

Package ‘cytolib’

September 15, 2024

Type Package

Title C++ infrastructure for representing and interacting with the gated cytometry data

Version 2.16.0

Date 2017-08-07

Author Mike Jiang

Maintainer Mike Jiang <mike@ozette.ai>

Description This package provides the core data structure and API to represent and interact with the gated cytometry data.

License AGPL-3.0-only

License_restricts_use no

LazyLoad yes

Imports RProtoBufLib

Depends R (>= 3.4)

Suggests knitr, rmarkdown

LinkingTo BH(>= 1.84.0.0), RProtoBufLib(>= 2.13.1),Rhdf5lib

biocViews ImmunoOncology, FlowCytometry, DataImport, Preprocessing, DataRepresentation

VignetteBuilder knitr

SystemRequirements GNU make, C++11

RoxygenNote 7.1.2

Biarch TRUE

git_url <https://git.bioconductor.org/packages/cytolib>

git_branch RELEASE_3_19

git_last_commit 939e51d

git_last_commit_date 2024-04-30

Repository Bioconductor 3.19

Date/Publication 2024-09-15

Contents

cytolib_LdFlags 2

Index **3**

cytolib_LdFlags	<i>Output the LD flags for building against cytolib. These flags are propagated to sourceCxx via the inlineCxxPlugin (defined below) and to packages via a line in Makevars[.win] like this:</i>
-----------------	--

Description

```
PKG_LIBS += $(shell "$R_HOME/bin/$R_ARCH_BIN/Rscript.exe" -e "cytolib::cytolib_LdFlags()")
```

Usage

```
cytolib_LdFlags()
```

Index

cytolib_LdFlags, [2](#)