# Package 'flowWorkspaceData'

July 10, 2025

Type I	Package
t	A data package containing two flowJo, one diva xml workspace and he associated fcs files as well as three GatingSets for esting the flowWorkspace, openCyto and CytoML packages.
Version	n 3.21.0
Date 2	2011-03-31
Author	r Greg Finak
Mainta	ainer Mike Jiang <wjiang2@fhcrc.org></wjiang2@fhcrc.org>
-	<b>ption</b> The necessary external data to run the flowWorkspace and openCyto vignette is found in this package.
biocVi	ews ExperimentData, FlowCytometryData
License	e GPL-2
LazyL	oad yes
git_url	https://git.bioconductor.org/packages/flowWorkspaceData
git_bra	anch devel
git_las	<b>t_commit</b> 0c71b0f
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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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