

Package ‘iSEHub’

January 30, 2026

Title iSEE for the Bioconductor ExperimentHub

Version 1.12.0

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Description This package defines a custom landing page for an iSEE app interfacing with the Bioconductor ExperimentHub. The landing page allows users to browse the ExperimentHub, select a data set, download and cache it, and import it directly into a Bioconductor iSEE app.

License Artistic-2.0

URL <https://github.com/iSEE/iSEHub>

BugReports <https://support.bioconductor.org/t/iSEHub>

biocViews DataImport, ImmunoOncology Infrastructure, ShinyApps, SingleCell, Software

Encoding UTF-8

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Depends SummarizedExperiment, SingleCellExperiment, ExperimentHub

Imports AnnotationHub, BiocManager, DT, iSEE, methods, rintrojs, S4Vectors, shiny, shinydashboard, shinyjs, utils

Suggests BiocStyle, covr, knitr, RefManageR, rmarkdown, sessioninfo, testthat (>= 3.0.0), nullrangesData

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Description

iSEEhub app

Usage

iSEEhub(ehub, runtime_install = FALSE)

Arguments

- ehub An [ExperimentHub\(\)](#) object.
- runtime_install A logical scalar indicating whether the app may allow users whether to install data set dependencies at runtime using [BiocManager::install\(\)](#) through a modal prompt.

Value

An [iSEE\(\)](#) app with a custom landing page interfacing with ehub.

Examples

```
library(ExperimentHub)
ehub <- ExperimentHub()

app <- iSEEhub(ehub)

if (interactive()) {
  shiny::runApp(app, port = 1234)
}
```

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