Package 'msPurityData'

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Type Package Title Fragmentation spectral libraries and data to test the msPurity package Version 1.34.0 Date 12-12-2018 Author Thomas N. Lawson Maintainer Thomas N. Lawson <thomas.nigel.lawson@gmail.com> Description Fragmentation spectral libraries and data to test the msPurity package License GPL (>= 2) LazyData TRUE VignetteBuilder knitr RoxygenNote 5.0.1 Suggests knitr biocViews ExperimentData, MassSpectrometryData NeedsCompilation no git_url https://git.bioconductor.org/packages/msPurityData git_branch RELEASE_3_20 git_last_commit b678757 git_last_commit_date 2024-10-29 **Repository** Bioconductor 3.20 Date/Publication 2024-12-19

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msPurityData-package Test data for the msPurity package

Description

This data package contains test data to be used with package msPurity, see folders lcms and dims. This contains LC-MS, LC-MS/MS and DI-MS datasets in mzML format. There are also various .csv files and .rds files representing model outputs from the msPurity package. The LC-MS, LC-MS/MS and DI-MS datasets have been reduced in size by reducing the number of scans and m/z range.

The data package also contains a fragmentation spectral library created by msp2db (https://msp2db.readthedocs.io/en/late with data from MassBank, GNPS, LipidBlast and HMDB. This is the default spectral library that is used with the spectral_matching with msPurity. The library data is from MoNA (http://mona.fiehnlab.ucdavis.edu/downloc downloaded on 5th November 2018.

The dataset also contains data relating to the msPurity publication.

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