Package 'Streamer'

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Type Package

Title Enabling stream processing of large files

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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.

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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
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DAGParam-methods.R DAGTeam-class.R Utility-classes.R
lapply-methods.R stream-methods.R plot-methods.R zzz.R

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Contents

Streamer-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.

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 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

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

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

p <- ReadTableProducer(fl, 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(fl, "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 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 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

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 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

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](#)

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)
```

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.

Methods

> **register** Invoked with an argument `param` stores the `param` for use in subsequent parallel computa-
> tion. 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 rep-
resented 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 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

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](#).

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)
```

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.

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("+"))
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 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().

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	*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.

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

x, X	For `Stream`, 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`.

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)
```

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

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.

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

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

Author(s)

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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.

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See Also

Stream, Producer, Consumer.

Examples

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

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