

Package ‘cytolib’

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Type Package

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

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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(>= 1.1.2)

VignetteBuilder knitr

SystemRequirements GNU make, C++11

RoxygenNote 6.0.1

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

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R topics documented: