

Package ‘BioImageDbs’

July 15, 2025

Type Package

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

Version 1.16.0

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

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

git_branch RELEASE_3_21

git_last_commit ab06d69

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Repository Bioconductor 3.21

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Contents

BioImageDbs-package	2
-------------------------------	---

Index	3
--------------	----------

BioImageDbs-package	<i>Bio- and biomedical imaging dataset for machine learning and deep learning (for ExperimentHub)</i>
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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.

Details

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

Index: This package was not yet installed at build time.

Author(s)

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Maintainer: Satoshi Kume <satoshi.kume.1984@gmail.com>

Examples

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

Index

* **package**

BioImageDbs-package, [2](#)

BioImageDbs (BioImageDbs-package), [2](#)

BioImageDbs-package, [2](#)