Package 'RedisParam'

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Title Provide a 'redis' back-end for BiocParallel

Version 1.14.0

Description This package provides a Redis-based back-end for BiocParallel, enabling an alternative mechanism for distributed computation. The The 'manager' distributes tasks to a 'worker' pool through a central Redis server, rather than directly to workers as with other BiocParallel implementations. This means that the worker pool can change dynamically during job evaluation. All features of BiocParallel are supported, including reproducible random number streams, logging to the manager, and alternative 'load balancing' task distributions.

Depends R (>= 4.2.0), BiocParallel (>= 1.29.12)

SystemRequirements hiredis

Imports methods, redux, withr, logger

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`bpstopall` *Deprecated functions in the RedisParam package*

Description

`bpstopall()` is provided for compatibility with previous versions of RedisParam, and will be defunct after the next release. Use `rpstopall()` instead.

Usage

```
bpstopall(x)
```

Arguments

`x`	a `RedisParam` object.

Value

See `?rpstopall` for return value.

Examples

```
if (FALSE) {
    ## bpstopall()
    ## deprecated -- use rpstopall() instead
}
```

RedisBackend *Creating the Redis backend*

Description

Creating the Redis backend

Usage

```
RedisBackend(
  RedisParam = NULL,
  jobname = "myjob",
  host = rphost(),
  port = rpport(),
  password = rppassword(),
  timeout = .Machine$integer.max,
  type = c("manager", "worker"),
  id = NULL,
  log = FALSE,
  redis.log = NULL,
  flushInterval = 5L
)

## S4 method for signature 'RedisBackend'
.recv(worker)

## S4 method for signature 'RedisBackend'
.send(worker, value)

## S4 method for signature 'RedisBackend'
.close(worker)

## S4 method for signature 'RedisBackend'
.send_to(backend, node, value)

## S4 method for signature 'RedisBackend'
.recv_any(backend)

## S4 method for signature 'RedisBackend'
.recv_all(backend)

## S4 method for signature 'RedisBackend'
bpjobname(x)

## S4 method for signature 'RedisBackend'
bpworkers(x)
```

Arguments

RedisParam	RedisParam, if this argument is not NULL, all the other arguments will be ignored except type.
jobname	character(1) The job name used by the manager and workers to connect.
host	character(1) The host of the Redis server.
port	integer(1) The port of the Redis server.
password	character(1) The password of the redis server.
timeout	integer(1) The waiting time in BLPOP.
type	character(1) The type of the backend (manager or worker?).
id	character(1) The manager/worker ID. If not given by the user and the environment REDISPARAM_ID is not defined, a random ID will be used
log	logical(1) Whether to enable the log
redis.log	logical(1) Whether to enable the redis server log
flushInterval	numeric(1) The waiting time between two flush operation.

Value

RedisBackend() returns an object of class RedisBackend. This object is not useful to the end user.

RedisParam *Enable redis-based parallel evaluation in BiocParallel*

Description

RedisParam() creates an object describing manager and worker configurations for parallel compuation using a Redis server back-end.

rpalive() tests whether it is possible to connect to a redis server using the host, port, and password in the RedisParam object.

rpstopall() is used from the manager to stop redis workers launched independently, with is.worker=TRUE.

rpworkers() determines the number of workers using snowWorkers() if workers are created dynamically, or a fixed maximum (currently 1000) if workers are listening on a queue.

rphost() reads the host name of the Redis server from the system environment variable REDISPARAM_HOST, if the variable is not defined, fallback to REDIS_HOST. Otherwise default to "127.0.0.1". rphost(x) gives the host name used by x.

rpport() reads the port of the Redis server from a system environment variable REDISPARAM_PORT, if the variable is not defined, fallback to REDIS_PORT. Otherwise default to 6379. rpport(x) gives the port used by x.

rppassword() reads an (optional) password from the system environment variable REDISPARAM_PASSWORD, if the variable is not defined, fallback to REDIS_PASSWORD. Otherwise default to NA_character_ (no password). rppassword(x) gives the password used by x.

Usage

```
RedisParam(
  workers = rpworkers(is.worker),
  tasks = 0L,
  jobname = ipcid(),
  log = FALSE,
  logdir = NA,
  threshold = "INFO",
  resultdir = NA_character_,
  stop.on.error = TRUE,
  timeout = NA_integer_,
  exportglobals = TRUE,
  progressbar = FALSE,
  RNGseed = NULL,
  queue.multiplier = 2L,
  redis.hostname = rphost(),
  redis.port = rpport(),
  redis.password = rppassword(),
  is.worker = NA
)

rpalive(x)

rpstopall(x)

rpworkers(is.worker)

rphost(x)

rpport(x)

rppassword(x)

rpisworker(x)

## S4 method for signature 'RedisParam'
bpisup(x)

## S4 method for signature 'RedisParam'
bpbackend(x)

## S4 method for signature 'RedisParam'
bpstart(x, ...)

## S4 method for signature 'RedisParam'
bpstop(x)

## S4 method for signature 'RedisParam'
```

```
bpworkers(x)

## S4 replacement method for signature 'RedisParam,logical'
bplog(x) <- value
```

Arguments

workers	integer(1) number of redis workers. For is.worker=FALSE, this parameter is the maximum number of workers expected to be available. For is.worker=NA, this is the number of workers opened by bpstart().
tasks	See ?"BiocParallelParam-class".
jobname	character(1) name (unique) used to associate manager & workers on a queue.
log	See ?"BiocParallelParam-class".
logdir	See ?"BiocParallelParam-class".
threshold	See ?"BiocParallelParam-class".
resultdir	See ?"BiocParallelParam-class".
stop.on.error	See ?"BiocParallelParam-class".
timeout	See ?"BiocParallelParam-class".
exportglobals	See ?"BiocParallelParam-class".
progressbar	See ?"BiocParallelParam-class".
RNGseed	See ?"BiocParallelParam-class".
queue.multiplier	numeric(1), The multiplier of the queue depth. The depth of the queue is calculated by queue.multiplier * bpnworkers(p). A proper queue depth can provide more performance benefit in task dispatching, but the improvement is likely to be marginal for an excessively large queue.multiplier.
redis.hostname	character(1) host name of redis server, from system environment variable REDISPARAM_HOST or REDIS_HOST, if both are not defined, the default "127.0.0.1" is used.
redis.port	integer(1) port of redis server, from system environment variable REDISPARAM_PORT or REDIS_PORT, if both are not defined, the default 6379 is used.
redis.password	character(1) or NULL, host password of redis server from system environment variable REDISPARAM_PASSWORD or REDIS_PASSWORD, if both are not defined, the default NA_character_ (no password) is used.
is.worker	logical(1) bpstart() creates worker-only (TRUE), manager-only (FALSE), or manager and worker (NA, default) connections.
x	A RedisParam object.
...	ignored.
value	The value you want to replace with

Details

Use an instance of RedisParam() for interactive parallel evaluation using bplapply() or bpiterate(). RedisParam() requires access to a redis server, running on manager.hostname (e.g., 127.0.0.1) at manager.port (e.g., 6379). The manager and workers communicate via the redis server, rather than the socket connections used by other BiocParallel back-ends.

When invoked with is.worker = NA (the default) bpstart(), bplapply() and bpiterate() start and stop redis workers on the local computer. It may be convenient to use bpstart() and bpstop() independently, to amortize the cost of worker start-up across multiple calls to bplapply() / bpiterate().

Alternatively, a manager and one or more workers can each be started in different processes across a network. The manager is started, e.g., in an interactive session, by specifying is.worker=FALSE. Workers are started, typically as background processes, with is.worker = TRUE. Both manager and workers must specify the same value for jobname =, the redis key used for communication. In this scenario, workers can be added at any time, including during e.g., bplapply() evaluation on the manager. See the vignette for possible scenarios.

Value

RedisParam() returns an object of class RedisParam, for use in controlling parallel evaluation with BiocParallel::bplapply() or BiocParallel::bpiterate().

Examples

```
param <- RedisParam()
if (rpalive(param)) {
    res <- bplapply(1:20, function(i) Sys.getpid(), BPPARAM = param)
    table(unlist(res))
}

## Not run:
## start workers in background proocess(es)
rscript <- R.home("bin/Rscript")
worker_script <- tempfile()
writeLines(c(
    'worker <- RedisParam::RedisParam(jobname = "demo", is.worker = TRUE)',
    'RedisParam::bpstart(worker)'
), worker_script)

for (i in seq_len(2))
    system2(rscript, worker_script, wait = FALSE)

## start manager
p <- RedisParam(jobname = "demo", is.worker = FALSE)
result <- bplapply(1:5, function(i) Sys.getpid(), BPPARAM = p)
table(unlist(result))

## stop all workers
rpstopall(p)

## End(Not run)
```

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