

# Package ‘NMRS’

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**Type** Package

**Title** NMR Spectroscopy

**Version** 1.0

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**Description** NMRS has been developed to load directly the spectra in the Bruker spectroscopy format. This application displays the spectrum reference and manages basic operations such as setting the chemical shift of a certain compound (TSP or DSS) to 0 ppm, zero order and first order phase corrections, baseline adjustment and spectral area selection

**License** GPL (>= 2)

**Depends** Rwave, FTICRMS, tcltk, tkrplot

**Repository** CRAN

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 NMRS-package

*NMR Spectra Preprocessing*


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### Description

This "NMRS" package permits the pre-processing of the raw NMR Bruker spectra. Data pre-processing transforms the data in a way that subsequent analysis and modelling are easier, more robust and more accurate. For the analysis of NMR spectra, pre-processing methods usually intend to reduce variances and influences as phase corrections of each spectrum, baseline corrections, etc.

### Details

Package: NMRS  
 Type: Package  
 Version: 1.0  
 Date: 2009-02-11  
 License: GPL version 2 or newer  
 LazyLoad: yes

The NMRS package has been designed as an interactive process. By Typing NMRS () the user has access to complete preprocessing of the data

### Author(s)

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 Baseline.Correction

*Baseline Correction*


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### Description

Interactive Baseline Correction of raw spectra based in the FTICRMS package.

### Usage

Baseline.Correction(coord)

### Arguments

coord Data.frame with the spectral information, where the first column holds the chemical shift positions and the different spectral intensities are in the followings columns

**Details**

Baseline correction is a very essential step to obtain high quality NMR spectra in some cases. Rolling baselines can make it difficult to identify peaks, as well as introduce significant errors into any quantitative measurements. This function based in the FTICRMS package launches a interactive graphical display to control the individual baseline correction. It computes an estimated baseline curve for a spectrum by a method of Rocke and Xi .

**Value**

a3	Estimated Baseline
datos	Data.frame with the baseline corrected

**Author(s)**

Jose L. Izquierdo <izquierdo@ieb.ucm.es>

**References**

FTICRMS package <http://cran.r-project.org/web/packages/FTICRMS/index.html>

**See Also**

[NMRS](#)

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Fid	<i>Bruker FID loading</i>
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**Description**

This function loads a Bruker FID file and applies the Fast Fourier Transform.

**Usage**

```
Fid()
```

**Details**

The Fid function has been developed to be launched with the [Met.FID](#) Interactive Display

**Value**

fidRaw	Original Bruker FID format
fid	Bruker FID in complex numbers format
SF	SF value
SWHz	SWHz value
XScaleHZ	X scale in Hz

PPM	X scale in PPM
spectrum	FFT of the FID file
tt	Internal value
si	Internal value

**Author(s)**

Jose L. Izquierdo <izquierdo@ieb.ucm.es>

**References**

R.R. Ernst, G. Bodenhausen, A. Wokaun. Principles of Nuclear Magnetic Resonance in One and Two Dimensions. Clarendon Press, Oxford, 2003.

**See Also**

[Met.FID](#), [NMRS](#)

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Manual.cut

*Spectral region selection*

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**Description**

This function selects the spectral region to the statistical analysis. Also, Manual.cut can be used to remove the water peak.

**Usage**

```
Manual.cut(coordenadas)
```

**Arguments**

coordenadas	Data.frame with the spectral information, where the first column holds the chemical shift positions and the different spectral intensities are in the followings columns
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**Details**

Interactive Display.

**Value**

datos	Data.frame with the spectral information
xlim	Internal value
xlim2	Internal value

**Author(s)**

Jose L. Izquierdo <izquierdo@ieb.ucm.es>

**See Also**

[NMRS](#)

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Met.FID

*Interactive Phase Correction and Peak reference*

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**Description**

Interactive graphical display that allow zero order and first order phase corrections. In addition the user can set the chemical shift of a certain compound (TSP or DSS) to 0 ppm as peak reference.

**Usage**

Met.FID()

**Details**

The Met.FID graphical Display runs into the NMRS main function.

**Value**

spect.fid      Preprocessed spectra

**Author(s)**

Jose L. Izquierdo <izquierdo@ieb.ucm.es>

**See Also**

[Fid, phase, NMRS](#)

Met.spectrum

*Spectrum Graphical Display*

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**Description**

Interactive Graphical Display

**Usage**

Met.spectrum(xCoords)

**Arguments**

xCoords            Data frame to plot

**Details**

Internal Function

**Author(s)**

Jose L. Izquierdo <izquierdo@ieb.ucm.es>

**See Also**

[Met.FID, NMRS](#)

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NMRS*Preprocessing of NMR Spectra*

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**Description**

Import.data loads directly the Bruker spectroscopy format (FID file) and applies the Fast Fourier Transform. Furthermore, This application displays the spectrum reference and manages basic operations such as setting the chemical shift of a certain compound (TSP or DSS) to 0 ppm, zero order and first order phase corrections, baseline adjustment,...

**Usage**

NMRS()

**Details**

Interactive application.

**Value**

Spectra            Preprocessed spectra

**Author(s)**

Jose L. Izquierdo <izquierdo@ieb.ucm.es>

**See Also**

[Baseline.Correction](#), [Fid](#), [Manual.cut](#), [phase](#)

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phase                            *Phase Correction*

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**Description**

Zero order and first order phase corrections.

**Usage**

```
phase(fid.out, phc0, phc1, pivot)
```

**Arguments**

fid.out	FID file
phc0	Zero orden correction
phc1	First orden correction
pivot	pivot

**Details**

The phase function has been developed to be launched with the [Met.FID](#) Interactive Display

**Author(s)**

Jose L. Izquierdo <izquierdo@ieb.ucm.es>

**References**

R.R. Ernst, G. Bodenhausen, A. Wokaun. Principles of Nuclear Magnetic Resonance in One and Two Dimensions. Clarendon Press, Oxford, 2003.

**See Also**

[Met.FID](#), [NMRS](#)

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