



Bruker 1D

Bruker CSI

Import multiple experiments

One click to rule them all → import multiple experiments into jMRUI at the same time

- if same parameters → one multiple-signals MRUI file
- if different parameters → multiple one-signal MRUI files
- possibility to individually exclude files

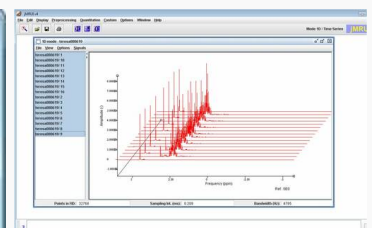
Scanner files compatible from a signal-processing point of view could be non-compatible according to other criteria → the user decides, not jMRUI

Modification made at kernel level that is available for all conversion plug-ins.

If user changes his/her mind, multiple-signal files can be split at any time into multiple one-signal files.

	n°	Experiment	Date modified	SW (Hz)	Transmitter Frequency	Nucleus	B0
<input checked="" type="checkbox"/>	1	1	2010.10.04	4000.000	161.975	1.0	3.804
<input checked="" type="checkbox"/>	2	12	2010.10.04	50000.000	161.975	1.0	3.804
<input checked="" type="checkbox"/>	3	18	2010.10.04	50000.000	161.975	1.0	3.804
<input checked="" type="checkbox"/>	4	24	2010.10.04	50000.000	161.975	1.0	3.804
<input checked="" type="checkbox"/>	5	6	2010.10.04	50000.000	161.975	1.0	3.804

Buttons: OK, Cancel, Select All, Deselect All



Case 1: Click on folder_level0 -folder_level0 - <expname_1> - <expno_1> - <i>fid, acqu...</i> - <expname_2> - <expno_2_1> - <i>fid, acqu...</i> - <expname_3> - <expno_3_1> - <i>fid, acqu...</i> - <expno_3_2> - <i>fid, acqu...</i>	Case 2: Click on <expname> - <expname> - <expno_1> - <i>fid, acqu...</i> - <expno_2> - <i>fid, acqu...</i> - <expno_3> - <i>fid, acqu...</i>	Case 3: Click on <expno> - <expno> - <i>fid, acqu...</i>	Case 4: Click on <i>fid</i> - <i>fid, acqu...</i>
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Import all Bruker fids in experiment folders (expno) pertaining to different "patients" (expname)

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Last update: **2010/11/24 17:43**

