Package 'adductData'

November 10, 2026

Title Data from untargeted MS of modifications to Cys34 of serum albumin

Version 1.28.0

Author Josie Hayes

Maintainer Josie Hayes <jlhayes1982@gmail.com>

Description mzXML files from Grigoryan et al 2016 (Anal Chem).

Depends R (>= 3.6)

Imports ExperimentHub (>= 1.9.0), AnnotationHub (>= 2.13.10), stats (>= 3.5.0), utils (>= 3.5.0), methods (>= 3.5.0), datasets (>= 3.5.0)

Suggests knitr (>= 1.15.1), rmarkdown (>= 1.5)

License Artistic-2.0

LazyData false

LazyLoad false

Encoding UTF-8

RoxygenNote 6.1.0

biocViews ExperimentData,MassSpectrometryData,ExperimentHub

VignetteBuilder knitr

git_url https://git.bioconductor.org/packages/adductData

git_branch RELEASE_3_23

git_last_commit 76f624f

git_last_commit_date 2026-04-28

Repository Bioconductor 3.23

Date/Publication 2026-11-10

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adductData *Adduct data from Grigoryan at al 2016. This package contains 2 mzXML files from Grigoryan et al that were used to perform adductomics.*

Description

Adduct data from Grigoryan at al 2016. This package contains 2 mzXML files from Grigoryan et al that were used to perform adductomics.

Value

2 mzXML files and processed data

Source

https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5555296/

Examples

```
## Not run:
hub <- ExperimentHub::ExperimentHub()
x <- AnnotationHub::query(hub, c("adductData"))
## download resource
data = x[[1]]

## End(Not run)
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