# Package 'flowWorkspaceData'

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Туре	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.
Versi	on 3.21.0
Date	2011-03-31
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Desci	<b>ription</b> The necessary external data to run the flowWorkspace and openCyto vignette is found in this package.
biocV	Views ExperimentData, FlowCytometryData
Licen	se GPL-2
Lazy	Load yes
git_u	rl https://git.bioconductor.org/packages/flowWorkspaceData
git_b	ranch devel
git_la	st_commit 0c71b0f
git_la	sst_commit_date 2025-04-15
Repo	sitory Bioconductor 3.22
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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()

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();

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