# Package 'RMassBankData'

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Title Test dataset for RMassBank  Version 1.28.0  Author Michael Stravs, Emma Schymanski, Steffen Neumann  Maintainer Michael Stravs, Emma Schymanski <massbank@eawag.ch>  Description Example spectra, example compound list(s) and an example annotation</massbank@eawag.ch>
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<b>Description</b> Example spectra, example compound list(s) and an example annotation
list for a narcotics dataset; required to test RMassBank. The package is described in the man page for RMassBankData. Includes new XCMS test data.
biocViews ExperimentData, MassSpectrometryData
License Artistic-2.0
Suggests RMassBank
Collate 'RMassBankData.R'
git_url https://git.bioconductor.org/packages/RMassBankData
git_branch RELEASE_3_12
git_last_commit 0fe2270
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## Description

This contains data suitable to test the RMassBank functionalities.

### **Details**

The package contains the folders:

- spectra LC-MS runs of 15 narcotics standards, in mzML format and deprofiled.
- listA CSV list with compound informations for the 15 narcotics, as needed by RMassBank.
- infolistsA complete CSV list with annotations for the 15 standards.
- infolists\_incompleteA partial list of infolists, to demonstrate the download of missing entries.
- infolists\_editedThe downloaded missing entries, subsequently checked and completed by hand.
- resultsThe intermediate and final results of the msms\_workflow runs. This serves to build the vignette, since it would take too long to run the whole workflow during the vignette build.

### Author(s)

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