

Package ‘cytolib’

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Type Package

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

Version 1.0.1

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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 FlowCytometry, DataImport, Preprocessing, DataRepresentation

LinkingTo BH(>= 1.62.0-1), RProtoBufLib

VignetteBuilder knitr

SystemRequirements GNU make, C++11

RoxygenNote 6.0.1

NeedsCompilation yes

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LdFlags

linker flags Generate linker flags This function prints the absolute path to the protobuf library file that is shipped with the package. So that user package can use it in their Makevar file.

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Usage

LdFlags()

Value

nothing.

Examples

LdFlags()

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