

Package ‘AHMeSHDBs’

December 10, 2025

Type Package

Title MeSHDBs for AnnotationHub

Version 1.8.0

Encoding UTF-8

Depends R (>= 4.1.0)

Imports AnnotationHub (>= 2.22.0)

Suggests BiocStyle, knitr, rmarkdown

BugReports <https://github.com/rikenbit/AHMeSHDBs/issues>

Description Supplies AnnotationHub with `MeSHDB` NIH MeSH annotation databases for many species. All the SQLite files and metadata.csv are generated by our Snakemake workflow [mesh-workflow](<https://github.com/rikenbit/mesh-workflow>).

VignetteBuilder knitr

License Artistic-2.0

LazyLoad yes

biocViews AnnotationData, FunctionalAnnotation, MeSHDB, AnnotationHub

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

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Repository Bioconductor 3.23

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AHMeSHDBs-package*MeSHDBs for AnnotationHub*

Description

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Details

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

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

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Examples

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

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