



Clarity™

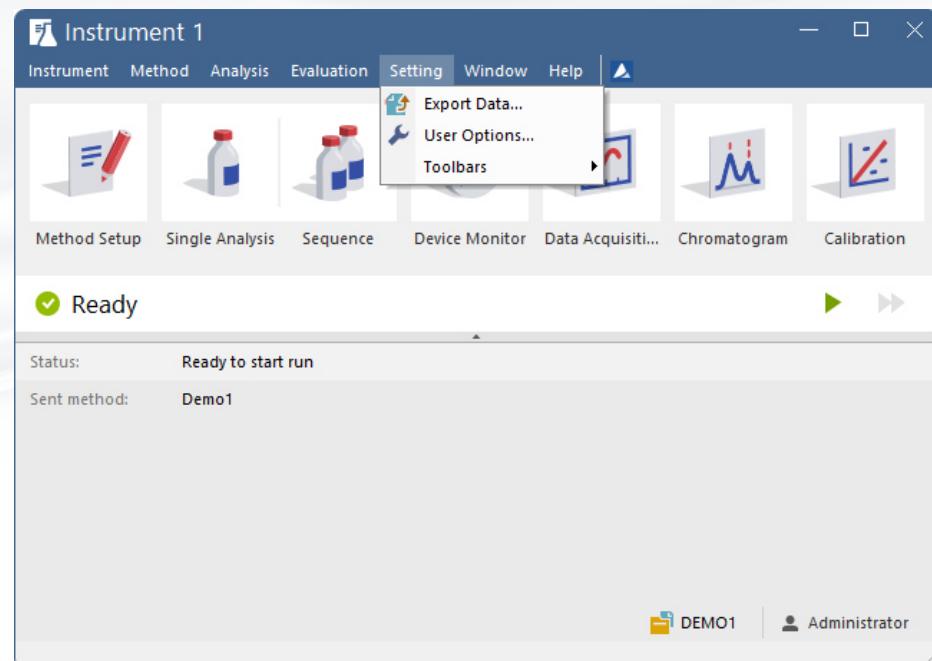
ADVANCED CHROMATOGRAPHY SOFTWARE

CLARITY TRAINING

ADVANCED – PART 1

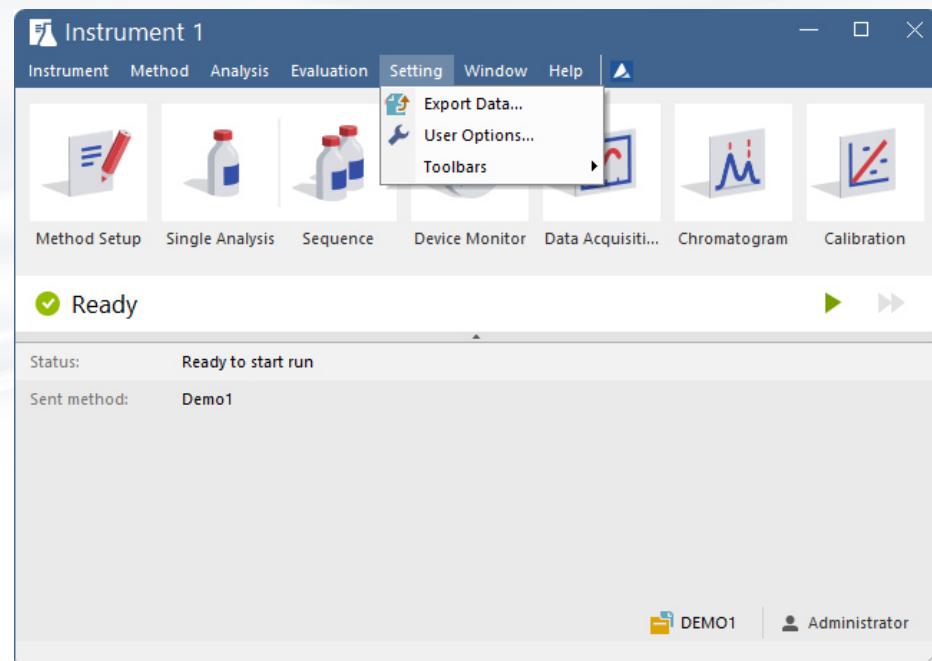
AGENDA

- Instrument window
 - User Options
- Single Analysis
 - Injected volume
 - Autosampler use
- Sequence table
 - Sequence Mode
 - Options
 - Import, Fill series



AGENDA

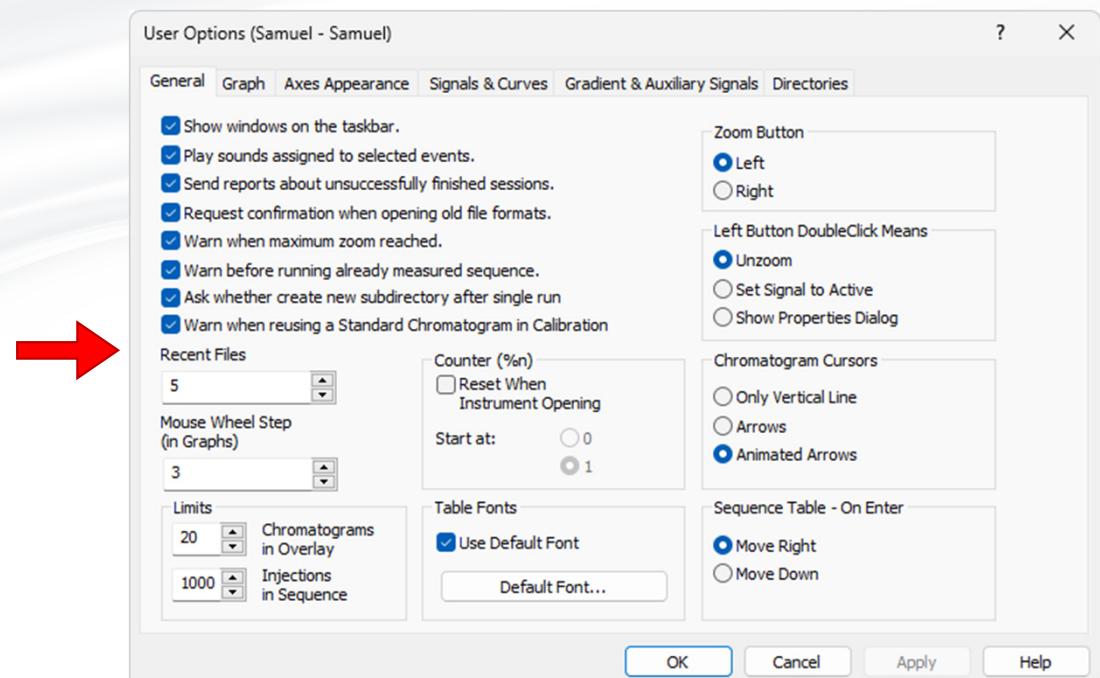
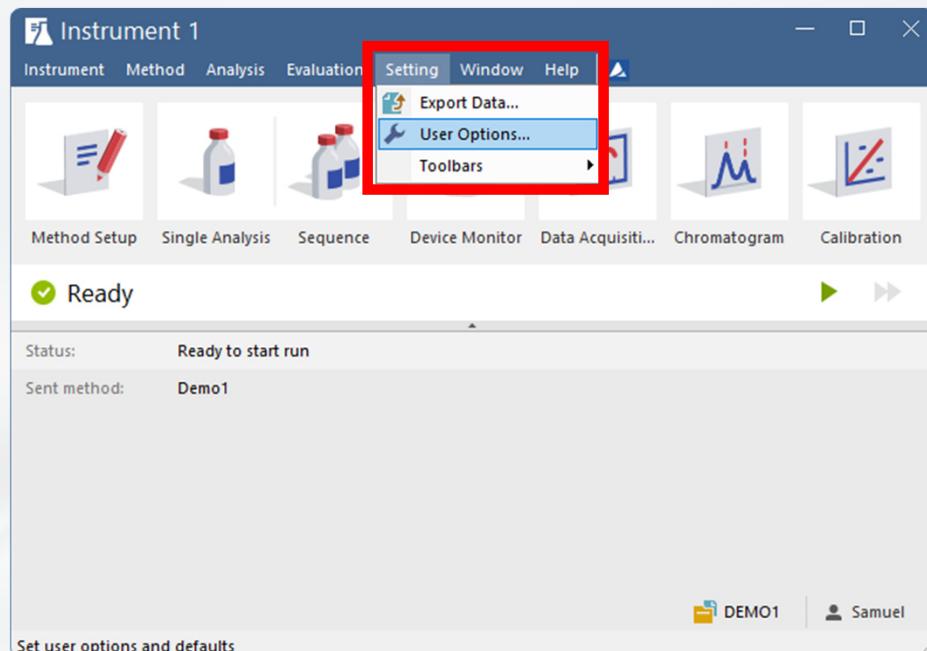
- Instrument window
 - User Options
- Single Analysis
 - Injected volume
 - Autosampler use
- Sequence table
 - Sequence Mode
 - Options
 - Import, Fill series



INSTRUMENT WINDOW

USER OPTIONS

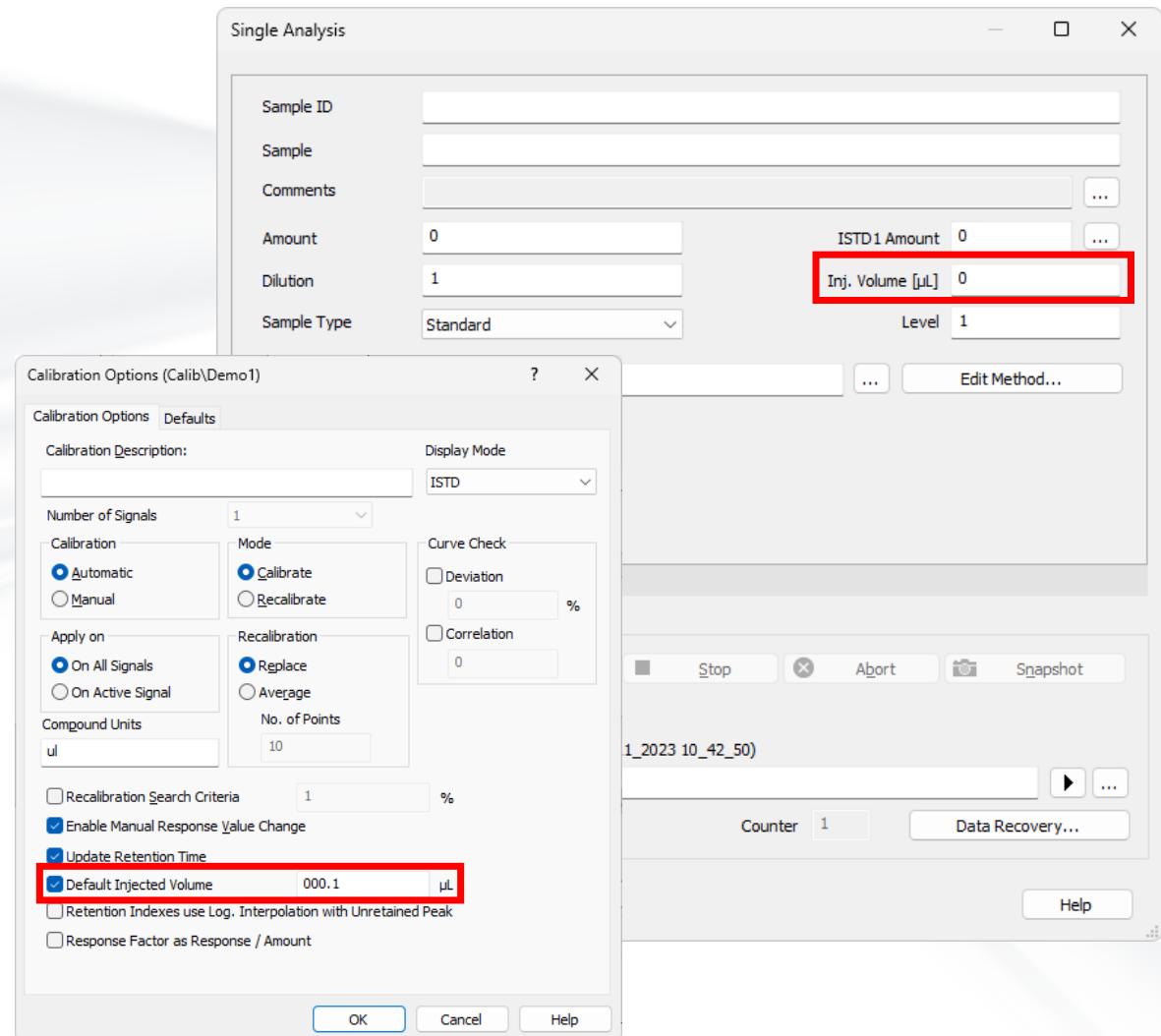
- Stored in desktop *.dsk files = can be set individually for each user
- Shared desktop file to share settings



SINGLE ANALYSIS

INJECTED VOLUME

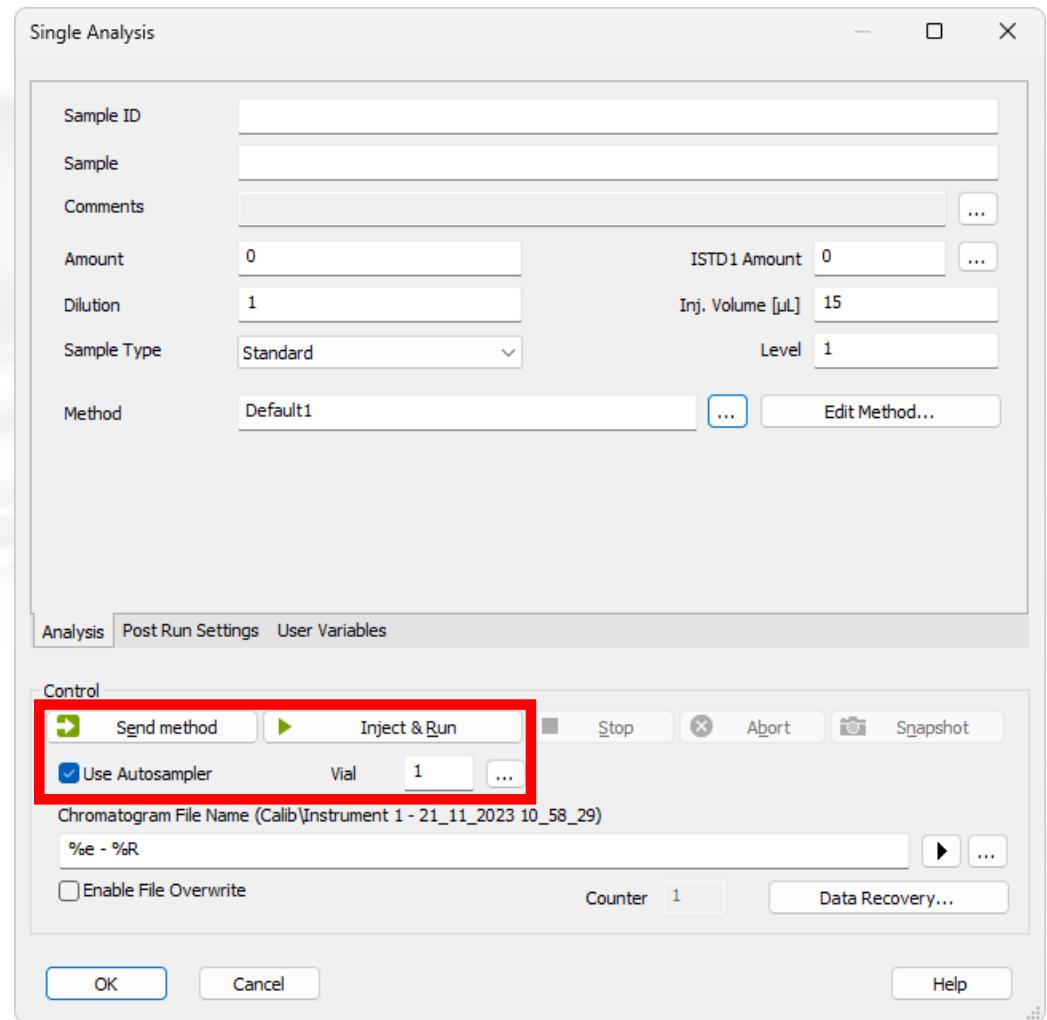
- **Linear** detector response is considered
- First, set Default Injected Volume in Calibration Options
- In Single Analysis, set the Injection volume for the sample and standard
- The responses are corrected by the ratio of Inj. Volume and Default Injected Volume in recalibration or calibration calculations



SINGLE ANALYSIS

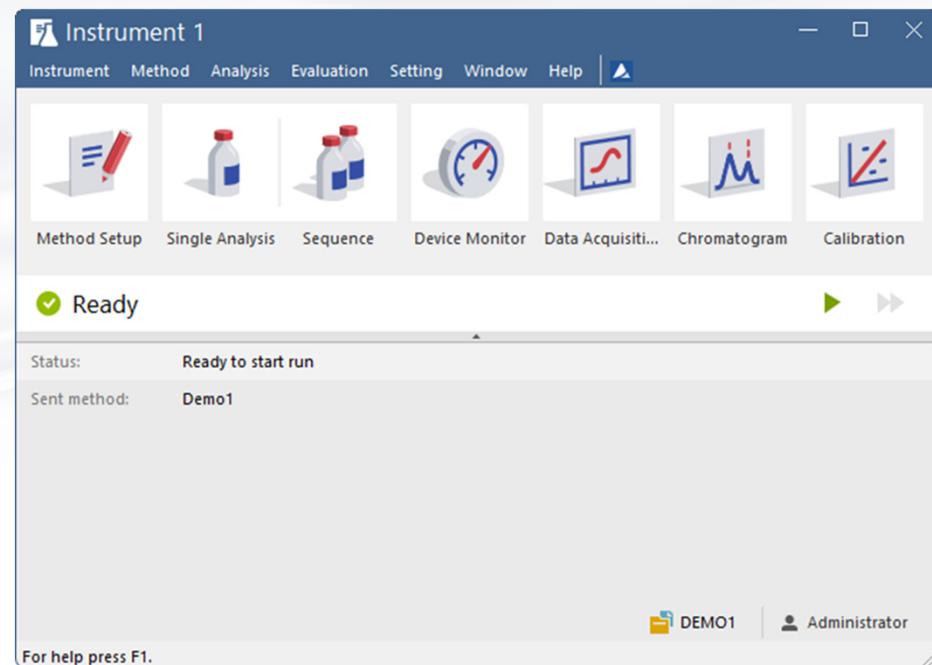
AUTOSAMPLER USE

- **Inject & Run** performs one injection from the specified vial and starts the acquisition
- Vial can be selected via a graphic interface if supported by the AS
- The **Use Autosampler** checkbox is present only if AS is configured



AGENDA

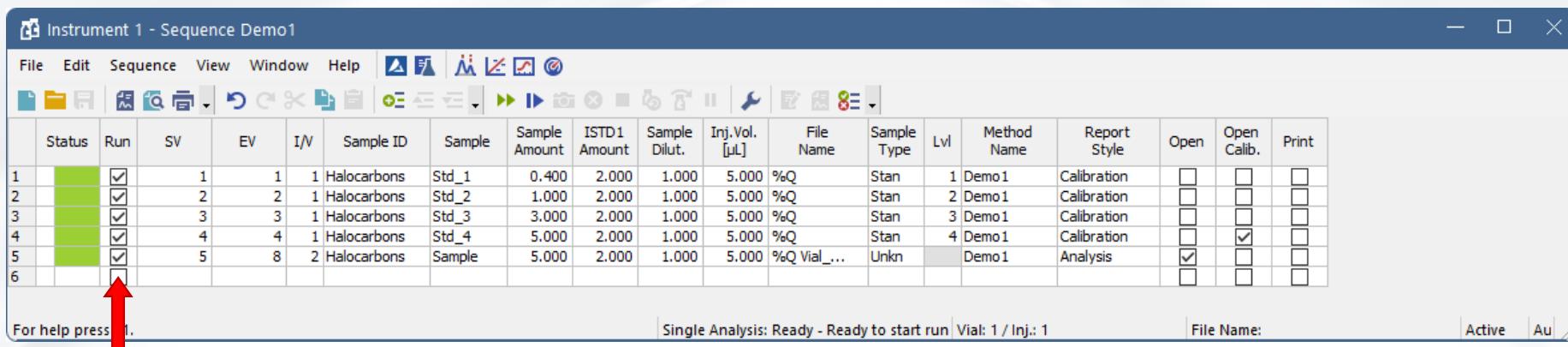
- Instrument window
 - User Options
- Single Analysis
 - Injected volume
 - Autosampler use
- Sequence
 - Description
 - Options and functions overview
 - Import, Fill series, Select vial



SEQUENCE TABLE OVERVIEW

ADD SAMPLES

- Samples are added by filling the **last empty row**
 - When any field is filled, the other fields are **copied** from the previous row
- Copy & Paste** one or more rows to add samples
- Import of sample data from **text files**
- Reprocessing** of the last measured sample set

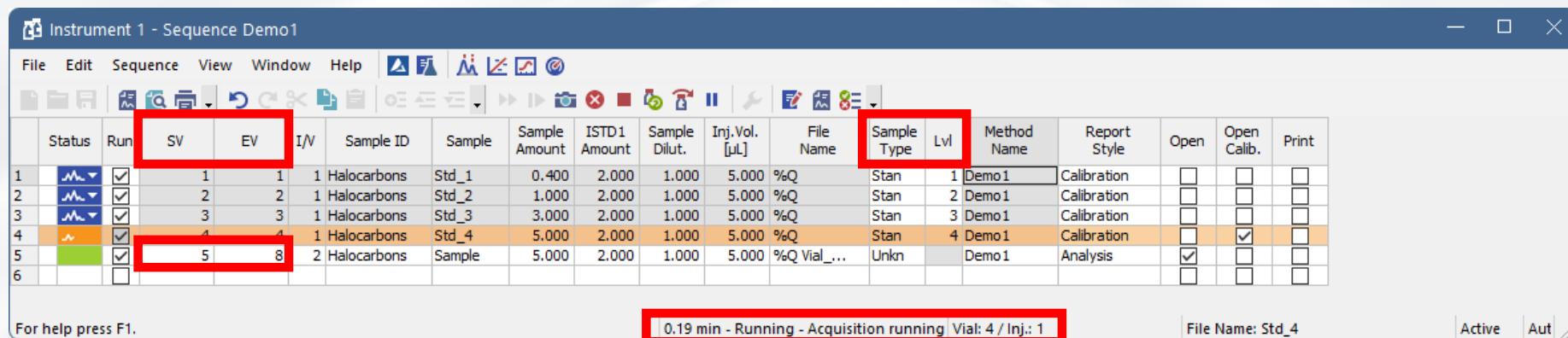


	Status	Run	SV	EV	I/V	Sample ID	Sample	Sample Amount	ISTD1 Amount	Sample Dilut.	Inj. Vol. [µL]	File Name	Sample Type	Lvl	Method Name	Report Style	Open	Open Calib.	Print
1				1	1	1 Halocarbons	Std_1	0.400	2.000	1.000	5.000	%Q	Stan	1	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
2				2	2	1 Halocarbons	Std_2	1.000	2.000	1.000	5.000	%Q	Stan	2	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
3				3	3	1 Halocarbons	Std_3	3.000	2.000	1.000	5.000	%Q	Stan	3	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
4				4	4	1 Halocarbons	Std_4	5.000	2.000	1.000	5.000	%Q	Stan	4	Demo1	Calibration	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>
5				5	8	2 Halocarbons	Sample	5.000	2.000	1.000	5.000	%Q	Vial_...	Unkn	Demo1	Analysis	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
6																	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

SEQUENCE WINDOW

DESCRIPTION

- **SV, EV** = start and end vial injection number
 - A single row can define **several** vials/injections
- **Sample Type, Lvl** = can be used for automatic **recalibration** via sequence
- Window bottom shows the **state** of the sequence (runtime, vial, etc.)



The screenshot shows the DataApex Sequence Demo1 software interface. The main window displays a sequence table with the following data:

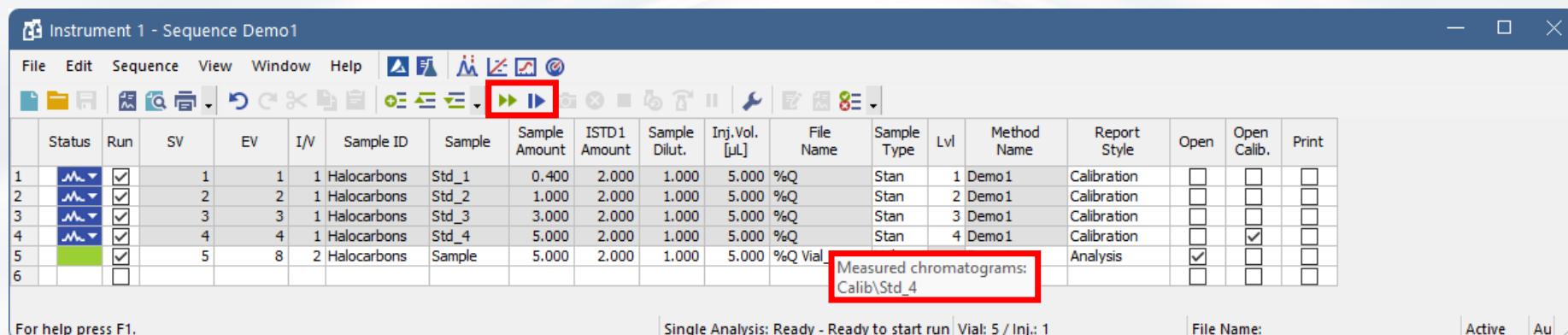
	Status	Run	SV	EV	I/V	Sample ID	Sample	Sample Amount	ISTD1 Amount	Sample Dilut.	Inj. Vol. [µL]	File Name	Sample Type	Lvl	Method Name	Report Style	Open	Open Calib.	Print		
1	W	✓			1	1	1	0.400	2.000	1.000	5.000	%Q	Stan	1	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>		
2	W	✓			2	2	1	Halocarbons	Std_2	1.000	2.000	1.000	5.000	%Q	Stan	2	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
3	W	✓			3	3	1	Halocarbons	Std_3	3.000	2.000	1.000	5.000	%Q	Stan	3	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
4	W	✓			4	4	1	Halocarbons	Std_4	5.000	2.000	1.000	5.000	%Q	Stan	4	Demo1	Calibration	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>
5	W	✓			5	8	2	Halocarbons	Sample	5.000	2.000	1.000	5.000	%Q	Vial_...	Unkn	Demo1	Analysis	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
6																		<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	

For help press F1. 0.19 min - Running - Acquisition running Vial: 4 / Inj: 1 File Name: Std_4 Active Aut

SEQUENCE WINDOW

DESCRIPTION

- **Run** sequence using Ctrl+Q or the  icon
 - Running sequence will lead to a reset of all rows (indicated by a message dialog)
- **Resume** sequence Ctrl+Space or the  icon
 - Continue from the first unmeasured row with the Run option checked
- Tooltips display additional details (e.g., actual file names of measured chromatograms, etc.)



Instrument 1 - Sequence Demo1

	Status	Run	SV	EV	I/V	Sample ID	Sample	Sample Amount	ISTD1 Amount	Sample Dilut.	Inj. Vol. [µL]	File Name	Sample Type	Lvl	Method Name	Report Style	Open	Open Calib.	Print
1		<input checked="" type="checkbox"/>		1	1	1 Halocarbons	Std_1	0.400	2.000	1.000	5.000 %Q	Stan	1	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
2		<input checked="" type="checkbox"/>		2	2	1 Halocarbons	Std_2	1.000	2.000	1.000	5.000 %Q	Stan	2	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
3		<input checked="" type="checkbox"/>		3	3	1 Halocarbons	Std_3	3.000	2.000	1.000	5.000 %Q	Stan	3	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
4		<input checked="" type="checkbox"/>		4	4	1 Halocarbons	Std_4	5.000	2.000	1.000	5.000 %Q	Stan	4	Demo1	Calibration	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
5		<input checked="" type="checkbox"/>		5	8	2 Halocarbons	Sample	5.000	2.000	1.000	5.000 %Q Vial				Analysis	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
6		<input type="checkbox"/>														<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	

For help press F1. Single Analysis: Ready - Ready to start run Vial: 5 / Inj.: 1 File Name: Active Au

Measured chromatograms: Calib\Std_4

SEQUENCE WINDOW

BASIC FUNCTIONS

- **Snapshot** : create a temporary snapshot (will be replaced by the complete chromatogram when measured)
- **Abort** sequence : terminate the sequence without saving the current chromatogram
- **Stop** sequence : 1st click halts the sequence after the current measurement, 2nd click stops analysis immediately
- **Repeat injection** : interrupt the actual measurement and repeat injection
- **Skip vial**
- **Check sequence** for errors

	Status	Run	SV	EV	I/V	Sample ID	Sample	Sample Amount	ISTD1 Amount	Sample Dilut.	Inj. Vol. [µL]	File Name	Sample Type	Lvl	Method Name	Report Style	Open	Open Calib.	Print
1		<input checked="" type="checkbox"/>		1	1	1 Halocarbons	Std_1	0.400	2.000	1.000	5.000	%Q	Stan	1	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
2		<input checked="" type="checkbox"/>		2	2	1 Halocarbons	Std_2	1.000	2.000	1.000	5.000	%Q	Stan	2	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
3		<input checked="" type="checkbox"/>		3	3	1 Halocarbons	Std_3	3.000	2.000	1.000	5.000	%Q	Stan	3	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
4		<input checked="" type="checkbox"/>		4	4	1 Halocarbons	Std_4	5.000	2.000	1.000	5.000	%Q	Stan	4	Demo1	Calibration	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>
5		<input checked="" type="checkbox"/>		5	8	2 Halocarbons	Sample	5.000	2.000	1.000	5.000	%Q	Vial_...	Unkn	Demo1	Analysis	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
6		<input type="checkbox"/>														<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	

SEQUENCE WINDOW

BASIC FUNCTIONS

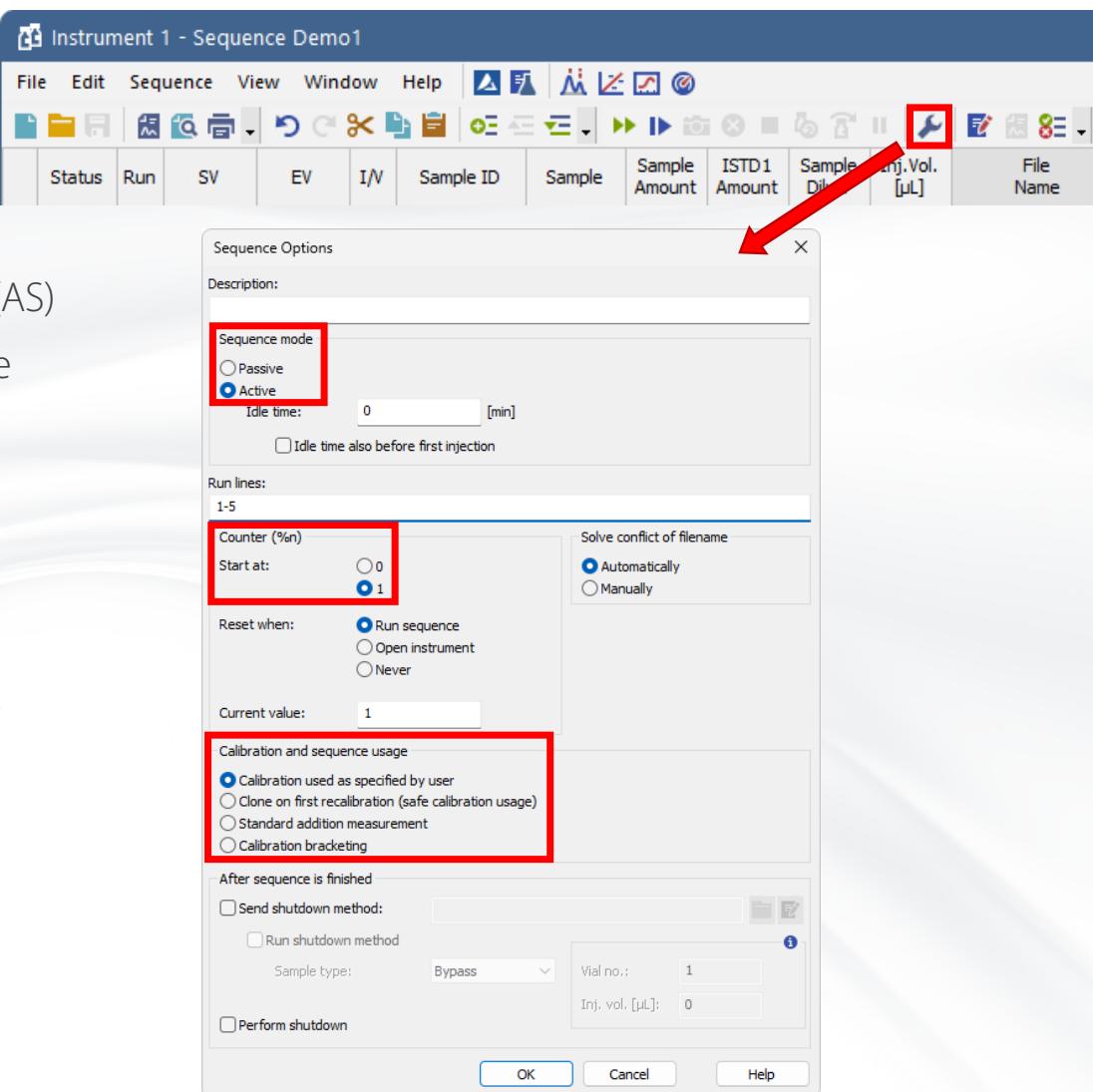
- Useful **file name variables**: %s = sequence, %J = Method, %P = project, %Q = Sample
- **Bypass** sample type = run without injection (for startup and shutdown methods)
- **Include in SST** – perform SST test when injection is finished
- **Stored Calib.** – chromatogram will be opened with stored calibration
- **Close All** – close all currently opened chromatograms in overlay

	Status	Run	SV	EV	I/V	Sample ID	Sample	Sample Amount	ISTD1 Amount	Sample Dilut.	Inj. Vol. [µL]	File Name	Sample Type	Lvl	Method Name	Report Style	Open	Open Calib.	Print	Include in SST	Stored Calib.	Close All
1				1	1	1 Bypass		0.000	0.000	1.000	0.000	%s %J %P %Q	Bypass		Demo1		<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
2				1	1	1 Halocarbons	Std_1	0.400	2.000	1.000	5.000	%s %J %P %Q	Standar	1	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
3				2	2	1 Halocarbons	Std_2	1.000	2.000	1.000	5.000	%s %J %P %Q	Standar	2	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
4				3	3	1 Halocarbons	Std_3	3.000	2.000	1.000	5.000	%s %J %P %Q	Standar	3	Demo1	Calibration	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
5				4	4	1 Halocarbons	Std_4	5.000	2.000	1.000	5.000	%s %J %P %Q	Standar	4	Demo1	Calibration	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
6				5	8	2 Halocarbons	Sample	5.000	2.000	1.000	5.000	%Q Vial_%2v-%i	Unknow		Demo1	Analysis	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>
7																	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

SEQUENCE WINDOW

SEQUENCE MODES AND OPTIONS

- **Passive sequence:** Clarity only expects the START signal, the time of analysis is set by the autosampler (AS)
- **Active sequence:** Clarity sends a READY signal to the AS and waits for the START signal
 - **AS control** in active sequence: AS injects according to SV, EX, I/V, and Inj. Volume in the Sequence Table
- Set the initial value for the **%on** variable
- Choose the **calibration mode** applied for the whole sequence



SEQUENCE WINDOW

IMPORT

- Import Sequence Step 1 dialog: select files to import
- Import Sequence Step 2 dialog: assign the respective fields to be imported
 - the Start Vial (SV) and File Name (both in bold) are mandatory

Import Sequence Step 1

File Name	C:\Clarity\DataFiles\DEMO1\Sequence.txt		...																																																		
Delimiter	<TAB>	Decimal Delimiter	. <DOT>																																																		
Preview:	<input checked="" type="checkbox"/> First Row Is Header																																																				
<table border="1"><thead><tr><th>1</th><th>Run</th><th>SV</th><th>EV</th><th>I/V</th><th>Sample ID</th><th>Sample</th><th>Sample Amount</th><th>ISTD1 Amount</th><th>Sample</th></tr></thead><tbody><tr><td>2</td><td>1</td><td>1</td><td>1</td><td>1</td><td>Halocarbons</td><td>Std_1</td><td>0.400</td><td>2.000</td><td></td></tr><tr><td>3</td><td>1</td><td>2</td><td>2</td><td>1</td><td>Halocarbons</td><td>Std_2</td><td>1.000</td><td>2.000</td><td></td></tr><tr><td>4</td><td>1</td><td>3</td><td>3</td><td>1</td><td>Halocarbons</td><td>Std_3</td><td>3.000</td><td>2.000</td><td></td></tr><tr><td>5</td><td>1</td><td>4</td><td>4</td><td>1</td><td>Halocarbons</td><td>Std_4</td><td>5.000</td><td>2.000</td><td></td></tr></tbody></table>				1	Run	SV	EV	I/V	Sample ID	Sample	Sample Amount	ISTD1 Amount	Sample	2	1	1	1	1	Halocarbons	Std_1	0.400	2.000		3	1	2	2	1	Halocarbons	Std_2	1.000	2.000		4	1	3	3	1	Halocarbons	Std_3	3.000	2.000		5	1	4	4	1	Halocarbons	Std_4	5.000	2.000	
1	Run	SV	EV	I/V	Sample ID	Sample	Sample Amount	ISTD1 Amount	Sample																																												
2	1	1	1	1	Halocarbons	Std_1	0.400	2.000																																													
3	1	2	2	1	Halocarbons	Std_2	1.000	2.000																																													
4	1	3	3	1	Halocarbons	Std_3	3.000	2.000																																													
5	1	4	4	1	Halocarbons	Std_4	5.000	2.000																																													
< Back		Next >																																																			
Cancel		Finish																																																			

Import Sequence Step 2

Column Matching:

Sequence Column	Imported Column
SV	SV
EV	EV
I/V	I/V
Sample ID	Sample ID
Sample	Sample
Comments	
Sample Amount	Sample Amount
ISTD1 Amount	ISTD1 Amount
ISTD2 Amount	
ISTD3 Amount	
ISTD4 Amount	
ISTD5 Amount	
ISTD6 Amount	
ISTD7 Amount	

Sequence

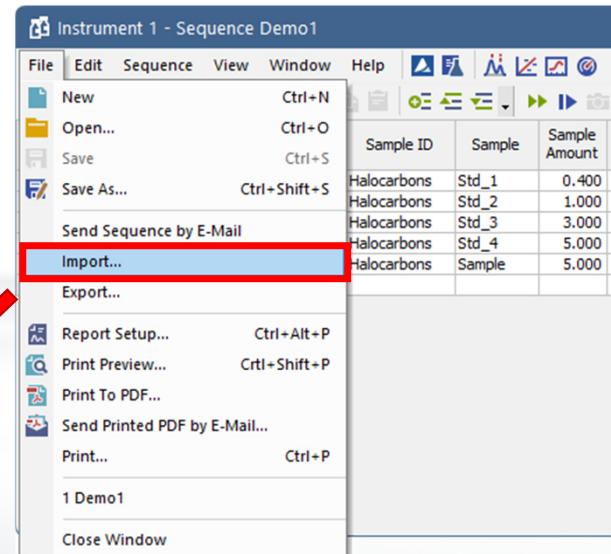
Create New
 Append to Existing

Save Sequence and Delete Import File

Save Import Settings

Show Sequence Options

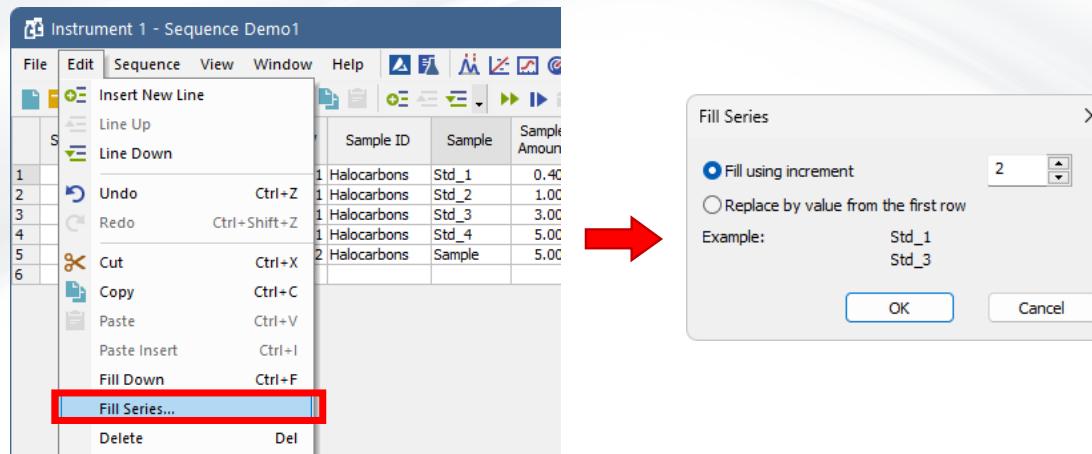
< Back Next > Cancel Finish



SEQUENCE WINDOW

FILL SERIES AND FILL DOWN

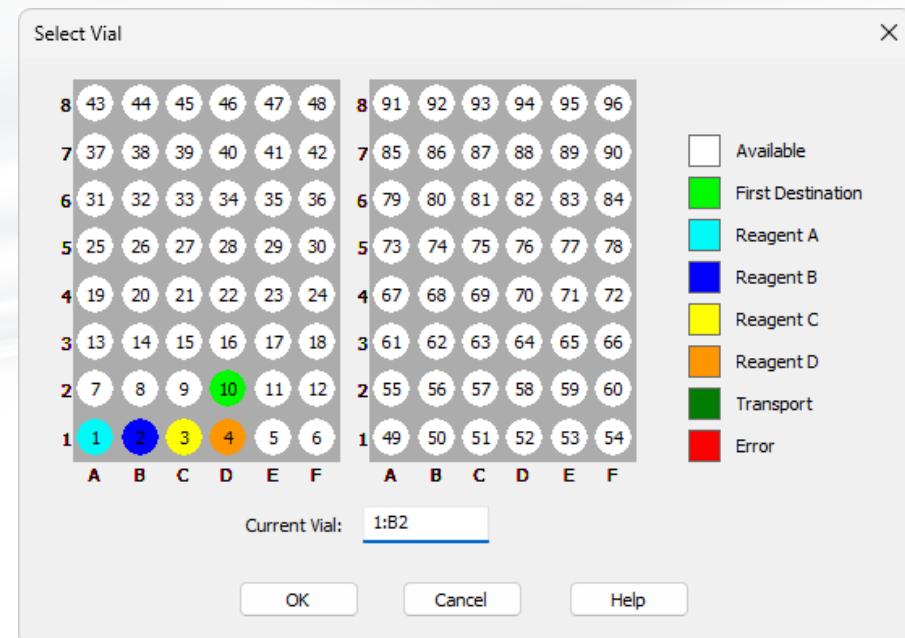
- Available in Sequence and Calibration
- Select a cell to be used for filling (numeric inputs, text inputs with numeric counters)
- Fill down (Ctrl+F):
 - Fill the cells below with the parameter in the selected cell
- Fill series:
 - Open the Fill Series dialog and specify the fill/replace parameters



SEQUENCE WITH AUTOSAMPLERS

SELECT VIAL

- The dialog with a graphic interface can be opened from Single Analysis/Sequence or viewed in Method Setup
- Convenient for working with **autosamplers** (available only in some AS)
- For controlled autosamplers, the **vial number** and **injection volume** are set in this dialog
- Allows **mid-run modifications** for not yet processed samples



23.0331



Clarity™

ADVANCED CHROMATOGRAPHY SOFTWARE



SHORT BREAK