Package 'flowWorkspaceData'

January 2, 2025

| 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 3.19.0 |
| Date 2011-03-31 |
| Author Greg Finak |
| Maintainer Mike Jiang <wjiang2@fhcrc.org></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 |
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| |
| Contents |
| flowWorkspaceData-package |
| Index |

flowWorkspaceData-package

A flowJo XML Workspace and Associated Data for testing the flow-Workspace package.

Description

This package contains fcs data files and xml workspaces for testing the flowWorkspace and open-Cyto 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()

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