Package 'flowWorkspaceData'

April 16, 2024

Type Package

Title A data package containing two flowJo, one diva xml workspace and the associated fcs files as well as three GatingSets for testing the flowWorkspace, openCyto and CytoML packages.

Version 3.15.0

Date 2011-03-31

Author Greg Finak

Maintainer Mike Jiang <wjiang2@fhcrc.org>

Description The necessary external data to run the flowWorkspace and openCyto vignette is found in this package.

biocViews ExperimentData, FlowCytometryData

License GPL-2

LazyLoad yes

git_url https://git.bioconductor.org/packages/flowWorkspaceData

git_branch devel

git_last_commit eba87ed

git_last_commit_date 2023-10-24

Repository Bioconductor 3.19

Date/Publication 2024-04-16

Contents

flowWorkspaceData-package 2
flowWorkspaceDataInfo 3

Index **4**

`flowWorkspaceData-package` *A flowJo XML Workspace and Associated Data for testing the flowWorkspace package.*

Description

This package contains fcs data files and xml workspaces for testing the flowWorkspace and openCyto packages and building the their vignettes. These can be located in the inst/extdata directory of the flowWorkspaceData package. The sample files and workspaces contain data from whole blood. The primary purpose of this data set is to test the import code and export code in the flowWorkspace package and running automated gating pipeline in the openCyto package.

Details

Package:	flowWorkspaceData
Type:	Package
Version:	1.1.0
Date:	2011-03-31
License:	GPL 2.0
LazyLoad:	yes

flowWorkspaceDataInfo() provides information about the package.

Author(s)

Greg Finak

Maintainer: <wjiang2@fhcrc.org>

See Also

`flowWorkspace` `openCyto`

Examples

```
flowWorkspaceDataInfo()
```

flowWorkspaceDataInfo *Print some information about this package and the data.*

Description

Contains the data necessary to build the flowWorkspace and openCyto vignettes.

Usage

```
flowWorkspaceDataInfo()
```

Author(s)

Greg Finak

See Also

flowWorkspace-package openCyto

Examples

```
flowWorkspaceDataInfo();
```

Index

∗ **package**
flowWorkspaceData-package, 2

flowWorkspace, *2*
flowWorkspaceData
*(*flowWorkspaceData-package*)*, 2
flowWorkspaceData-package, 2
flowWorkspaceDataInfo, 3

openCyto, *2*, *3*