


AVANCE IVDr

- IVDr Data Browser
User Manual
Version 001



Copyright © by Bruker Corporation

All rights reserved. No part of this publication may be reproduced, stored in a retrieval system, or transmitted, in any form, or by any means without the prior consent of the publisher. Product names used are trademarks or registered trademarks of their respective holders.

This manual was written by

Applied, Industrial & Clinical MR Market Division

© September 21, 2016 Bruker Corporation

Document Number: 10000057225

P/N: H161805

1 Introduction

The Bruker In Vitro Diagnostics (IVDr) Data Browser is a convenient program to scan through NMR data. It simplifies selecting and sorting spectral data by gathering metadata from a locally existing repository of experimental NMR data. Additionally, the metadata of the NMR spectra can be augmented by metadata, e. g. concerning the origin of the measured sample.

Several tools to import and export data, to sort the data and to open data in TopSpin™ or AMIX™ are available.

2 Getting Started

2.1 Searching for Spectral Data

After the initial start of the IVDr Data Browser the tables of the tool (i.e. **Main List** and **Work List**), which represent the main workbench, are empty. In order for data to show up here, select **Search for spectra...** and choose a directory to search in the local file system:

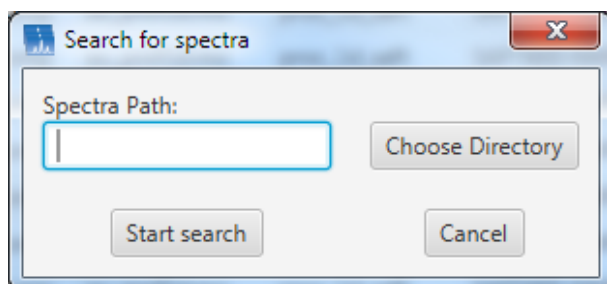


Figure 2.1: Searching for Spectral Data

Press **Start search** to begin the search, it may take several minutes depending on the folder selected and the computer performance.

2.2 Workbench Overview

The workbench consists of two tabs representing two tables with spectral data (see [Figure 2.2 \[▶ 6\]](#)).

- Main List
- Work List

The Main List contains the main spectra data. When a spectra search is performed, all of the metadata from the spectra found will be placed here. The Work List is a second table that can be used as a work area. Data from the Main List can be easily copied to the Work List, e.g. by selecting spectral data in the Main List and pressing the keys **Ctrl+C**. The menu item **Copy spectra to work list** is also available from the context menu of the Main List and from the main menu of the IVDr Data Browser.

As you will notice, certain functions are exclusively available for either the Main List or the Work List. For example, the **Search for spectra...** is only available for the Main List and the execution of TopSpin commands on spectra is only available from the Work List.

Directory	Sample name	Exp.No.	Proc.No.	Experiment	Pulse program	AUNM	AUNMP	Instrument	Probehead	NS	DS	P1	P4B 1	P4B 9	RG
I:\Experiments\Bruker\IVDrForum\S..._urine3_0013		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.11	-12.000018...	45.6479257...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0014		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.87	-12.000018...	45.8084107...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0015		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.36	-12.000018...	45.4838516...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0016		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	12.58	-12.000018...	46.0003662...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0017		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.47	-12.000018...	45.4126276...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0018		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	12.85	-12.000018...	45.8129179...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0019		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.24	-12.000018...	45.5623211...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0020		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	11.68	-12.000018...	46.6511237...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0021		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	11.87	-12.000018...	46.5109669...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0022		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.01	-12.000018...	45.7144459...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0023		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.51	-12.000018...	45.3888793...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0024		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	12.77	-12.000018...	46.8791646...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0025		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	12.55	-12.000018...	46.0271070...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0026		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.65	-12.000018...	46.2973287...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0027		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	12.36	-12.000018...	46.1596108...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0028		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	11.99	-12.000018...	46.4239554...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0029		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.74	-12.000018...	45.2402462...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0030		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	11.88	-12.000018...	46.5003607...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0031		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.47	-12.000018...	45.4126276...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0032		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.87	-12.000018...	45.1349511...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0033		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	12.29	-12.000018...	46.2094917...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0034		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	11.94	-12.000018...	46.4598957...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0035		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.23	-12.000018...	45.5687867...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0036		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	13.91	-12.000018...	45.1334386...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0037		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	12.4	-12.000018...	46.1315462...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0038		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	12.57	-12.000018...	46.0123752...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0039		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	11.81	-12.000018...	46.5598808...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0040		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	11.79	-12.000018...	46.5697052...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0041		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	11.79	-12.000018...	46.5697052...	86.63
I:\Experiments\Bruker\IVDrForum\S..._urine3_0042		10	1	PROF_URINE_NOE5Y	noesyggpr1d	au_urine_noesy	proc_urine_noesy	AV600_HASC	5 mm PABBI 1H-D-88 Z-GRD Z814601.0063	32	4	12.09	-12.000018...	46.3145571...	86.63

Figure 2.2: The IVDr Data Browser Workbench

2.3 Configuration

Before the functions from TopSpin or AMIX can be used, the IVDr Data Browser needs to be configured. To do this go to the configuration menu item and enter the correct path to TopSpin and AMIX (if available):

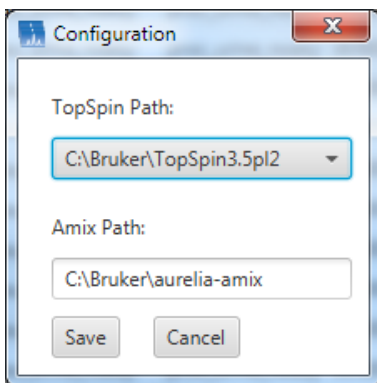


Figure 2.3: Configuration: Selecting the TopSpin and AMIX Path

3 Description of Menu Items and Functions

3.1 Miscellaneous Functions

Open and Save

The IVDr Data Browser saves the current data in the tables when the program is closed. After restarting the IVDr Data Browser, the data in the Main List and the Work List is reloaded.

To store the current data in the program to a file, select **Save** from the **File** menu item. Stored files have the ending *.idb and can be easily loaded by selecting **Open** from the same menu item.

Copy spectra to work list and remove spectra from table

When spectra are selected in the Main List, the list item **Copy spectra to work list** is available.

The item **Remove spectra from table** removes the selected spectra from the current tab.

Copy experiment directories to clipboard

This menu item copies the directories of the selected spectra to the clipboard. At the file system level the directories will be copied to the selected folder by clicking e.g. **Paste** from the context menu of the file browser.

Show in explorer

This menu item allows all the directories of the selected spectra to be opened at once. Use this function with care, because for each selected spectra a new window will be opened.

Filter Spectra and Reset (all) filter

The **Filter Spectra** menu item is available for the Main List and the Work List. It allows certain spectra to be filtered and selected in the foreground table. Through the use of filtering you can search for specific text in a selected column or all available columns.

Filters can also be combined and the complex filter queues can be saved to a file and reloaded from disk. The functions for removing a filter from the queue, loading and saving filters from or to disk are available via the context menu of the filter table:

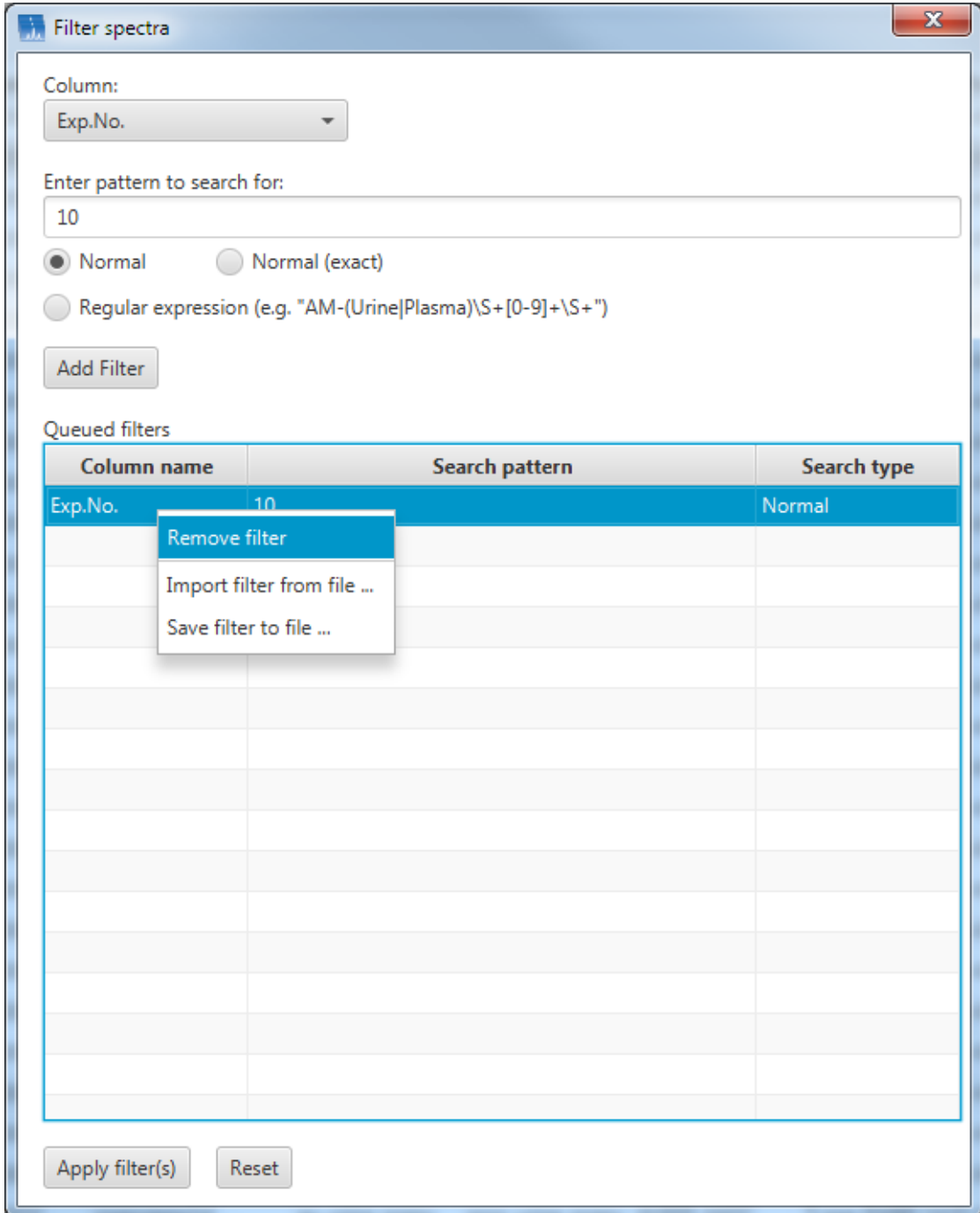


Figure 3.1: Filter Spectra

The menu item **Reset (all) filters** resets all the filters and shows the data of the complete table again. This menu item is only available if a filter was applied in advance.

3.2 Export Spectral Data

Execute TopSpin commands on spectra

The menu item for the execution of TopSpin commands is only available for the Work List. A list of commands can be entered and is executed consecutively for each selected spectra in the Work List:

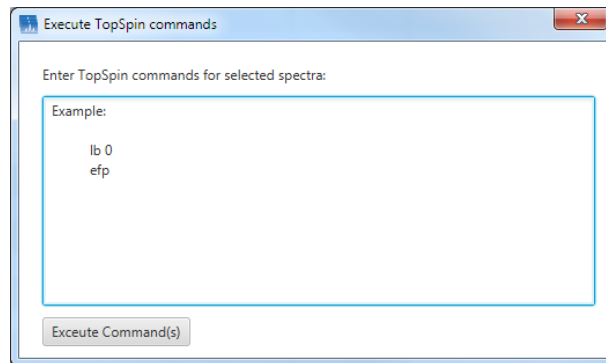


Figure 3.2: Execute TopSpin Commands on Spectra

This function is only available if the configuration for TopSpin was pre-assigned correctly.

Copy experiments to directory

The menu item **Copy experiments to directory ...** copies the experiments folder of each selected spectra to a directory that has to be selected.

Export selected/all experiments to file

The menu items **Export selected experiments to file** and **Export all experiments to file** copy the complete data of the spectra that is observable in the table to a tab-delimited text file.

Export paths of selected/all experiments as list

The menu items **Export paths of selected experiments as list** and **Export paths of all experiments as list** copy the complete paths of the "1r" or "2r" file of the spectra to a text file.

Open in AMIX

The menu item **Open in AMIX** opens the selected spectra of the table in AMIX. To use this function it is necessary to assign the path to AMIX in the IVDr Data Browser configuration.

Open in TopSpin

The menu item **Open in TopSpin** opens the selected spectra of the table in TopSpin. To use this function it is necessary to assign the path to TopSpin in the IVDr Data Browser configuration.

Scan multiple spectra in TopSpin

The menu item **Scan multiple spectra in TopSpin** opens the selected spectra of the table consecutively in TopSpin. A left limit and a right limit for the display of the spectra can be specified and also an interrupt that pauses the loading of the next spectra for a certain time (see figure below). If you interrupt the loading of spectra by pressing the **Pause** button, it is possible to jump forward or backward in the list of selected spectra. To use this function it is necessary to assign the path to TopSpin in the IVDr Data Browser configuration.

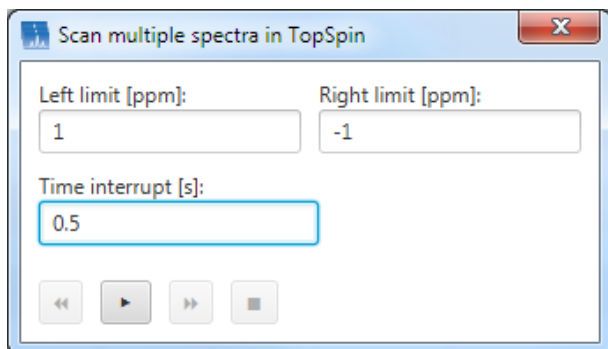


Figure 3.3: Scan Multiple Spectra in TopSpin

3.3 Metadata

Besides the metadata that is gathered directly from the spectral data, it is possible to augment the data with additional data. For example, the data may contain information concerning the origin of the measured sample for example.

Import metadata from file

Metadata can be easily imported from a tab-delimited file. The sample name is used as an identifier and must be unique. The first row of the file must contain the identifiers of the given metadata. All consecutive rows contain the data.

Example file:

```
SampleName <Tab> Gender (male/female) <Tab> Weight (kg) <Tab> Body
length (cm) ...
Urine_001 <Tab> male <Tab> 5.89 <Tab> 58.5 ...
Urine_002 <Tab> female <Tab> 4.53 <Tab> 51.5 ...
...
```

The importing of metadata is available for Main List and Work List. The other functions concerning metadata are only available for the Work List.

Export metadata of selected spectra / all metadata to file

This menu item exports the metadata to a tab-delimited text-file. Besides the metadata the sample name is used as an identifier, thus the sample name should be unique. The identifiers for the columns are given in the first row of the file. See above for an example file output.

Add metadata to XML

The **Add metadata to XML** augments existing XML files with the metadata or creates new XML files that contain the metadata of the spectrum. This function is only available if spectra in the Work List are selected.

When using this function, you have to enter the name of the source XML file and specify the export path. When the source XML file does not exist, then you should check **Create XML-File with metadata if source XML-File is not existent** to create the file with metadata:

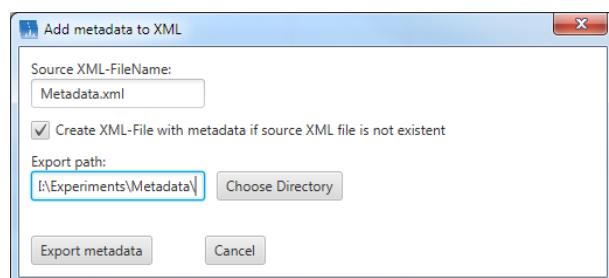


Figure 3.4: Add Metadata to XML

3.4 Customizing the Table View

Select columns in view

The **Select columns in view** menu item offers the opportunity to show or hide certain columns of the table in the foreground:

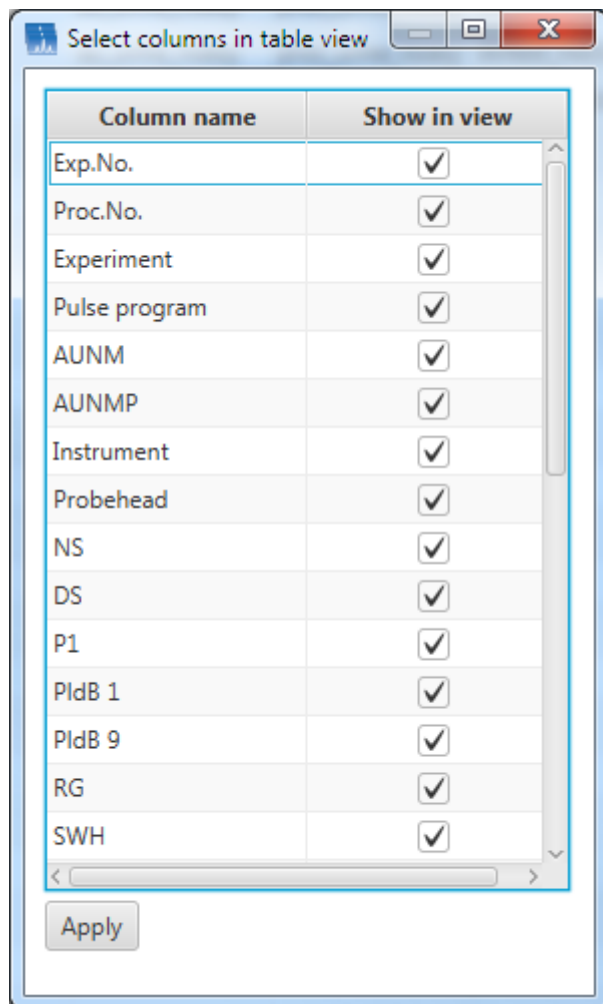


Figure 3.5: Select Columns in View

Show all/show selected columns in table view

These menu items toggle between the view of selected table columns and the view of all columns in the table in the foreground. The list of selected columns can either be modified via the menu item **Select columns in view** or by deleting columns from the table via the context menu of the column title. It is not possible to delete the columns “Directory” and “Sample name” from the view.

3.5 Persistence and Exiting of IVDr Data Browser

Whenever the IVDr Data Browser is closed, the current data in the table is written to a file and is reloaded when the IVDr Data Browser is restarted. Depending on the amount of data in the table and the performance of the computer, this may take a few seconds.

4 Contact

Manufacturer

Bruker BioSpin GmbH
Silberstreifen 4
D-76287 Rheinstetten
Germany
<http://www.bruker.com>

WEEE DE43181702

NMR Hotlines

Contact our NMR service centers.

Bruker BioSpin NMR provides dedicated hotlines and service centers, so that our specialists can respond as quickly as possible to all your service requests, applications questions, software or technical needs.

Please select the NMR service center or hotline you wish to contact from our list available at:

<https://www.bruker.com/service/information-communication/helpdesk.html>

Phone: +49 721-5161-6155

E-mail: nmr-support@bruker.com





Bruker Corporation

info@bruker.com
www.bruker.com

Order No: H161805