# Package 'flowWorkspaceData'

October 12, 2023

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.
Versi	on 3.12.0
Date	2011-03-31
Autho	or Greg Finak
Main	tainer Mike Jiang <wjiang2@fhcrc.org></wjiang2@fhcrc.org>
Descr	<b>ription</b> The necessary external data to run the flowWorkspace and openCyto vignette is found in this package.
biocV	Yiews ExperimentData, FlowCytometryData
Licen	se GPL-2
Lazyl	Load yes
git_uı	rl https://git.bioconductor.org/packages/flowWorkspaceData
git_bı	ranch RELEASE_3_17
git_la	st_commit b5e006c
git_la	st_commit_date 2023-04-25
Date/	<b>Publication</b> 2023-10-12
R to	opics documented:
	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**