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응용 자료

# A Highly Selective Method for the Analysis of Drospirenone in Human Plasma

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**Abstract** 

Several common birth control formulations contain both drospirenone and ethinyl estradiol. A highly selective and sensitive analytical method for the analysis of drospirenone in human plasma has been developed for use in bioequivalence studies. The solid-phase extraction (SPE) and UPLC-MS/MS methodologies are described as well as performance against validation parameters.

#### Introduction

The challenges of developing a bioanalytical method revolve around meeting the rigorous criteria set forth in the FDA Guidance for Industry for Bioanalytical Method Validation. Methods need to be acceptable in terms of linearity, sensitivity, selectivity, accuracy and precision, stability and carry-over. Recently, the FDA has also included a more comprehensive set of guidelines relating specifically to the assessment and quantitation of matrix effects. Effective, highly selective sample preparation is an important component in the minimization of matrix effects and sample variability.

A bioanalytical method for drospirenone, a component of common birth control formulations, was developed in order to perform bioequivalence studies on generic replacements. Several methods exist for the analysis of ethinyl estradiol,<sup>1-5</sup> but this is believed to be the first reported method for drospirenone.

We describe sample preparation and chromatographic techniques for the analysis of the two synthetic hormones (drospirenone and ethinyl estradiol), and validation data for the analysis of drospirenone.

In this work, we employ mixed-mode SPE to more selectively separate drospirenone and ethinyl estradiol from matrix components. Subsequent analysis was performed by UPLC-MS/MS. Structures for the analytes are shown in Figure 1.

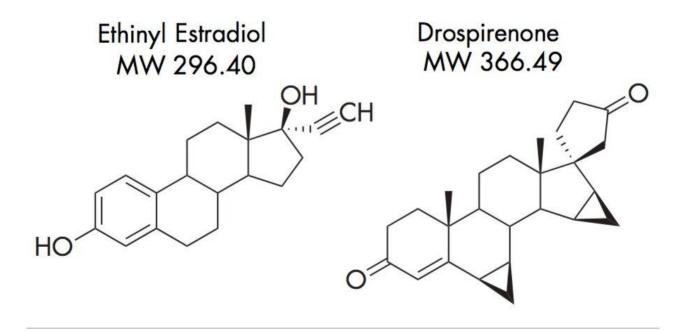


Figure 1. Structures for ethinyl estradiol (17-ethynylestradiol) and drospirenone.

### Experimental

#### **Experimental Conditions**

System:	Waters ACQUITY UPLC System with a Waters	
	Quattro Premier Triple Quadrupole Mass	
	Spectrometer operated in positive ion MRM	
	mode	
Column:	ACQUITY UPLC BEH C <sub>18</sub> Column, 2.1 x 50 mm,	
	1.7 µm Part Number: 186002350	
Mobile phase:	A: 0.1% HCOOH in water	
	B: MeOH (drospirenone) or ACN (ethinyl	
	estradiol)	

Gradient:	70% A to 98% B over 2 min, hold 0.5 min, reset	
	to initial (3 min total cycle time)	
Flow rate:	0.6 mL/min	
Injection:	40 μL	
Temperature:	55 °C	
Desolvation gas flow:	750 L/hr	
_		
Source temperature:	120 °C	
Deschustion temperature	250 °C	
Desolvation temperature:	350 ℃	
Collision cell pressure:	2.6 x 10 <sup>(-3)</sup> mbar	
Complete con processes	2.6 x 105d.	
MRM transitions:	drospirenone 367 -> 96.6	
	ethinyl estradiol 530 -> 171	
Solid-Phase Extraction (SPE)		
Solid I Hase Extraction (of E)		
SPE Device:	Oasis MAX, 30 mg 96-well plate	
	Part Number:186000373	
Condition:	500 μL MeOH	
Condition.	300 με MeOΠ	
Equilibrate:	500 μL water	
Load:	250 μL human plasma diluted and acidified 1:1	
	with 4% H <sub>3</sub> PO <sub>4</sub> in water	
Wash 1:	500 μL 5% NH <sub>4</sub> OH in water	

Elute 1:  $2 \times 125 \mu L 100\% ACN$ 

Elute 2:  $2 \times 125 \mu L$  2% formic acid in 100% ACN

Directly inject elute 1. Dry down elute 2 and reconstitute in 50  $\mu$ L of 10 mM sodium bicarbonate buffer, pH 10.5, add 50  $\mu$ L of 1 mg/mL dansyl chloride solution in acetone and derivatize at 60 °C for 20 minutes.

#### Results and Discussion

Mixed-mode SPE is highly selective and has been shown to be a very powerful tool in reducing matrix effects for bioanalytical assays. As this method is to be used in the determination of bioequivalence, it must meet all of the FDA requirements for validation, including the recent guideline relating to variability of matrix effects. The FDA now recommends that matrix effects be quantitated in 6 sources of matrix and that the variability in matrix effects not exceed  $\pm$  15%. Variability of matrix effects was determined to be less than  $\pm$  15% across all sources (Table 1).

Human Plasma Source	% Matrix Effects for Drospirenone
Donor 1	7.4
Donor 2	5.9
Donor 3	3.8
Donor 4	-6
Pooled Sample 1	-5.1
Pooled Sample 2	-6

Table 1. Calculated matrix effects for drospirenone in 6 sources of human plasma prepared by mixed-mode anion exchange SPE.

We included 4 individual donors and 2 lots of pooled plasma in the analysis to prove that the assay is selective for drospirenone. Minimization of variability across different sources of matrix (i.e., individual subjects in clinical studies), has been a major point of discussion in relation to Incurred Sample Reanalysis (ISR.) Many researchers believe matrix effects arising from sample variability are one of the primary causes for failure in repeat analysis.

SPE recovery was calculated for both drospirenone and ethinyl estradiol. Drospirenone was found in eluate 1 (bound by reversed phase) as expected and ethinyl estradiol was found in eluate 2 (bound by anion exchange to the resin). Analyte SPE recovery and RSDs are shown in Table 2.

	% SPE Recovery	% RSD
Drospirenone	109	7.2
Ethinyl Estradiol	89	4.5

Table 2. SPE recoveries (n=8) and RSDs for drospirenone and ethinyl estradiol.

The method is linear (1/x weighting) for the determination of drospirenone in human plasma from 0.5 to 250 ng/mL. Correlation coefficients were greater than 0.997, and deviations from the actual concentrations were less than 15%, including at the lower limit of quantitation (LLOQ). Sensitivity at the LLOQ of 0.5 ng/mL is shown in Figure 2. Calculated concentrations for QC's at 0.75, 25 and 200 ng/mL were all within 15% of expected values.

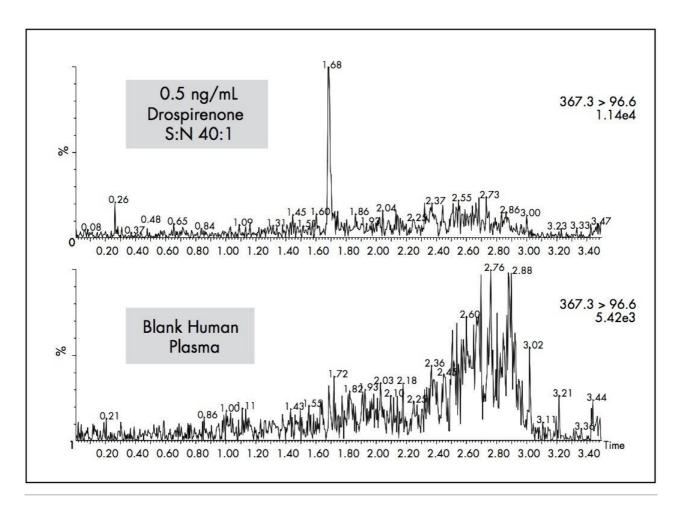


Figure 2. UPLC-MS/MS analysis of a blank plasma sample (A) and 0.5 ng/mL drospirenone extracted from human plasma (B).

#### Conclusion

A sensitive, robust, and reliable UPLC-MS/MS method was developed for quantification of drospirenone in human plasma to support a bioequivalence study. Matrix effects were virtually eliminated through the combination of mixed-mode SPE and UPLC-MS/MS. Combining SPE in a 96-well format with 3-minute cycle times yielded sample throughput of over 400 samples a day without compromising data quality.

#### References

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