

Package ‘geuvStore2’

October 18, 2017

Title demonstrate storage discipline for eQTL enumerations, revised

Version 1.6.0

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

Description demonstrate storage discipline for eQTL enumerations and analyses based on a selection of GEUVADIS results

Suggests Homo.sapiens, knitr (>= 1.7), rmarkdown

Imports methods, gQTLBase

Depends BatchJobs, GenomicRanges

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

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LazyLoad yes

biocViews ExperimentData, SequencingData, MicroarrayData

VignetteBuilder knitr

NeedsCompilation no

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geuvStore2-package	<i>demonstrate storage discipline for eQTL enumerations, revised</i>
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Description

demonstrate storage discipline for eQTL enumerations and analyses based on a selection of GEUVADIS results

Details

The DESCRIPTION file:

```
Package:      geuvStore2
Title:       demonstrate storage discipline for eQTL enumerations, revised
Version:     1.6.0
Author:      VJ Carey <stvjc@channing.harvard.edu>
Description: demonstrate storage discipline for eQTL enumerations and analyses based on a selection of GEUVADIS
Suggests:   Homo.sapiens, knitr (>= 1.7), rmarkdown
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biocViews:  ExperimentData, SequencingData, MicroarrayData
VignetteBuilder: knitr
```

Index of help topics:

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geuvStore2-package  demonstrate storage discipline for eQTL
                    enumerations, revised
gv2pm               components for constructing the distributed
                    store for the subset of eQTL association
                    statistics based on GEUVADIS
```

This package illustrates an approach to managing large numbers of tests, in this case, eQTL association statistics and annotation. The idea is that the report is broken into small chunks, indexed by feature names and ranges, and can be filtered using parallel computing if desired.

The `makeGeuvStore2` function constructs the basic object for working with a subset of tests related to the GEUVADIS study [PMID 24037378].

Author(s)

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Examples

```
mm = makeGeuvStore2()
mm
data(geuvStore2Desc)
geuvStore2Desc # describes distributed computing resources used
```

gv2pm

components for constructing the distributed store for the subset of eQTL association statistics based on GEUVADIS

Description

gv2pm and *gv2rm* are used to index distributed *GRanges* objects; *geuvStore2Desc* is a serialized description of store contents. *kpp* is the vector of gene identifiers retained in this subset (about 5

Usage

```
data("gv2pm")
```

Format

A data frame with 920 observations on the following 2 variables.

probeid a character vector

jobnum a numeric vector

Examples

```
data(gv2rm)  
gv2rm
```

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