

Package: BioImageDbs (via r-universe)

October 10, 2026

Type Package

Title Bio- and biomedical imaging dataset for machine learning and deep learning (for ExperimentHub)

Version 1.21.0

Encoding UTF-8

Depends R (>= 4.1.0)

Imports ExperimentHub, AnnotationHub, markdown, rmarkdown, EBImage, magick, magrittr, filesstrings, animation, einsum

Suggests knitr, BiocStyle, purrr

Description The package provides a bioimage dataset for the image analysis using machine learning and deep learning. The dataset includes microscopy imaging data with supervised labels. The data is provided as R list data that can be loaded to Keras/tensorflow in R.

VignetteBuilder knitr

License Artistic-2.0

URL <https://kumes.github.io/BioImageDbs/>

LazyLoad yes

biocViews ExperimentHub, ExperimentData, CellCulture, Tissue

RoxygenNote 7.1.1

Config/pak/sysreqs cmake libfftw3-dev make libmagick++-dev gsfonts texlive libicu-dev libjpeg-dev libpng-dev libtiff-dev libuv1-dev libssl-dev zlib1g-dev

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

Date/Publication 2026-04-28 12:35:08 UTC

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

RemoteRef HEAD

RemoteSha 5064ecaa874cf7248c035dacd2c04748cfd60544

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Details

The DESCRIPTION file: This package was not yet installed at build time.

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Examples

```
ls("package:BioImageDbs") # This is empty package
```

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