

Package: imageTCGA (via r-universe)

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Type Package

Title TCGA Diagnostic Image Database Explorer

Version 1.4.0

Description A Shiny application to explore the TCGA Diagnostic Image Database.

License Artistic-2.0

Encoding UTF-8

LazyData false

biocViews ShinyApps

Imports DT, dplyr, bslib, bsicons, ggplot2, viridis, tidyr, leaflet, clipr, rlang

Suggests BiocManager, BiocStyle, knitr, curl, glue, rmarkdown, sessioninfo, testthat, tibble, GenomicDataCommons

RoxygenNote 7.3.2

Depends shiny

BugReports <https://github.com/billila/imageTCGA/issues>

URL <https://github.com/billila/imageTCGA>

VignetteBuilder knitr

Config/pak/sysreqs libabsl-dev cmake libgdal-dev gdal-bin libgeos-dev make libicu-dev libpng-dev libuv1-dev libssl-dev libproj-dev libsqlite3-dev libudunits2-dev libx11-dev zlib1g-dev

Repository <https://bioc-release.r-universe.dev>

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`imageTCGA`*imageTCGA: A Shiny application to explore the TCGA Diagnostic Image Database*

Description

This function launches a Shiny application that allows users to explore and download data from the TCGA Diagnostic Image Database.

Usage

```
imageTCGA()
```

Value

A Shiny application object

Examples

```
if (interactive()) {  
  imageTCGA()  
}
```

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