

A Novel Clinical Research Method for the Unambiguous Measurement of Low-Level Testosterone in Serum using Differential Ion Mobility Spectrometry-Tandem Mass Spectrometry (DMS-MS/MS)



Michael J. Y. Jarvis
SCIEX, Concord, Canada

INTRODUCTION

It has been well documented that liquid chromatography-tandem mass spectrometry (LC-MS/MS) provides excellent accuracy, precision and sensitivity for measurements of steroids in biological matrices compared to traditional techniques such as immunoassays, which may suffer from cross-reactivity.

Nevertheless, there are numerous uncharacterized, endogenous components in biological fluids which have the potential to interfere with the measurement of low-level steroids such as testosterone. In this work, we present a novel method employing differential ion mobility spectrometry (DMS) in conjunction with LC-MS/MS analysis to eliminate potential interferences, thereby simplifying sample pre-treatment and enabling reduced LC run-times.

MATERIALS AND METHODS

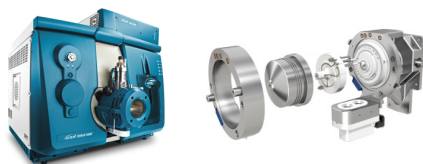


Figure 1. SCIEX Triple Quad™ 6500+ system, equipped with the SelexION+ ion mobility device.

We have applied liquid chromatography (LC), differential ion mobility spectrometry (DMS), and tandem mass spectrometry (MS/MS) to enable the accurate quantitation of low-level testosterone in human serum. The DMS cell filters out potential interferences prior to detection by MS/MS, ensuring that isobaric components do not obfuscate the analysis.

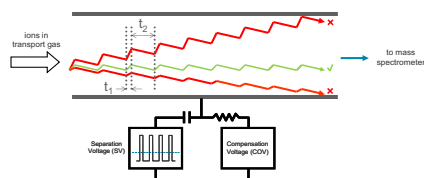


Figure 2. The SelexION+ differential ion mobility device consists of two planar electrodes. Voltages are applied, which serve to filter out isobaric interferences prior to detection by tandem mass spectrometry.

Sample Preparation:

Testosterone and Testosterone-d3 were obtained as 1mg/mL standards in methanol, from Cerilliant Corporation (Round Rock, TX). Calibration curves were prepared by spiking known amounts of Testosterone into steroid-free serum obtained from Golden West Biologicals.

Sample preparation consisted of a one-step liquid-liquid extraction. 200µL of serum sample was combined with 50µL of internal standard solution and 1000µL of 90:10 hexane ethyl acetate in a micro-centrifuge tube. The sample was vortex mixed, centrifuged at 14,000 rpm for 15 minutes, and then 900 µL of the organic supernatant was removed and evaporated to dryness under a stream of nitrogen gas. The dried sample was reconstituted in 100µL of methanol, and then further diluted with 100 µL of deionized water.

HPLC Conditions:

Chromatographic separation was accomplished using the Shimadzu Prominence 20AD-XR HPLC system, with a Phenomenex Kinetex C18 column (100x2.1mm, 2.6µm), at a flow rate of 0.6 mL/min. Mobile phase A consisted of water containing 0.2mM NH₄F. Mobile phase B consisted of methanol. The method run-time was 7 minutes.

MS/MS Conditions:

MS/MS detection was performed using the SCIEX Triple Quad™ 6500+ system equipped with IonDrive™ Turbo V source and operated in electrospray ionization mode. Multiple Reaction Monitoring (MRM) mode was employed, with 2 MRM transitions monitored.

Q1	Q3	ID	DP	CE	CXP	SV	COV
289.1	97	Testosterone 1	70	30	12	3700	8.9
289.1	109	Testosterone 2	70	32	11	3700	8.9

Optimization of SelexION+ parameters was performed using T-infusion of Testosterone at mobile phase flow rate of 0.6 mL/min. At a fixed separation voltage (SV) of 3700 V, the compensation voltage (COV) was ramped across a broad voltage range using a step-size of 0.5V. The optimum COV value, producing a maximum in signal intensity, was observed at a value of 8.9 V.

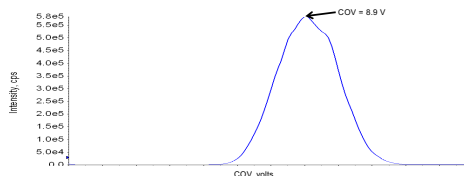


Figure 3. Optimization of the COV parameter, for Testosterone.

RESULTS

We have applied our LC-DMS-MS/MS method to the measurement of anonymized serum samples. As shown in Figure 4, the method displayed excellent sensitivity, and low chemical background due to the application of DMS. The limit of quantitation (LOQ) was observed to be <1 pg/mL, and therefore this LC-DMS-MS/MS method is suitable for the measurement of total Testosterone, as well as the measurement of 'free' testosterone.

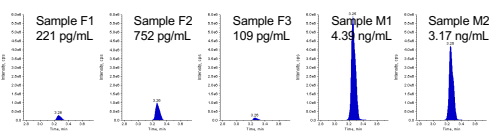


Figure 4a. Example data for analysis of Testosterone in serum.

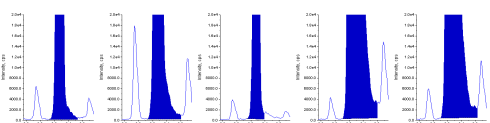


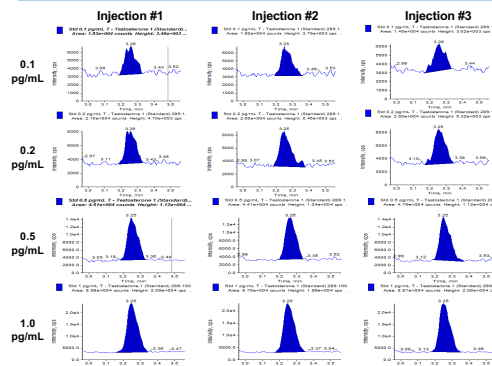
Figure 4b. Zoom on y-axis, to visualize separation of interferences, and low chemical background due to the use of the ion mobility filter.

DISCUSSION

The second generation SelexION+ ion mobility cell incorporates a jet injector lens, and provides >2x improvement in ion transmission vs the original design.

To demonstrate the improved sensitivity when operating the ion mobility cell, a sensitivity comparison in neat solvent is presented in Figure 5. The total counts (cps) are lower when SelexION+ is employed, however the S/N and LOQ are equivalent, or better, when the ion mobility device is used.

Analysis on SCIEX 6500+ LC-MS/MS system (no ion mobility)



Analysis on SCIEX 6500+ LC-MS/MS system, equipped with SelexION+ ion mobility device

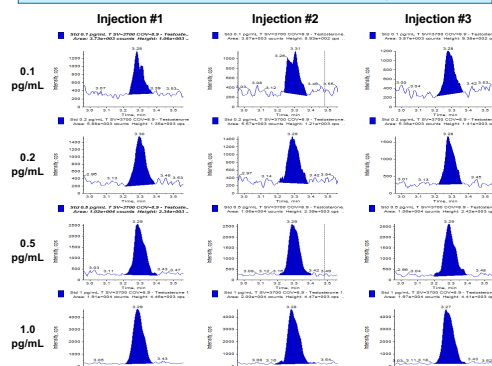


Figure 5. Comparison of LOQ and S/N, in neat solvent, with and without the SelexION+ ion mobility device.

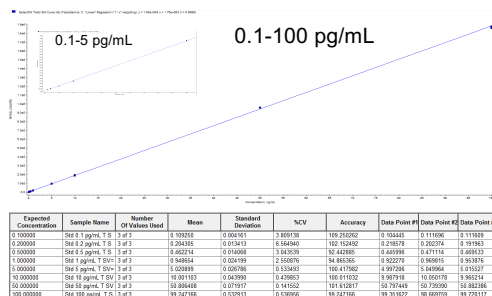


Figure 6. Linear calibration curve (0.1 - 100 pg/mL) for Testosterone in neat solvent, using SelexION+ ion mobility device.

CONCLUSIONS

The LC-DMS-MS/MS method presented here enabled the quantitation of Testosterone in human serum at <1pg/mL. No compromise in analytical sensitivity (LOQ) was observed when employing the ion mobility cell. This method boasts the added advantage of improved specificity, and therefore the possibility of simplified sample preparation.

TRADEMARKS/LICENSING

AB Sciex is doing business as SCIEX. For Research Use Only. Not for use in diagnostic procedures. The trademarks mentioned herein are the property of AB Sciex Pte. Ltd. or their respective owners. AB SCIEX™ is being used under license.