Package ‘AHMassBank’

May 17, 2024

Type Package
Title MassBank Annotation Resources for AnnotationHub
Version 1.4.0
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
git_url https://git.bioconductor.org/packages/AHMassBank
git_branch RELEASE_3_19
git_last_commit bcc7fa7
git_last_commit_date 2024-04-30
Repository Bioconductor 3.19
Date/Publication 2024-05-17
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Author(s)

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See Also

Useful links:

- https://github.com/jorainer/AHMassBank
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