

HPLC 2013 MASS-09

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1. Introduction

With the development of highly sensitive and fast LC-MS/MS instruments, the triple quadrupole technology has found its way into clinical drug monitoring and is the method of choice for a number of assays. The steadily increasing number of applications in the clinical sector demands fast and efficient development of new LC-MS/MS methods. The foundation for high quality data is made through optimized chromatographic separations. Fully

automated optimization of the UHPLC method using Shimadzu's method scouting software (Fig. 1) in combination with automated MS optimization for MRM parameters are the perfect platform for the generation of new triple quad MS methods. Here we report a new and fast procedure for the LC-MS/MS method optimization for clinical drug monitoring.

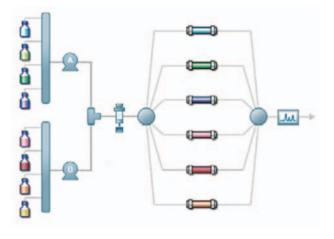


Fig. 1 Combination of columns and solvents during the method scouting process.



LCMS-8040 triple quadrupole mass spectrometer

2. Methods

2-1. LC-MS/MS parameters

One of the first steps during this automated process is the precursor ion selection, followed by the m/z adjustment of the precursor. The collision energy is optimized for the most abundant fragments and finally the fragment m/z

adjustment. Six optimization steps were performed via flow injection analysis, each taking 30 seconds (Fig. 2). The result of these automated steps was the automatic generation of a final MRM method (Table 1).

2-2. UHPLC parameters

Choosing the best HPLC column and composition of eluents are often the most important but time-consuming steps during method development. This can influence sensitivity and separation from potentially interfering matrix effects. Shimadzu Method Scouting was used to determine the best HPLC parameters for the analysis of 14 different

drugs. This allowed the combination of 6 HPLC columns with up to 16 different eluents, resulting in the investigation of up to 96 different combinations, requiring only a fraction of the time required by traditional approaches.



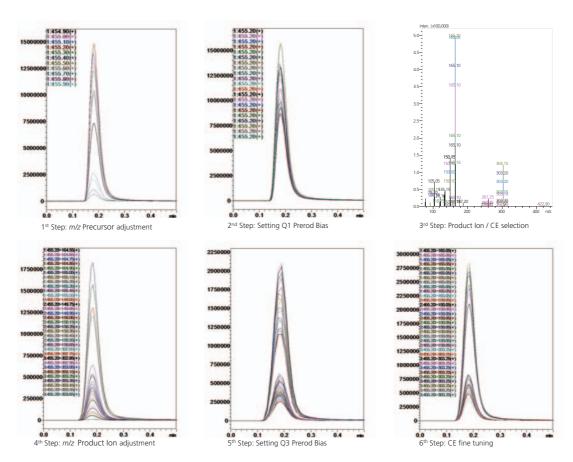


Fig. 2 Automated MRM Optimization of the drug Verapamil on the LCMS 8040

Table 1 Optimized MRM transitions of 14 drugs

Compound	Mode	MRM transitions	Collision energy (kV)
Disopyramide	ESI positive	340.3 > 239.10 / 340.3 > 195.10	-19 / -35
Lidocaine	ESI positive	235.10 > 86.20 / 235.10 > 58.05	-22 / -39
Mexiletine	ESI positive	180.20 > 105.10 / 180.20 > 121.20	-22 / -18
Quinidine	ESI positive	325.30 > 307.10 / 325.30 > 172.05	-26 / -40
Losartan	ESI positive	423.20 > 207.00 / 423.20 > 405.00	-24 / -13
Amiodarone	ESI positive	645.90 > 100.20 / 645.90 > 86.20	-34 / 41
Amitriptyline	ESI positive	278.10 > 105.10 / 278.10 > 233.10	-27 / -18
Chlorpromazine	ESI positive	319.20 > 86.20 / 319.20 > 239.10	-23 / -28
Haloperidol	ESI positive	376.05 > 165.15 / 376.05 > 123.10	-25 / -45
Imipramine	ESI positive	281.25 > 208.00 / 281.25 > 193.10	-27 / -46
Metoprolol	ESI positive	268.25 > 116.15 / 268.25 > 133.00	-20 / -28
Nortriptyline	ESI positive	264.25 > 91.20 / 264.25 > 233.15	-30 / -15
Verapamil	ESI positive	455.20 > 165.05 / 455.20 > 150.05	-34 / -46
Warfarin	ESI negative	307.25 > 160.85 / 307.25 > 249.90	21 / 24



3. Results

3-1. Method development

Traditional method development in HPLC is extremely time consuming. The combination of automated HPLC and MS method development allows the development of complete LC-MS/MS methods within a single day. In this study we show an automated method scouting procedure including the search for optimum column and mobile phase and the gradient conditions. The combination with the fully automated MRM-optimization by flow injection allows a

fast development of a final method for the analysis of clinical drugs. Here we show methods automatically generated for the separation, identification and quantification of a mixture of drugs by the use of 7 different solvents and 6 different columns. The primary step to elucidate the best HPLC conditions out of various combinations is the automated batch creation via method scouting software from Shimadzu (Fig. 3).

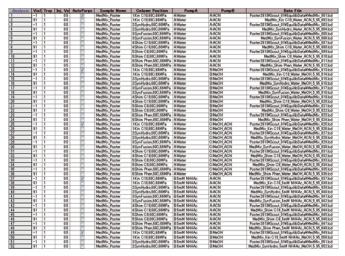


Fig. 3 Batch generated by the method scouting software

Table 2 Solvents and Columns used for method development

Solvent	Column	
AA: Water	Kinetex 2.6 μ C18 100 × 2.10 mm (Phenomenex)	
AB: 5 mM CH ₃ COO ₋ NH ₄ +; pH 8	Synergie 2.5 µ Fusion-RP, 100 × 2.00 mm (Phenomenex)	
AC: 0.1% Formic acid	Synergie 2.5 µ Hydro-RP, 100 × 2.00 mm (Phenomenex)	
AD:10 mM CH₃COO-NH4+; pH 4.5	Shim-pack XR-ODS II 2.2 μ, 100 × 2.00 mm (Shimadzu)	
BA: Acetonitrile		
BB: Methanol	Shim-pack XR-C8 2.2 μ, 100 × 2.00 mm (Shimpack)	
BC: Acetonitrile / Methanol 50/50 (v/v)	Shim-pack XR-Phenyl 2.2 μ , 100 \times 2.00 mm (Shimpack)	

3-2. Data Recording

The first step is the evaluation of the optimal column / solvent combination using a generic gradient starting with 5% of organic solvent increasing to 95% within a specified time. This initial stage generates a viable method requiring some further optimization. The second step optimizes the

slope of the gradient as well as the solvent conditions. Several different conditions were then performed during an overnight analysis. Table 2 shows the used columns and solvents.



3-3. Data comparison

A total number of 162 different combinations were analyzed and evaluated for the best separation and peak intensities (Fig. 4).

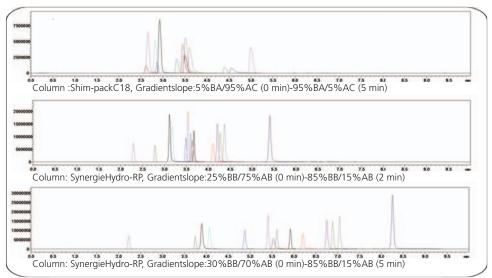


Fig. 4 Examples for poor, medium and good results

3-4. Final method

Flow rate : 0.4 ml / min
Column : Synergie Hydro-RP

Solvent A : 5 mM Ammonium acetate, pH 8

Solvent B : Methanol Oven temp. : 50°C Gradient:

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to change without notice

0 min : 30% B 5 min : 85% B 5.01 min: 95% B 8 min : 95% B 8.01 min: 30% B 10 min : Stop

4. Conclusion

The Combination of the method scouting software tool coupled Shimadzu's ultrafast LCMS 8040 Triple Quad Mass analyzer is a unique tool for fast and easy method development of LC-MS/MS methods. The chromatographic

separation of 14 different drugs as well as their identification and quantification was established successfully within one working day.

First Edition: June, 2013



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