

Package: AHMassBank (via r-universe)

September 21, 2026

Type Package

Title MassBank Annotation Resources for AnnotationHub

Version 1.13.0

Encoding UTF-8

Depends R (>= 4.2)

Imports AnnotationHubData (>= 1.5.24)

Suggests BiocStyle, knitr, AnnotationHub (>= 2.7.13), rmarkdown, methods, CompoundDb (>= 1.1.4)

Description Supplies AnnotationHub with MassBank metabolite/compound annotations bundled in CompDb SQLite databases. CompDb SQLite databases contain general compound annotation as well as fragment spectra representing fragmentation patterns of compounds' ions. MassBank data is retrieved from <https://massbank.eu/MassBank> and processed using helper functions from the CompoundDb Bioconductor package into redistributable SQLite databases.

VignetteBuilder knitr

License Artistic-2.0

LazyLoad yes

BugReports <https://github.com/jorainer/AHMassBank/issues>

URL <https://github.com/jorainer/AHMassBank>

biocViews MassSpectrometry, AnnotationHubSoftware

Roxygen list(markdown=TRUE)

RoxygenNote 7.2.3

Config/pak/sysreqs

make libbz2-dev libicu-dev liblzma-dev libpng-dev libxml2-dev libssl-dev xz-utils zlib1g-dev

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

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