

Package: highthroughputassays (via r-universe)

October 4, 2026

Title Using Bioconductor with High Throughput Assays

Version 1.36.0

Date 2017-09-05

Description The workflow illustrates use of the flow cytometry packages to load, transform and visualize the flow data and gate certain populations in the dataset. The workflow loads the `flowCore`, `flowStats` and `flowWorkspace` packages and its dependencies. It loads the ITN data with 15 samples, each of which includes, in addition to FSC and SSC, 5 fluorescence channels: CD3, CD4, CD8, CD69 and HLADR.

Depends R (>= 3.3.0), flowCore, flowStats, flowWorkspace

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Encoding UTF-8

LazyData true

VignetteBuilder knitr

biocViews ImmunoOncologyWorkflow, Workflow, ProteomicsWorkflow

Suggests knitr, rmarkdown, BiocStyle

NeedsCompilation no

URL <https://www.bioconductor.org/help/workflows/highthroughputassays/>

Config/pak/sysreqs make libjpeg-dev libpng-dev libxml2-dev libssl-dev zlib1g-dev

Repository Bioconductor 3.23

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RemoteUrl <https://github.com/bioc/highthroughputassays>

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