${\bf Package~`flow Work space Data'}$

October 16, 2018

Type	Package Package	
Title	ta package containing two flowJo, one diva xml workspace and ssociated fcs files as well as three GatingSets for the flowWorkspace, openCyto and CytoML packages.	
Version 2.16.0		
Date	03-31	
Auth	or Greg Finak	
Main	ntainer 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	
Licen	nse GPL-2	
Lazy	Load yes	
git_url https://git.bioconductor.org/packages/flowWorkspaceData		
git_b	oranch RELEASE_3_7	
git_la	ast_commit 62864da	
git_la	ast_commit_date 2018-04-30	
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R to	opics documented:	
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flo	owWorkspaceData-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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