

Package ‘gQTLBase’

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Title gQTLBase: infrastructure for eQTL, mQTL and similar studies

Version 1.0.0

Author VJ Carey <stvjc@channing.harvard.edu>

Description Infrastructure for eQTL, mQTL and similar studies.

Suggests geuvStore, knitr, rmarkdown, BiocStyle, RUnit, GGtools,
Homo.sapiens, IRanges

Imports GenomicRanges, methods, BatchJobs, BBmisc, S4Vectors, ff,
ffbase, BiocGenerics, foreach, doParallel

Depends

Maintainer VJ Carey <stvjc@channing.harvard.edu>

License Artistic-2.0

LazyLoad yes

VignetteBuilder knitr

BiocViews SNP, GenomeAnnotation, Genetics, DataImport,
FunctionalGenomics

NeedsCompilation no

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gQTLBase-package

*gQTLBase: infrastructure for eQTL, mQTL and similar studies***Description**

Infrastructure for eQTL, mQTL and similar studies.

Details

The DESCRIPTION file:

```

Package:      gQTLBase
Title:        gQTLBase: infrastructure for eQTL, mQTL and similar studies
Version:      1.0.0
Author:       VJ Carey <stvjc@channing.harvard.edu>
Description:  Infrastructure for eQTL, mQTL and similar studies.
Suggests:    geuvStore, knitr, rmarkdown, BiocStyle, RUnit, GGtools, Homo.sapiens, IRanges
Imports:      GenomicRanges, methods, BatchJobs, BBmisc, S4Vectors, ff, ffbase, BiocGenerics, foreach, doParallel
Depends:
Maintainer:   VJ Carey <stvjc@channing.harvard.edu>
License:      Artistic-2.0
LazyLoad:    yes
VignetteBuilder: knitr
BiocViews:    SNP, GenomeAnnotation, Genetics, DataImport, FunctionalGenomics

```

Index of help topics:

```

ciseStore-class      Class '"ciseStore"'
extractByProbes      retrieve eqtlTest results from a ciseStore
                     instance
gQTLBase-package     gQTLBase: infrastructure for eQTL, mQTL and
                     similar studies
storeApply           apply a function over job results in a
                     ciseStore instance
storeMapResults      use batchMapResults infrastructure to process
                     results in a ciseStore instance
storeToff            extract a vector from store results as ff (out
                     of memory reference); support statistical
                     reductions

```

Purpose is to define infrastructure on a comprehensive archive of eQTL, mQTL, dsQTL, etc., association statistics.

Package will complement gQTLStats. geuvStore is a basic illustration relative to GEUVADIS paper. [matprint](#) is exported from package ff.

Author(s)

VJ Carey <stvjc@channing.harvard.edu>

Maintainer: VJ Carey <stvjc@channing.harvard.edu>

ciseStore-class	Class "ciseStore"
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Description

wrap a BatchJobs registry that manages results of a cis-eQTL search

Objects from the Class

Objects can be created by calls of the form `new("ciseStore", reg, ...)`.

We can also use `ciseStore(reg, addProbeMap = TRUE, addRangeMap = TRUE)` and the `probeMap` and `rangeMap` slots will be populated appropriately

Slots

`reg`: Object of class "Registry" BatchJobs Registry instance

`validJobs`: Object of class "integer" vector of valid job identifiers for the registry

`probemap`: Object of class "data.frame" a map from expression probe identifiers to job identifiers where results for the probe are stored

`rangeMap`: Object of class "GRanges" a map from ranges on chromosomes, to job identifiers, in `mcols()$jobid`

Methods

`show`

Note

the construction of the maps occurs via [storeApply](#), which

will use [foreach](#), so that registration of a parallel back end using, e.g., [registerDoParallel](#), will determine the speed of construction

Examples

```
showClass("ciseStore")
# get the global assignment back
require(BatchJobs)
if (require(geuvStore)) {
  reg = partialRegistry()
  store = ciseStore(reg, addProbeMap=TRUE, addRangeMap=FALSE)
  store
}
```

extractByProbes *retrieve eqtlTest results from a ciseStore instance*

Description

retrieve eqtlTest results from a ciseStore instance

Usage

```
extractByProbes(store, probeids, extractTag = "probeid")
extractByRanges(store, gr)
```

Arguments

store	instance of ciseStore-class
probeids	vector character tokens
gr	instance of GRanges-class
extractTag	character atom telling what field in the archived GRanges is regarded as the probe or gene identifier

Details

an index will be searched if created by the ciseStore constructor

Value

a GRanges instance

Author(s)

VJ Carey <stvjc@channing.harvard.edu>

Examples

```
if (require(geuvStore)) {
  reg = partialRegistry()
  store = ciseStore(reg, addProbeMap=TRUE, addRangeMap=TRUE)
  ebp = extractByProbes(store, c("ENSG00000183814.10", "ENSG00000174827.9"))
  ebp
  rr = range(ebp)
  ebr = extractByRanges(store, rr)
  ebr
}
```

storeApply	<i>apply a function over job results in a ciseStore instance</i>
------------	--

Description

apply a function over job results in a ciseStore instance

Usage

```
storeApply(store, f, n.chunks, ids=NULL, ..., verbose = FALSE)
```

Arguments

store	instance of ciseStore-class
f	function on GRanges stored in ciseStore
n.chunks	Number of chunks into which the jobs are to be broken; the series of chunks is handed to foreach to extract results and apply f to them. If missing, the value of <code>getDoParWorkers()</code> used.
ids	defaults to NULL; if non-null, the jobs to be processed are limited to those identified in this vector.
...	additional arguments to <code>foreach</code>
verbose	if TRUE will allow progressbars and other messages to display

Details

The chunking of job identifiers will determine the degree of parallelization of application, and the form of the list that is returned.

Value

A list whose structure depends on the chunking of job identifiers. See the examples.

Note

`eqtlStore` imports `BiocParallel`'s `bpparam` function, and this determines in real time the number of workers to be employed by `storeApply`.

See Also

[storeMapResults](#) will apply over the store using the batch jobs submission infrastructure and can target specific results via `ids`; `storeApply` uses `bplapply` over the entire store

Examples

```

if (require(geuvStore)) {
  require(BatchJobs)
  reg = partialRegistry()
  store = ciseStore(reg, addProbeMap=FALSE, addRangeMap=FALSE)
  storeApply(store, length)
  storeApply(store, length, ids=c(1:3,603))
}

```

storeMapResults	<i>use batchMapResults infrastructure to process results in a ciseStore instance</i>
-----------------	--

Description

use batchMapResults infrastructure to process results in a ciseStore instance

Usage

```

storeMapResults(store, reg2, fun, ...,
  ids = NULL, part = NA_character_, more.args = list())
loadAndFilterResult(reg,
  id, filter=force, part = NA_character_, missing.ok = FALSE)

```

Arguments

store	an instance of ciseStore-class
reg	instance of BatchJobs Registry class
reg2	an empty instance of the Registry class (see makeRegistry)
fun	A function to map over results in store, with formals (job, res, ...).
filter	a function that accepts and returns a GRanges instance, to be applied just after loading a result from the store
...	additional arguments to vectorize over (should be same length as length(findDone(store@reg)))
ids	ids of job results to be mapped; if missing, map all job results
id	a single job id
part	see batchMapResults
missing.ok	see loadResult
more.args	a list of other arguments to be passed to fun; default is empty list.

Value

integer vector with job ids. Main purpose is to prepare the registry for submitJobs.

Note

loadAndFilterResult is not intended to be exported and may be removed in future versions.

Author(s)

VJ Carey <stvjc@channing.harvard.edu>

Examples

```
if (require(geuvStore)) {
  require(BatchJobs)
  reg = partialRegistry()
  store = ciseStore(reg, addProbeMap=FALSE, addRangeMap=FALSE)
  fd = tempfile()
  tempreg = makeRegistry("tempSMR", file.dir=fd)
  storeMapResults( store, tempreg, fun=function(job, res, ...) length(res) )
  showStatus(tempreg)
  submitJobs(tempreg, 1:2)
  loadResults(tempreg)
  unlink(fd)
}
```

storeToFf	<i>extract a vector from store results as ff (out of memory reference); support statistical reductions</i>
-----------	--

Description

extract a vector from store results as ff (out of memory reference); support statistical reductions

Usage

```
storeToFf(store, field, ids = NULL, filter=force, ..., checkField = FALSE,
  ischar=FALSE)
```

Arguments

store	instance of ciseStore-class
field	character tag, length one. If name of a numeric field in the result set (typically something like 'chisq' in the GRanges generated by cisAssoc), ff is applied directly. Character variables are converted to factors before ff is applied.
ids	job ids to be used; if NULL, process all jobs
filter	function to be applied when GRanges is loaded from results store, should accept and return a GRanges instance
...	supplied to makeRegistry for a temporary registry: typically will be a vector of package names if additional packages are needed to process results

<code>checkField</code>	if TRUE steps will be taken to verify that the tag to which 'field' evaluates is present in result in the first job
<code>ischar</code>	must be true for character vector to be handled properly as a factor, otherwise NA will be returned

Details

uses current BatchJobs configuration to parallelize extraction; `reduceResults` could be used for a sequential solution

Value

a vector as ff reference

Note

uses `ffbase:::c.ff` explicitly to concatenate outputs; there is no guarantee of order among elements

Examples

```
if (require(geuvStore)) {  
  require(BatchJobs)  
  reg = partialRegistry()  
  store = ciseStore(reg, addProbeMap=FALSE, addRangeMap=FALSE)  
  smchisq = storeToFf( store, "chisq", ids=store@validJobs[1:3])  
  smchisq  
}
```


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