



1.7 Setting up a Batch (2/3) – Data Analysis Method

Std. Familiarization Time: 0:10:00
Approximate Video Time: 0:06:30

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MassHunter Basics for Agilent ICP-MS
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Startup

- [Data Analysis Method]
 - [FullQuant Analysis]
 - [SemiQuant Analysis]
 - [Analysis Mode]

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Analyte List

- [Delete]
- [Load List from Acquisition Method]
- Set elements for [ISTD]

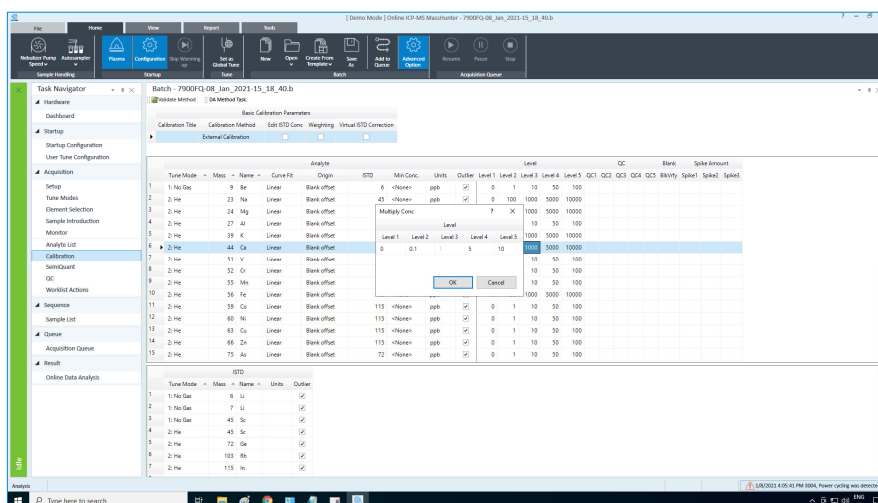
Calibration (1/3)

- [Curve Fit]
- [Origin]
- [ISTD]
- [Units]
- [Level]

Calibration (2/3)

- [Conc Multiply]
- [Add Levels]
- [Advanced Info] (In MassHunter 5.1, perform operations from Task Navigator)
 - [FullQuant Outlier]
 - [Worklist Actions]

Calibration (3/3)



The screenshot displays the 'Basic Calibration Parameters' dialog box in the MassHunter software. The 'Calibration Method' tab is selected, showing a table of calibration levels and a table of acquisition parameters.

Level	Level 1	Level 2	Level 3	Level 4	Level 5	QC1	QC2	QC3	QC4	QC5	Blank	Spike Amount
1	0	1	10	50	100							
2	0	1	10	50	100							
3	0	1	10	50	100							
4	0	1	10	50	100							
5	0	1	10	50	100							
6	0	1	10	50	100							
7	0	1	10	50	100							
8	0	1	10	50	100							
9	0	1	10	50	100							
10	0	1	10	50	100							
11	0	1	10	50	100							
12	0	1	10	50	100							
13	0	1	10	50	100							
14	0	1	10	50	100							
15	0	1	10	50	100							

The 'Acquisition' tab is also visible, showing a table of acquisition parameters:

Level	Level 1	Level 2	Level 3	Level 4	Level 5	QC1	QC2	QC3	QC4	QC5	Blank	Spike Amount
1	0	1	10	50	100							
2	0	1	10	50	100							
3	0	1	10	50	100							
4	0	1	10	50	100							
5	0	1	10	50	100							
6	0	1	10	50	100							
7	0	1	10	50	100							
8	0	1	10	50	100							
9	0	1	10	50	100							
10	0	1	10	50	100							
11	0	1	10	50	100							
12	0	1	10	50	100							
13	0	1	10	50	100							
14	0	1	10	50	100							
15	0	1	10	50	100							