Package 'adductData'

June 1, 2023

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

Version 1.16.0

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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

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R topics documented:

| 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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