

Package ‘AHLRBaseDbs’

December 10, 2025

Type Package

Title LRBaseDbs 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/AHLRBaseDbs/issues>

Description Supplies AnnotationHub with `LRbaseDb` Ligand-Receptor annotation databases for many species. All the SQLite files are generated by our Snakemake workflow [lrbase-workflow](<https://github.com/rikenbit/lrbase-workflow>). For the details, see the README.md of lrbase-workflow.

VignetteBuilder knitr

License Artistic-2.0

LazyLoad yes

biocViews AnnotationData, FunctionalAnnotation, LRBaseDb, AnnotationHub

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

git_branch devel

git_last_commit 0ee3f5c

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Repository Bioconductor 3.23

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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:AHLRBaseDbs") # This is empty package
```

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