### Package ‘flowWorkspaceData’

February 1, 2017

**Type** Package

**Title** A data package containing two flowJo xml workspaces, associated fcs files and three GatingSets for testing the flowWorkspace and openCyto packages.

**Version** 2.10.0

**Date** 2011-03-31

**Author** Greg Finak

**Maintainer** Mike Jiang <wjiang2@fhcrc.org>

**Description** The necessary external data to run the flowWorkspace and openCyto vignette is found in this package.

**biocViews** ExperimentData, FlowCytometryData

**License** GPL-2

**LazyLoad** yes

**NeedsCompilation** no

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### flowWorkspaceData-package

_A flowJo XML Workspace and Associated Data for testing the flowWorkspace package._

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### Description

This package contains fcs data files and xml workspaces for testing the flowWorkspace and openCyto 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.

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### Details
flowWorkspaceDataInfo() provides information about the package.

Author(s)

Greg Finak
Maintainer: <wjiang2@fhcrc.org>

See Also

flowWorkspace openCyto

Examples

flowWorkspaceDataInfo()

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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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