

Package ‘cytolib’

April 15, 2020

Type Package

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

Version 1.8.0

Date 2017-08-07

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

License Artistic-2.0

LazyLoad yes

Collate 'zzz.R'

Depends R (>= 3.4)

Suggests knitr

biocViews ImmunoOncology, FlowCytometry, DataImport, Preprocessing, DataRepresentation

VignetteBuilder knitr

SystemRequirements GNU make, C++11

RoxygenNote 6.0.1

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

git_branch RELEASE_3_10

git_last_commit 47dcb83

git_last_commit_date 2019-10-29

Date/Publication 2020-04-14

R topics documented: