

Version

C

LEOSON BV

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QuantBrowser startershandbook

LEOSON BV

Product manual

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Table of Contents

Concept.....	1
Supported platform.....	2
Installation.....	2
Launch	2
License.....	2
Prepare a dataset.....	3
Calibrate.....	5
Quantitate	9
Review integration.....	10
QEdit.....	11
Column configuration	13
Column information	14
Sample batch	15
Copy to clipboard	17
Export data.....	18

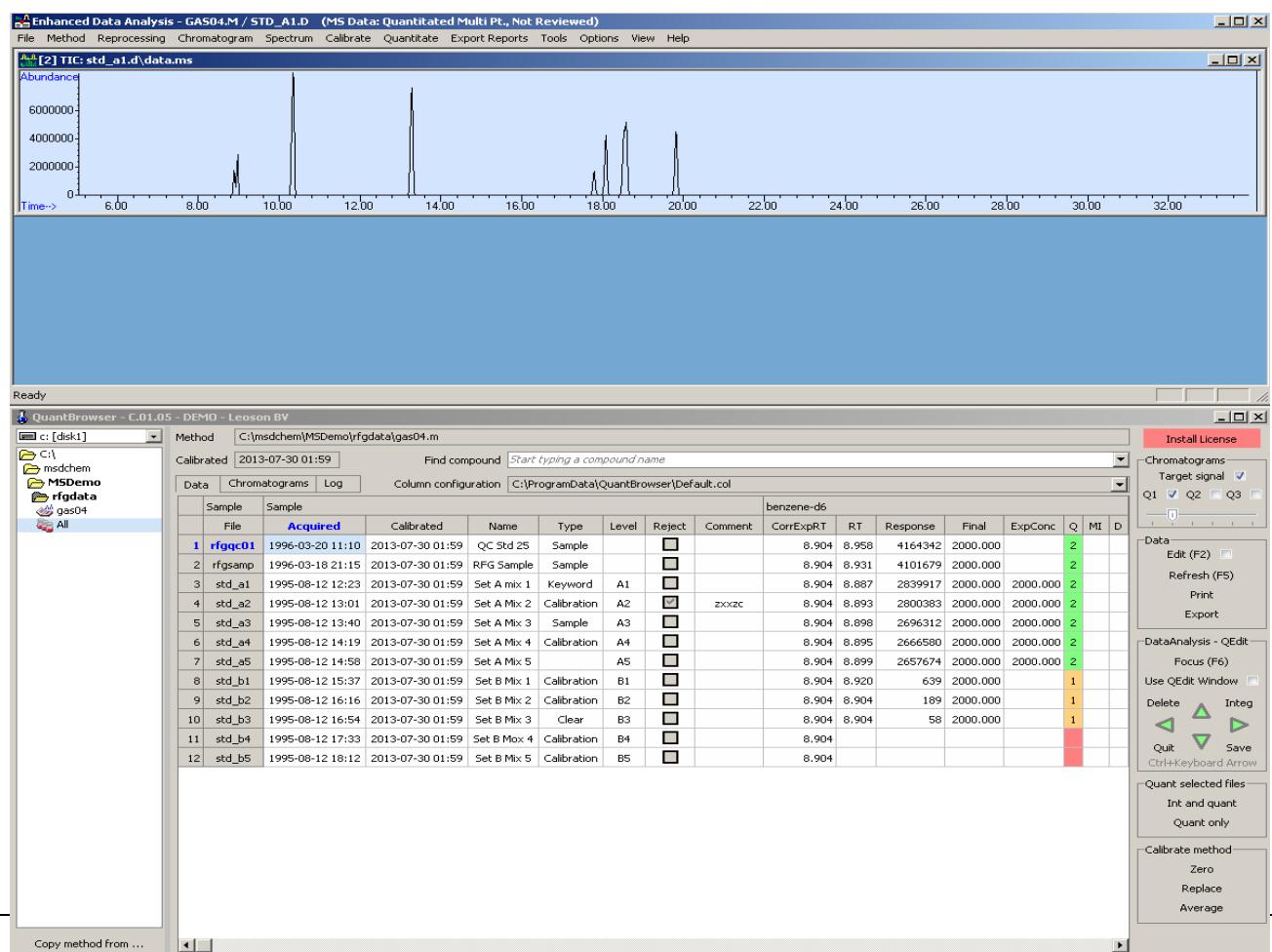
Introduction

Concept

QuantBrowser is an integrated add-on to Agilent DataAnalysis.

DataAnalysis operates on single samples with functions like Quantitation, Calibration, Reporting and QEdit.

QuantBrowser operates on a complete sequence. The results are displayed in a excel-like sheet, as shown in an example below.

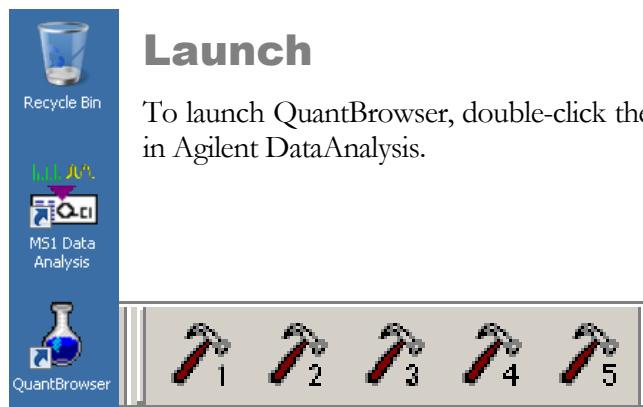


Supported platform

The platforms that support QuantBrowser are Agilent GCMS ChemStation C.00-E.02

Installation

To install QuantBrowser, execute the QuantBrowserSetup.exe. Additional files that are stored in the same folder are automatically installed by the setup program. The additional files are Column configuration files (.col) and a license file (.lic).



License

Install the license file to unlock demo mode. A license file can be obtained through www.waleson.eu.



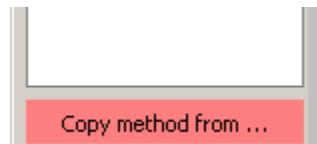
Quick Start

Prepare a dataset

1. Navigate to the data folder



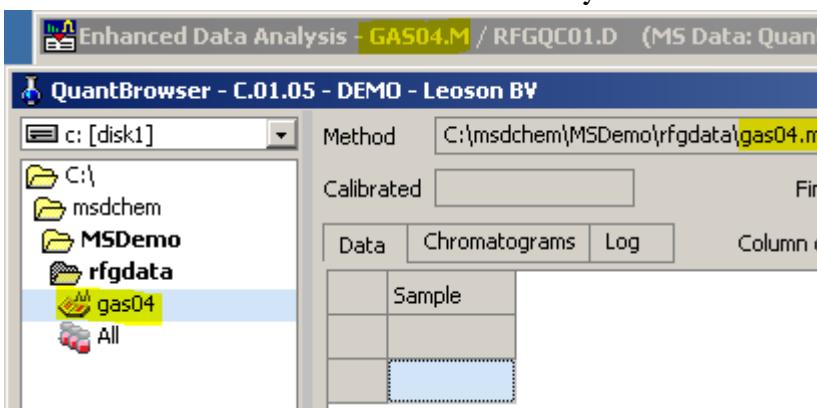
2. Copy a method to the data folder



3. Load the method in DataAnalysis, double-click on the method



4. Find the loaded method name in DataAnalysis title



5. List the samples and results, double click on All

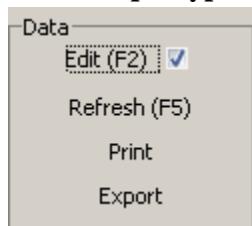


6. Find the samples and results in the table

	Sample	benzene-d6														
		File	Acquired	Calibrated	Name	Type	Level	Reject	Comment	Response	Final	ExpConc	Accuracy	Q	MI	D
1	rfgqc01	1996-03-20 11:10	2013-07-30 01:59	QC Std 25				<input type="checkbox"/>		4164342	2000.000			2		
2	rfgsamp	1996-03-18 21:15	2013-07-30 01:59	RFG Sample				<input type="checkbox"/>		4101679	2000.000			2		
3	std_a1	1995-08-12 12:23	2013-07-30 01:59	Set A mix 1				<input type="checkbox"/>		2839917	2000.000			2		
4	std_a2	1995-08-12 13:01	2013-07-30 01:59	Set A Mix 2				<input type="checkbox"/>		2800383	2000.000			2		
5	std_a3	1995-08-12 13:40	2013-07-30 01:59	Set A Mix 3				<input type="checkbox"/>		2696312	2000.000			2		
6	std_a4	1995-08-12 14:19	2013-07-30 01:59	Set A Mix 4				<input type="checkbox"/>		2666580	2000.000			2		
7	std_a5	1995-08-12 14:58	2013-07-30 01:59	Set A Mix 5				<input type="checkbox"/>		2657674	2000.000			2		
8	std_b1	1995-08-12 15:37	2013-07-30 01:59	Set B Mix 1				<input type="checkbox"/>		639	2000.000			1		
9	std_b2	1995-08-12 16:16	2013-07-30 01:59	Set B Mix 2				<input type="checkbox"/>		189	2000.000			1		
10	std_b3	1995-08-12 16:54	2013-07-30 01:59	Set B Mix 3				<input type="checkbox"/>		58	2000.000			1		
11	std_b4	1995-08-12 17:33	2013-07-30 01:59	Set B Mix 4				<input type="checkbox"/>								
12	std_b5	1995-08-12 18:12	2013-07-30 01:59	Set B Mix 5				<input type="checkbox"/>								

Calibrate

1. Edit sample type and level



2. Set calibration levels, right click in level column

The screenshot shows a context menu titled "Load data file" open over a table. The menu includes options like "Copy" (Ctrl+C), "Windows explorer" (Ctrl+E), "Integrate and quantitate selected datafiles", "Quantitate selected datafiles", "Batch editor ...", "Add to batch ...", "Remove from batch", "Delete batch ...", "Clear", and several level names: A1, A2, A3, A4, A5, B1, B2, B3, B4, and B5. The "B5" option is highlighted with a dark blue background. To the left of the menu is a table with columns "Type" and "Level". The rows show various entries: Control, Sample, Calibration (A1, A2, A3, A4, A5), Calibration (B1, B2, B3, B4, B5). The "Level" column is bolded.

Type	Level
Control	
Sample	
Calibration	A1
Calibration	A2
Calibration	A3
Calibration	A4
Calibration	A5
Calibration	B1
Calibration	B2
Calibration	B3
Calibration	B4
Calibration	B5
Calibration	B5

3. Set sample type, right click in type column

Type	Level	Reject	Comment
Control		<input type="checkbox"/>	
Sample		<input type="checkbox"/>	

Calibrat **Load data file**

Calibrat Copy Ctrl+C

Calibrat Windows explorer Ctrl+E

Calibrat Integrate and quantitate selected datafiles

Calibrat Quantitate selected datafiles

Calibrat Batch editor ...

Calibrat Add to batch ...

Calibrat Remove from batch

Calibrat Delete batch ...

Calibrat Clear

Calibrat Sample

Calibrat Blank

Calibrat Calibration

Calibrat Negative

Calibrat Control

Calibrat Keyword

Calibrat BFB

Calibrat DFTPP

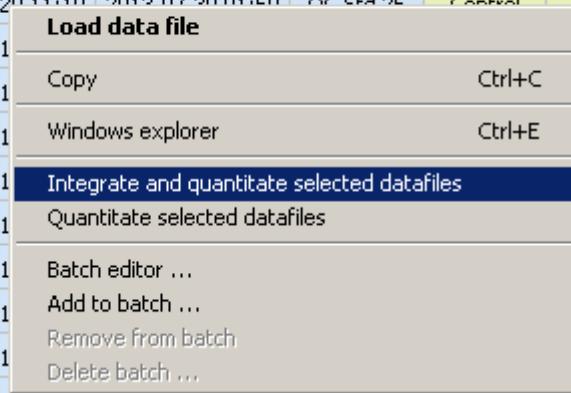


4. Select all samples, click in upper left corner

Data	Chromatograms	Log
	Sample	Sample
	File	Acquired
1	rfgqc01	1996-03-20 11:10 20
2	rfgsamp	1996-03-18 21:15 20

5. Integrate and quantitate selected samples

	Sample	Sample					
	File	Acquired	Calibrated	Name	Type	Level	
1	rfgqc01	1996-03-20 11:10:20	2012-07-20 01:50:00	OC STD 25	Control	Low	
2	rfgsamp	1996-03-1					
3	std_a1	1995-08-1					
4	std_a2	1995-08-1					
5	std_a3	1995-08-1					
6	std_a4	1995-08-1					
7	std_a5	1995-08-1					
8	std_b1	1995-08-1					
9	std_b2	1995-08-1					
..



6. Zero all responses in the calibration table, click on Zero

Calibrate method
Zero
Replace
Average

7. Select the calibration samples

Type	Level
Control	
Sample	
Calibration	A1
Calibration	A2
Calibration	A3
Calibration	A4
Calibration	A5
Calibration	B1
Calibration	B2
Calibration	B3
Calibration	B4
Calibration	B5

8. Replace the responses in the calibration table, click on Replace

Calibrate method
Zero
Replace
Average

8. Note that samples were calculated with an older version of the calibration table

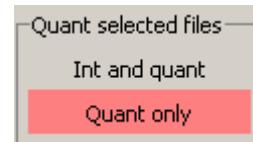
Method C:\msdchem\MSDemo\rfgdata\gas04.m			
Calibrated 2013-08-28 20:53		Find com	
	Data	Chromatograms	Log
	Sample	Sample	
	File	Acquired	Calibrated
1	rfgqc01	1996-03-20 11:10	2013-08-28 20:48
2	rgfsamp	1996-03-18 21:15	2013-08-28 20:48
3	std_a1	1995-08-12 12:23	2013-08-28 20:48
4	std_a2	1995-08-12 13:01	2013-08-28 20:48
5	std_a3	1995-08-12 13:40	2013-08-28 20:48
6	std_a4	1995-08-12 14:19	2013-08-28 20:48
7	std_a5	1995-08-12 14:58	2013-08-28 20:48
8	std_b1	1995-08-12 15:37	2013-08-28 20:48
9	std_b2	1995-08-12 16:16	2013-08-28 20:48
10	std_b3	1995-08-12 16:54	2013-08-28 20:48
11	std_b4	1995-08-12 17:33	2013-08-28 20:48
12	std_b5	1995-08-12 18:12	2013-08-28 20:48

Quantitate

1. Select all samples

	Data	Chromatograms	Log
	Sample	Sample	
1	rfgqc01	1996-03-20 11:10	20
2	rfgsamp	1996-03-18 21:15	20

2. Requantitate all samples, click on Quant only

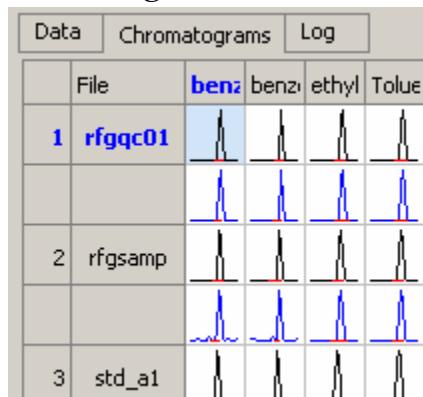


3. Note that the samples were quantitated with the current calibration table

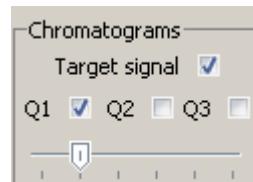
Method	C:\msdchem\MSDemo\rfgdata\gas04.m		
Calibrated	2013-08-28 20:53		
	Find com		
Data	Chromatograms	Log	Column config.
	Sample	Sample	
	File	Acquired	Calibrated
1	rfgqc01	1996-03-20 11:10	2013-08-28 20:53
2	rfgsamp	1996-03-18 21:15	2013-08-28 20:53

Review integration

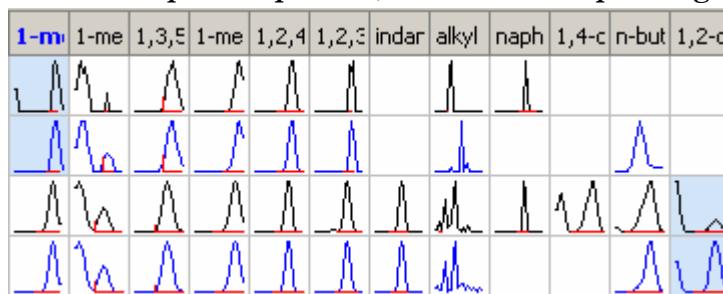
1. Review integration, click on the Chromatogram tab



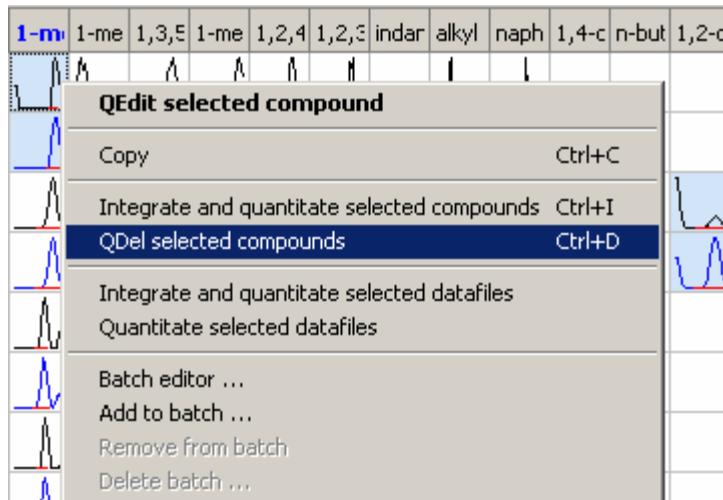
2. Select the signals and resize the plots



3. Select multiple compounds, left click while pressing the control key



4. Delete false peaks



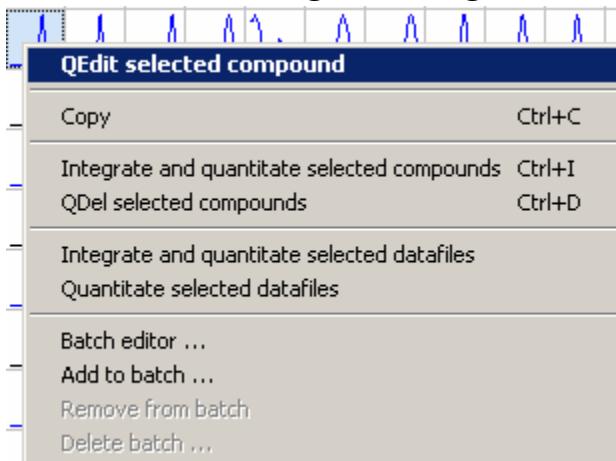
QEdit

1. Note the Response and MI value for o-xylene of std_a1

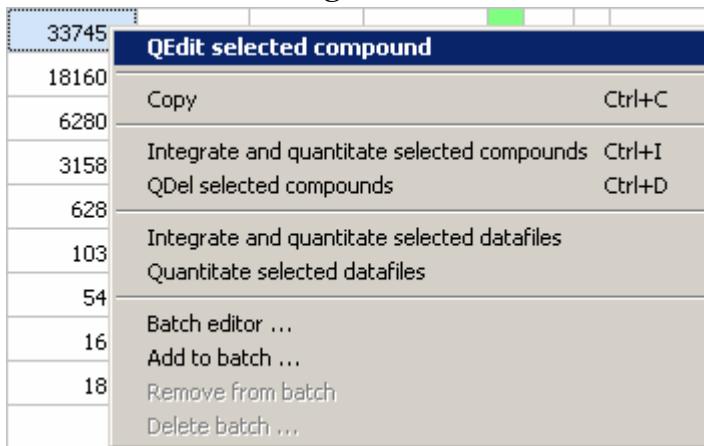
o-xylene						
Response	Final	ExpConc	Accuracy	Q	MI	D
2155357	2327.603			2		
2078636	2111.687			2		
3374554	5871.171	6000.000	97.85	2		
1816025	3218.328	3000.000	107.28	2		

2. Start QEdit

- a. QEdit from Chromatogram tab, right click menu, or double click on a plot



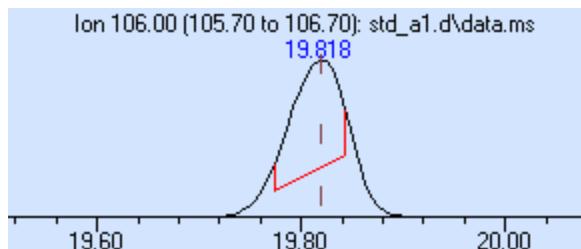
- b. QEdit from Data tab, right click menu, or double click on a compound cell



- c. QEdit with navigation arrow



3. Perform a manual integration on o-xylene of std_a1 in DataAnalysis, using the right mouse-button



4. Save QEdit in DataAnalysis, or click on navigation buttons or Save button in QuantBrowser

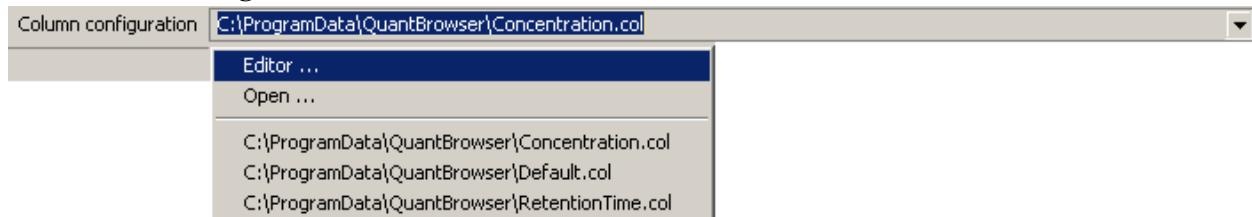


5. Note the new values for o-xylene of std_a1.

o-xylene						
Response	Final	ExpConc	Accuracy	Q	MI	D
2155357	2327.603			2		
2078636	2111.687			2		
1738070	3023.957	6000.000	50.40	1	1	
1816025	3218.328	3000.000	107.28	2		

Column configuration

1. New column configurations can be created



2. Column can be selected and re-order by drag and drop.

Sample columns are green, compound columns are red.

The screenshot shows the "Column set editor" dialog box. The title bar says "Column set editor" and the path "C:\ProgramData\QuantBrowser\Concentration.col" is in the title bar. The main area is a grid table with the following columns:

Column	Title	Format	Fixed columns	Data columns	Chrom columns	Export columns
DataFile	File	%s	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>
AcqTime	Acquired	yyyy-mm-dd hh:nn	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
AcqSequence	Sequence	%s	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
AcqMethod	AcqMethod	%s	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Folder	Folder	%s	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Multiplier	Multiplier	%,.4f	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
SampleAmount	Amount	%,.4f	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
QuantMethod	QntMethod	%s	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Calibrated	Calibrated	yyyy-mm-dd hh:nn	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
SampleName	Name	%s	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>
SampleInfo	Info	%s	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
SampleVial	Vial	%d	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
SampleType	Type	%s	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
CalibLevel	Level	%s	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
SampleReject	Reject	%d	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
SampleComment	Comment	%s	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
CompoundName	Compound	%s	<input type="checkbox"/>		<input checked="" type="checkbox"/>	
ExpRetentionTime	ExpRT	%.3f	<input type="checkbox"/>		<input type="checkbox"/>	
CorrExpRetentionTime	CorrExpRT	%.3f	<input type="checkbox"/>		<input type="checkbox"/>	
RetentionTime	RT	%.3f	<input type="checkbox"/>		<input type="checkbox"/>	
PercentDeltaRetentionTime	%Delta RT	%.3f	<input type="checkbox"/>		<input type="checkbox"/>	
AccuracyRetentionTime	Accuracy RT	%.3f	<input type="checkbox"/>		<input type="checkbox"/>	
RelativeRetentionTime	RRT	%.4f	<input type="checkbox"/>		<input type="checkbox"/>	
Index	Index	%d	<input type="checkbox"/>		<input type="checkbox"/>	
IsIstd	IsIstd	%d	<input type="checkbox"/>		<input type="checkbox"/>	
Istd	Istd	%d	<input type="checkbox"/>		<input type="checkbox"/>	
Response	Response	%.0f	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
IstdResponse	IResp	%.0f	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
RelativeResponse	RR	%.6f	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	

At the bottom of the dialog box, there is a note "Drag and drop rows to re-order" and buttons for "Open ...", "Save as ...", and "Close".

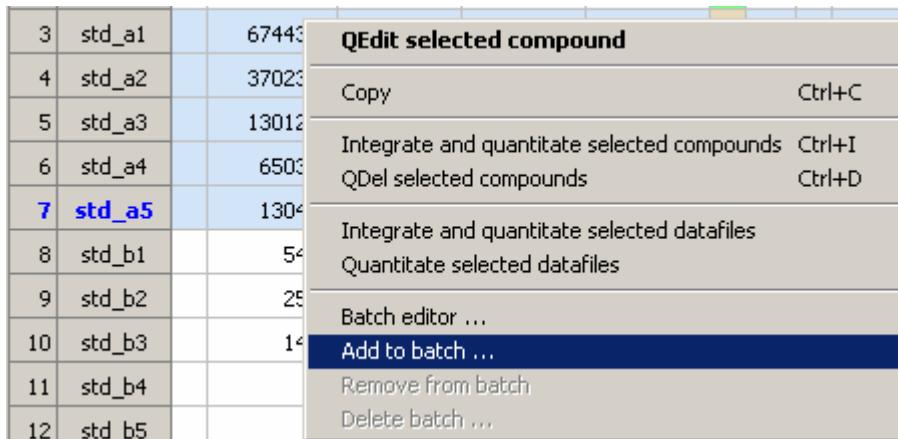
Column information

1. Rest the mouse above a column header

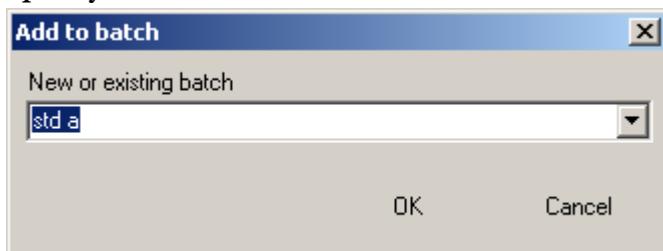
1-methylethylbenzene					
Q	MI	D	Response	Final	ExpConc
2	2	Integration			
2	2	0: By method			
0	1	1: Manually			
0	1	2: Corresponding ISTD was manual integrated			
8	2	3: Both this and ISTD manual integrated			
		5119	6.276		

Sample batch

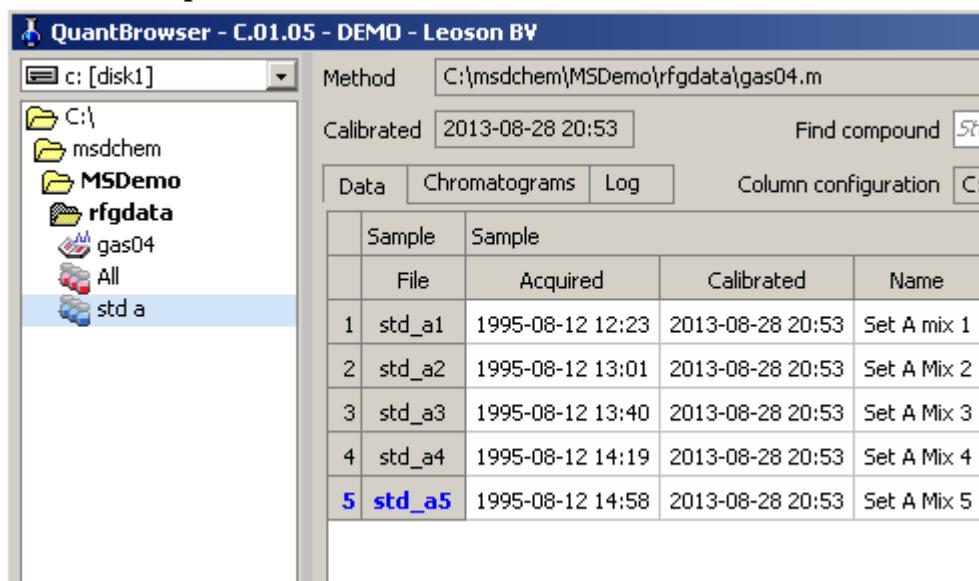
- Select sample for the batch, right click on the samples and Add to batch



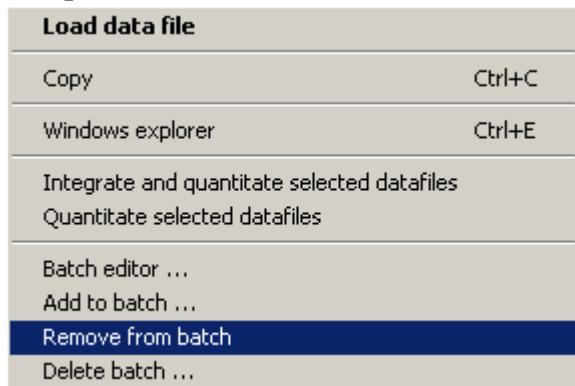
- Specify the batch name



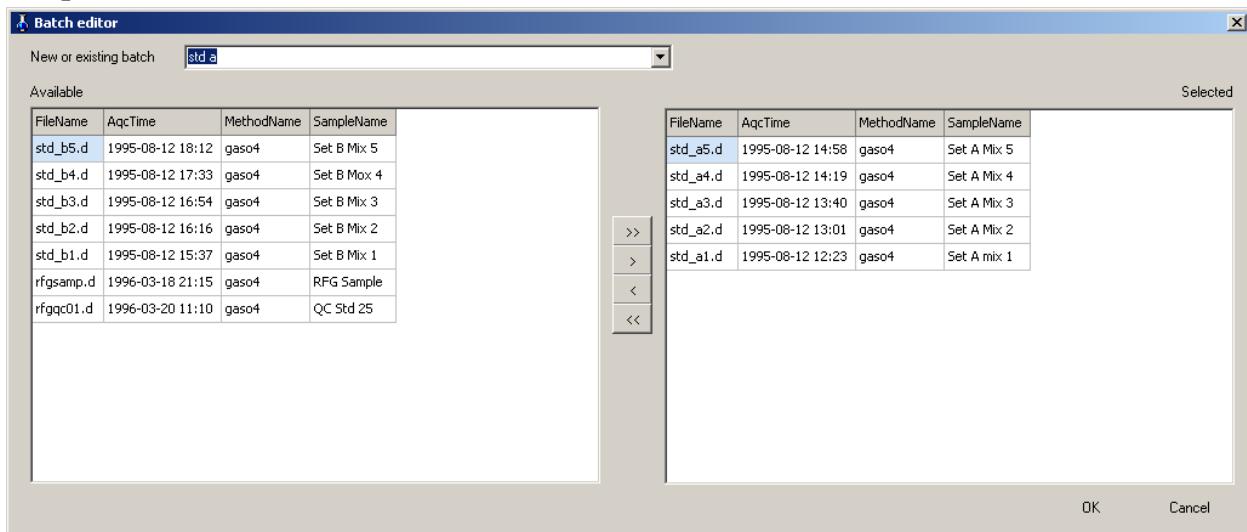
- Find the samples in the new batch



4. Samples can be removed from the batch



5. The batch editor helps to add or remove samples, click on the column header to sort the samples

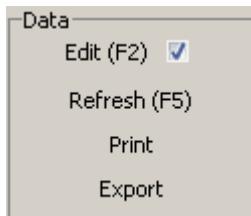


Copy to clipboard

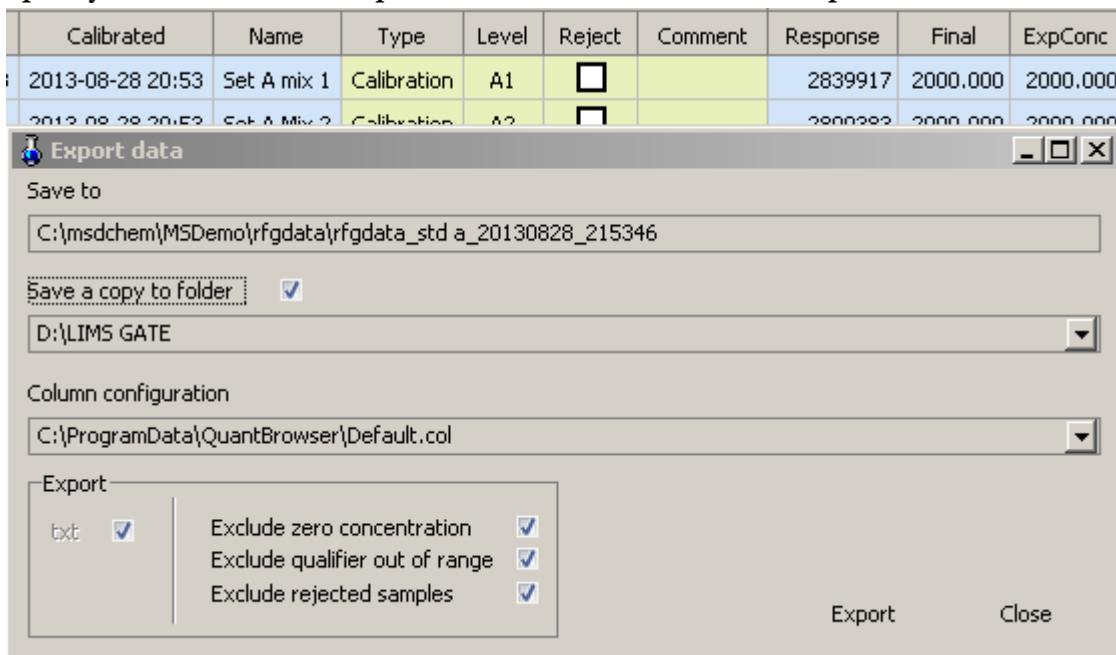
1. Select table cells and copy to clipboard. Use the right click menu in the table.

Export data

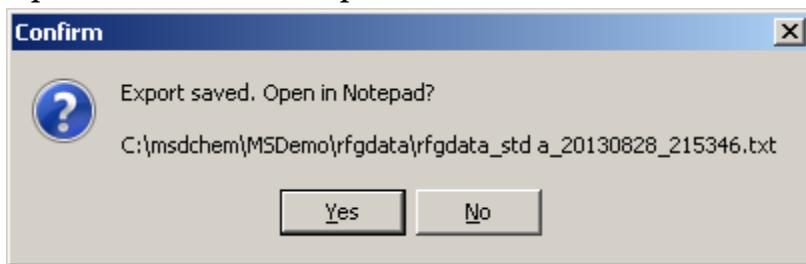
1. Open the export dialog, click on export button



2. Specify the folder for the export file and the columns to be exported



3. Open the results in Notepad



4. View the results

rfgdata_std a_20130828_215346.txt - Notepad		
File Edit Format View Help		
Name	Compound	Final
Set A mix 1	benzene-d6	2000.000
Set A mix 1	benzene	2947.396
Set A mix 1	ethylbenzene-d10	2000.000
Set A mix 1	Toluene	13917.123
Set A mix 1	ethylbenzene	4904.068
Set A mix 1	1-methylethylbenzene	12.769
Set A mix 1	1-methyl-3-ethylbenzene	3.829
Set A Mix 2	benzene-d6	2000.000
Set A Mix 2	benzene	1527.785
Set A Mix 2	ethylbenzene-d10	2000.000
Set A Mix 2	Toluene	10753.207
Set A Mix 2	ethylbenzene	2660.650
Set A Mix 2	o-xylene	3218.328
Set A Mix 2	1-methylethylbenzene	6.276
Set A Mix 2	1-methyl-3-ethylbenzene	2.008
Set A Mix 3	benzene-d6	2000.000
Set A Mix 3	benzene	1050.842
Set A Mix 3	ethylbenzene-d10	2000.000
Set A Mix 3	Toluene	6135.020
Set A Mix 2	ethylbenzene	1061.107