreTeam-class (Team), 18
reset, 14	riartitor cream crass (ream), 10
* package	ParallelParam, 9, 19
Streamer-package, 2	ParallelParam-class (ParallelParam), 9
[[,Stream,numeric-method(Stream), 17	ParallelRegister-class (ParallelParam) 9
close.ConnectionProducer	plot.DAGParam (DAGTeam), 5
(ConnectionProducer), 3	plot.DAGTeam (DAGTeam), 5
connection, 12	Producer, 2, 4, 5, 8, 10, 12, 15, 16, 18, 21
ConnectionProducer, 3	Producer-class (Producer), 10
ConnectionProducer-class	
(ConnectionProducer), 3	RawInput, <i>12</i> , 12
ConnectionProducer-classes	RawInput-class (RawInput), 12
(ConnectionProducer), 3	rawParserFactory, <i>13</i>
connections, $3$	rawParserFactory (RawInput), 12
Consumer, 2, 5, 5, 6–8, 11, 13–16, 18–21	rawReaderFactory, <i>12</i>
Consumer-class (Consumer), 5	rawReaderFactory (RawInput), 12
	RawToChar (Utility), $20$
DAGParam (DAGTeam), 5	RawToChar-class(Utility), 20
DAGParam, data.frame-method (DAGTeam), 5	readBin, <i>12</i>
DAGParam, graphNEL-method (DAGTeam), 5	ReadLinesProducer (ConnectionProducer)
DAGParam, matrix-method (DAGTeam), 5	3
DAGParam, missing-method (DAGTeam), 5	ReadLinesProducer-class
DAGParam-class (DAGTeam), 5	(ConnectionProducer), 3
DAGTeam, 5, 10, 19	${\sf ReadTableProducer}({\sf ConnectionProducer})$
DAGTeam-class (DAGTeam), 5	3
Downsample, 7	ReadTableProducer-class
Downsample-class (Downsample), 7	(ConnectionProducer), 3

INDEX 23

```
Reducer, 13
Reducer-class (Reducer), 13
register, 19
register (ParallelParam), 9
reset, 4, 12, 14
reset, Streamer-method (reset), 14
reset-methods (reset), 14
Rev (Utility), 20
Rev-class (Utility), 20
sapply, Producer-method (Producer), 10
sapply, Stream-method (Stream), 17
ScanProducer (ConnectionProducer), 3
ScanProducer-class
         (ConnectionProducer), 3
Seq, 15
Seq-class (Seq), 15
show, Consumer-method (Consumer), 5
status, 16
status, Streamer-method (status), 16
status-methods (status), 16
Stream, 2, 5, 7, 8, 10, 12–16, 17, 21
Stream, Consumer-method (Stream), 17
Stream, Producer-method (Stream), 17
Stream-class (Stream), 17
Stream-methods (Stream), 17
Streamer, 4, 5, 11, 20
Streamer (Streamer-package), 2
Streamer-class (Streamer-package), 2
Streamer-package, 2
Team, 6, 9, 10, 18
Team, missing-method (Team), 18
Team, MulticoreParam-method (Team), 18
Team-class (Team), 18
tryCatch, 11, 18
Utility, 20
Utility-class (Utility), 20
Utility-classes (Utility), 20
yield, 2, 12, 21
yield, Streamer-method (yield), 21
\verb|yield-methods| (\verb|yield|), 21
```