Package 'flowWorkspaceData'

September 1, 2022

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 3.9.0
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
Licen	ase GPL-2
Lazy	Load yes
git_u	rl https://git.bioconductor.org/packages/flowWorkspaceData
git_b	ranch master
git_la	nst_commit 90c0661
git_la	nst_commit_date 2022-04-26
Date/	Publication 2022-09-01
R to	opics documented:
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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)

Greg Finak

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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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