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Application Note

Automated Creation of Chromatographic Methods for Analysis with an ACQUITY™ QDa Detector Using Empower™ Sample Set Generator

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This is an Application Brief and does not contain a detailed Experimental section.

Abstract

Optimizing ion source-mass spectrometry (MS) parameters to enhance signal requires creation of multiple chromatographic methods over a range of ionization variables. This technology brief illustrates the use of an Empower™ Sample Set Generator Software to automatically create chromatographic methods for analysis with an ACQUITY QDa Mass Detector, to study the impact of probe temperature and cone voltage on the sensitivity of memantine hydrochloride.

Benefits

- Automated and quick creation of Empower instrument methods, methods sets, and sample set methods using
 Empower Sample Set Generator, while varying chromatographic parameters
- Generation of chromatographic methods for Waters ACQUITY LC instruments, optical detectors, and ACQUITY QDa Mass Detector

Introduction

Optimization of MS source ionization parameters plays an important role in obtaining high sensitivity analysis in liquid chromatography-mass spectrometry (LC-MS). The ionization parameters (e.g., ionization mode, cone voltage, or probe temperature) are often investigated to enhance the MS signal and to achieve low detection limits for target analytes. To study the impact of different ionization parameters and their ranges on the sensitivity requires careful creation of multiple chromatographic methods over set of variables. Manually creating and verifying these methods can be time-consuming and prone to errors process.

The Empower Sample Set Generator (SSG) capabilities automates the creation of instrument methods, method sets, and sample set methods, while varying the chromatographic parameters. The Empower method sets and instruments methods are automatically created and structured in the sample set method according to the experiment design as a ready-to-run injection sequence. Automating creation of chromatographic methods minimizes transcription errors that may arise during the manual process and time spent generating methods.

This technology brief illustrates the use of Empower SSG to automatically create chromatographic methods for analysis with an ACQUITY QDa Mass Detector coupled to an Arc™ Premier System. The ACQUITY QDa is used for the detection of non-chromophoric memantine hydrochloride, a drug commonly used to treat dementia often associated with Alzheimer's disease. ¹ The impact of QDa ionization parameters (probe temperature and cone voltage) on the method sensitivity for analysis of memantine is investigated.

Results and Discussion

Memantine hydrochloride lacks a chromophore required for UV detection but produces a robust MS signal using an ACQUITY QDa Mass Detector. ² Method conditions used in this work are summarized in Table 1. A standard solution at 10 ng/mL in 90:10 water/methanol was prepared from a concentrated stock solution containing 1 mg/mL in methanol and used to carry-out the sensitivity study with MS detection.

Method conditions for analysis of memantine hydrochloride

System: Arc Premier System, 2998 PDA and

ACQUITY QDa Column: CORTECS $^{\text{\tiny M}}$ C₁₈₊, 2.7 μ m, 3 mm x 75 mm (p/n: 186007401) Column temperature: 45 °C A: 0.1% Formic acid in water Mobile phase: B: 0.1% Formic acid in acetonitrile Injection volume: 3.0 µL MS detection: ACQUITY QDa Mass Detector (Extended Performance) •Ionization mode: ESI+ -Single ion recording (SIR): 180.2 Da •Probe temperature: 600 °C •Cone voltage: 15 V Capillary voltage: 0.8 kV Data: centroid

Wash solvents:

Purge/sample wash: 50:50

Seal wash: 90:10 water/acetonitrile

water/methanol

Gradient Table

| Step | Time | Flow rate (mL/min) | %A | %B | Curve |
|------|---------|-----------------------|------|------|-------|
| 1 | Initial | 1.00 | 95.0 | 5.0 | 6 |
| 2 | 2.5 | 1.00 | 10.0 | 90.0 | 6 |
| 3 | 3.5 | 1.00 | 10.0 | 90.0 | 6 |
| 4 | 3.6 | 1.00 | 95.0 | 5.0 | 6 |
| 5 | 6.0 | 1.00 | 95.0 | 5.0 | 6 |

Chromatographic methods were automatically created using the Empower SSG to study the impact of probe temperature (600, 500, and 400 °C) and cone voltage (5, 10, 15, and 20 Volts) on the sensitivity of memantine HCl following the steps described below.

- A comma separated value (CSV) file with the QDa variables was imported to the Empower SSG (Figure 1).
 The experimental design included a combination of MS conditions with different probe temperatures and cone voltages.
- · A base sample set method that included method set and instrument method with the system configuration for the analysis was loaded from the Empower project to Empower SSG.
- · Study factors probe temperature and cone voltage were associated with the Empower settings (Figure 2).
- · Final settings (injection panel, equilibration time, and method names) were defined before completing generation of the sample set method (Figure 3).

After completing the above steps, the Empower SSG automatically created a sample set method according to the experiment design as a ready-to-run injection sequence (Figure 4). The Empower instruments methods and method sets were automatically created and built into the sample set method. The sample set method included the experiment name and 12 method sets with different probe temperatures and cone voltages. An equilibration step and injections of blank were added at the beginning of the run as instructed by the user. Without the Empower SSG, a user would need to manually create 12 instrument methods and method sets, which is time consuming and prone to errors process. Using automated generation with Empower SSG reduced the time and transcription errors, providing confidence that all chromatographic runs were completed with correctly created methods. The Empower data acquired using ACQUITY QDa Detector is shown in Figure 5. The peak data across

the experimental runs indicated that the best sensitivity for memantine HCl was achieved with a probe temperature of 600°C and cone voltage of 15 Volts, resulting in highest signal-to-noise (s/n) value (Figure 5A).

A. Import the CSV file with the experimental design with variables for testing



B. Review the content of the experimental study

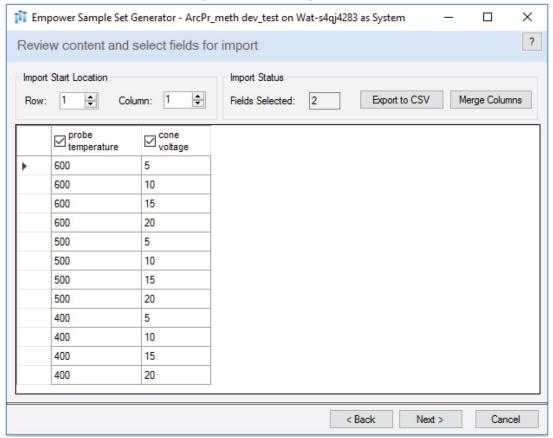


Figure 1. Creating chromatographic methods with Empower SSG. Import CSV file (A) and review experiment design (B).

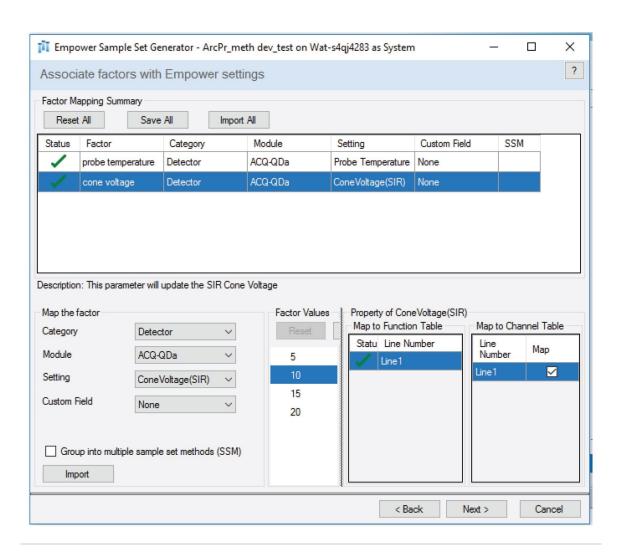


Figure 2. Empower SSG. Associate factors or instrument variables with Empower settings.

| The Empower Sample Set Generator - ArcPr_meth dev_test on Wat-s4qj4283 as System — — — X | | | | | | | | | |
|--|---|--------------------------------|---------------------------------|--------------------------------------|-------------------------|-----|--|--|--|
| Final Generation Settings ? | | | | | | | | | |
| Preparation Informat Number of Preparati 1 Number of Injections 1 | ons/Experiment | | | | | | | | |
| Injections | At the Beginning | On Instrument Method Change | On Preparation Factor Change | On Sample Concentration Change | Number of Injections | | | | |
| Blank | \square | | | | 2 | | | | |
| System Suitability1 | | | | | | | | | |
| System Suitability2 | | | | | | | | | |
| Standard | | | | | | | | | |
| Equilibration Run Tir | or will insert an Equilib me (Minutes) | | - | mistry between samp | le lines | | | | |
| | ethod, Method Set a | nd Instrument Method | Name | | | | | | |
| Sample Set Method | | | | | | | | | |
| Memantine_QDa | | | | | | | | | |
| Method Set Name Memantine_QDa | | | | | | | | | |
| Instrument Method Name | | | | | | | | | |
| Memantine_QDa | | | | | | | | | |
| | | | < | Back Genera | ate Can | cel | | | |

Figure 3. Empower SSG final settings for generation of methods.

| Ē | Plate /Well | # of Injs | SampleName | Function | Method Set / Report or Export Method | Probe Temp | Cone Voltage | Run Time (Minutes) |
|----|----------------|--------------|--------------|----------------|--|---------------|-----------------|--------------------------|
| 1 | | | | Equilibrate | Memantine_QDa_1 | | | 10.00 |
| 2 | 1:A,1 | 2 | Memantine_1 | Inject Samples | Memantine_QDa_1 | 600 | 5 | 6.50 |
| 3 | 1:A,2 | 1 | Memantine_1 | Inject Samples | Memantine_QDa_1 | 600 | 5 | 6.50 |
| 4 | 1:A,2 | 1 | Memantine_2 | Inject Samples | Memantine_QDa_2 | 600 | 10 | 6.50 |
| 5 | 1:A,2 | 1 | Memantine_3 | Inject Samples | Memantine_QDa_3 | 600 | 15 | 6.50 |
| 6 | 1:A,2 | 1 | Memantine_4 | Inject Samples | Memantine_QDa_4 | 600 | 20 | 6.50 |
| 7 | 1:A,2 | 1 | Memantine_5 | Inject Samples | Memantine_QDa_5 | 500 | 5 | 6.50 |
| 8 | 1:A,2 | 1 | Memantine_6 | Inject Samples | Memantine_QDa_6 | 500 | 10 | 6.50 |
| 9 | 1:A,2 | 1 | Memantine_7 | Inject Samples | Memantine_QDa_7 | 500 | 15 | 6.50 |
| 10 | 1:A,2 | 1 | Memantine_8 | Inject Samples | Memantine_QDa_8 | 500 | 20 | 6.50 |
| 11 | 1:A,2 | 1 | Memantine_9 | Inject Samples | Memantine_QDa_9 | 400 | 5 | 6.50 |
| 12 | 1:A,2 | 1 | Memantine_10 | Inject Samples | Memantine_QDa_10 | 400 | 10 | 6.50 |
| 13 | 1:A,2 | 1. | Memantine_11 | Inject Samples | Memantine_QDa_11 | 400 | 15 | 6.50 |
| 14 | 1:A,2 | 1 | Memantine_12 | Inject Samples | Memantine_QDa_12 | 400 | 20 | 6.50 |

Figure 4. Sample set method generated using Empower SSG.

A. Peak data across experimental runs [Fmonwer's Peak summary report]

| EIII | linner a I Cak_ | Surrier | ary report | | | |
|------|---------------------------------|-----------------|------------------|---------------------|--------|---------|
| | annel Name: c. Chnl. Descr.: | QDa 1 1: QDa | 0.20 | | | |
| | SampleName | RT | Probe Temp(C) | Cone Voltage (V) | Height | USP s/n |
| 1 | Memantine_3 | 1.775 | 600 | 15 | 36023 | 72 |
| 2 | Memantine_2 | 1.797 | 600 | 10 | 35715 | 68 |
| 3 | Memantine_4 | 1.832 | 600 | 20 | 32793 | 40 |
| 4 | Memantine_7 | 1.792 | 500 | 15 | 31467 | 60 |
| 5 | Memantine_6 | 1.818 | 500 | 10 | 29286 | 36 |
| 6 | Memantine_8 | 1.771 | 500 | 20 | 28327 | 48 |
| 7 | Memantine_11 | 1.782 | 400 | 15 | 25978 | 50 |
| 8 | Memantine_1 | 1.761 | 600 | 5 | 24986 | 40 |
| 9 | Memantine_10 | 1.803 | 400 | 10 | 24413 | 39 |
| 10 | Memantine_12 | 1.822 | 400 | 20 | 23552 | 28 |
| 11 | Memantine_5 | 1.775 | 500 | 5 | 20856 | 39 |
| 12 | Memantine_9 | 1.779 | 400 | 5 | 17530 | 33 |

B. Chromatogram of 10 ng/mL memantine standard solution in 90:10 water/methanol

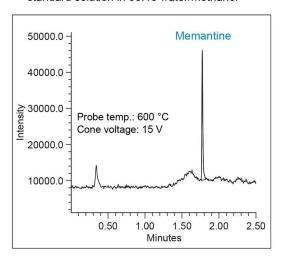


Figure 5. Empower data. Peak data across the experimental runs (A) and representative chromatogram of the memantine standard solution (B).

Conclusion

Using the Empower Sample Set Generator, chromatographic methods were automatically created for analysis with an ACQUITY QDa Mass Detector, to investigate the effect of ionization parameters (probe temperature and cone voltage) on the sensitivity of memantine hydrochloride. The Empower instrument methods and method sets were automatically created and built into the a sample set method as a ready-to-run injection sequence.

Automating methods generation reduced transcriptions errors and time associated with manual process.

The Empower Sample Set Generator automates creation of chromatographic methods for wide range of

| applications | including met | hod development | and validation, | performed on | Waters A | CQUITY LC | Systems and |
|--------------|---------------|-----------------|-----------------|--------------|----------|-----------|-------------|
| detectors | | | | | | | |

References

- 1. https://www.webmd.com/drugs/2/drug-77932-377/memantine-oral/memantine-oral/details < https://www.webmd.com/drugs/2/drug-77932-377/memantine-oral/memantine-oral/details>.
- 2. Maziarz M, Wrona M, McCarthy SM. Benefits of Using QDa Mass Detection for Quantitative Analysis of Non-Chromophoric Memantine HCl in Tablet Formulation. Waters Application Note, 720005179, 2014.

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Arc Premier System https://www.waters.com/waters/nav.htm?cid=135083359

2998 Photodiode Array (PDA) Detector https://www.waters.com/1001362

ACQUITY UPLC PDA Detector https://www.waters.com/514225

Empower Chromatography Data System https://www.waters.com/10190669>

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