Package 'flowWorkspaceData'

August 21, 2025

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	
Auth	or Greg Finak	
Main	tainer Mike Jiang <wjiang2@fhcrc.org></wjiang2@fhcrc.org>	
Desci	ription The necessary external data to run the flowWorkspace and openCyto vignette is found in this package.	
biocV	Views ExperimentData, FlowCytometryData	
Licer	ase GPL-2	
Lazy	Load yes	
git_u	rl https://git.bioconductor.org/packages/flowWorkspaceData	
git_b	ranch devel	
git_la	ast_commit 0c71b0f	
git_la	ast_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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