# Package 'flowWorkspaceData'

May 30, 2024

Type Package
<b>Title</b> 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.
<b>Version</b> 3.16.0
<b>Date</b> 2011-03-31
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<b>Description</b> 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 RELEASE_3_19
git_last_commit ed6a37c
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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)

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#### 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();

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