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Note d'application

Rapid Quantification of Codeine by Desorption/Ionization on Silicon (DIOS) Time-of-Flight Mass Spectrometry

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

Abstract

- 1. Perform relative quantification of codeine by DIOS-Tof MS and LC-MS
- 2. Compare the sample cycle time, precision, and accuracy of quantification by DIOS-Tof MS and LC-MS
- 3. Quantify codeine in a pain medication elixir

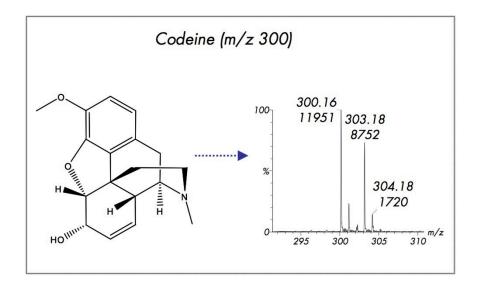
Benefits

- · Rapid sample cycle times are generally less than 1 minute
- · DIOS is a matrix-free method, so the analyte signal is not obscured by chemical noise from the matrix
- · Method development is minimal and simple providing for quick results
- · The technique provides both qualitative and DIOS-Tof MS with the Waters Micromass MassPREP DIOS-

target Plate adds a new dimension to the capabilities of the Waters Micromass MALDI micro MX Time-of-Flight Mass Spectrometer quantitative information

Introduction

Quantitative analysis of small molecules using laser desorption methods has long been an appealing concept due to high throughput capabilities of laser desorption time-of-flight mass spectrometers combined with the inherent mass selectivity. Both matrix-assisted laser/desorption ionization (MALDI)¹⁻³ and desorption/ionization on silicon mass spectrometry (DIOS)⁴⁻⁷ have a number of applications showing small molecule quantification. In most cases, the key to successful quantification is the presence of an internal standard and appropriate sample preparation.⁸





DIOS-Tof MS with the Waters Micromass
MassPREP DIOS-target Plate adds a new
dimension to the capabilities of the Waters
Micromass MALDI micro MX Time-of-Flight
Mass Spectrometer.

Results and Discussion

DIOS-Tof MS Codeine Quantification

Codeine was quantified relative to a deuterated internal standard. The standard curve is based on the correlation between the mass spectral intensity ratio (codeine/codine-d3) and the concentration of codeine (Figure 1A). Examples of the DIOS-Tof MS spectra are shown in Figure 1B with mono-isotopic peaks for codeine (m/z 300) and the internal standard codeine-d3 (m/z 303).

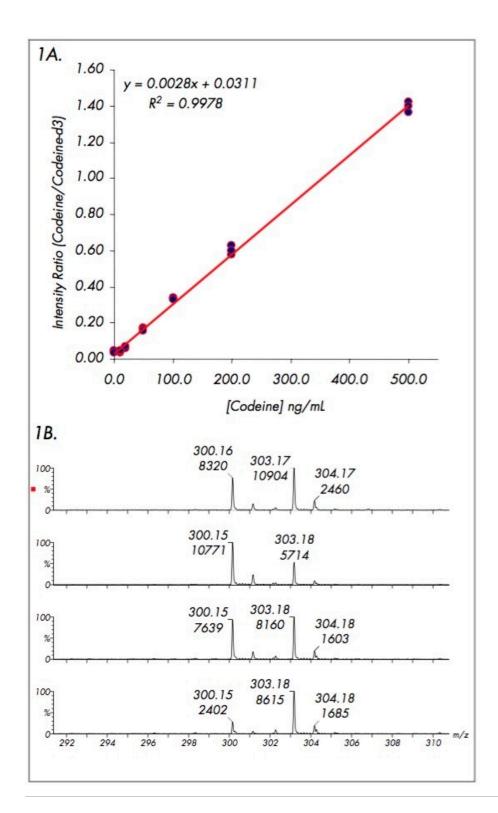


Figure 1A. The standard calibration curve for quantification of codeine derived from DIOS-Tof MS peak intensity

ratios. Figure 1B. Representative DIOS-Tof MS mass spectra of quality control (QC) and pain medication elixir samples.

LC-MS Codeine Quantification

The LC-MS method was operated in SIR (selected ion recording) mode and all data was analyzed using QuanLynx Application Manager for MassLynx v4.0 Software. The standard curve extends from 10.0 to 2000.0 ng/mL of codeine and the linear regression R² value of 0.9997 (Figure 2A), a significant improvement over the DIOS-Tof MS method calculations. However, the method development time and sample cycle times for the LC-based approach were markedly longer than those for the DIOS method. The DIOS method provides a significant advantage of increased sample throughput.

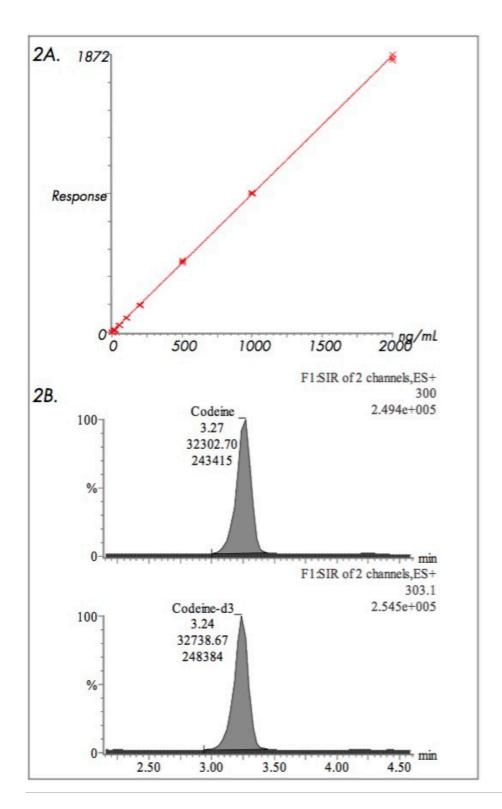


Figure 2A. The standard calibration curve for quantification of codeine from LC-MS SIR peak area ratios. Figure

2B. Peak integration of examples of codeine (m/z 300) and codeine-d3 (m/z 303).

Comparison of DIOS-Tof MS and LC-MS Codeine Quantification

The data in Table 1 indicate that for the DIOS-Tof MS method, from 15.0 to 800.0 ng/mL, the average % error and %RSD are 8.6 and 10.0 respectively while the sample cycle time is 66 seconds. The LC-MS approach, from 10.0 to 2000.0 ng/mL provided an average % error of 2.3%, %RSD of 1.1% and sample cycle time of 20 minutes.

| | | A. DIOS-TOF MS: MS Intensity Ratio | | | B. LC/MS: Peak Area Ratio | | |
|----------------------|------------------------------|---------------------------------------|---------|-------|-------------------------------|---------|-------|
| Sample | Actual [Codeine] ng/mL | Average [Codeine] ng/mL | % Error | % RSD | Average [Codeine] ng/mL | % Error | % RSD |
| Elixir | 240.0 | 270.8 | 12.8 | 0.9 | 239.6 | -0.2 | 0.8 |
| QC 15 | 15.0 | 15.3 | 1.7 | 43.2 | 15.7 | 4.4 | 0.4 |
| QC 90 | 90.0 | 91.8 | 2.0 | 3.9 | 89.5 | -0.6 | 0.9 |
| QC 300 | 300.0 | 328.6 | 9.5 | 1.5 | 317.6 | 5.9 | 2.4 |
| QC 800 | 800.0 | 662.5 | -17.2 | 0.6 | 803.6 | 0.5 | 0.8 |
| Root Mean Squared | | | 8.6 | 10.0 | | 2.3 | 1.1 |
| Linear Dynamic Range | | 15.0 to 800.0 ng/mL | | | 10.0 to 2000.0 ng/mL | | |
| | Cycle Time | 66 seconds | | | 20 minutes | | |

Table 1. Summary of quantification results for codeine using (A.) DIOS-Tof MS intensity ratios and (B.) LC-MS selected ion recording (SIR) peak area ratios.

Conclusion

- DIOS-Tof MS provides for quantification of codeine over roughly 2 orders of magnitude dynamic range with 8.6% error and 10.0% RSD.
- · LC-MS analysis of the same samples run by DIOS-Tof MS provides for at least 2.5 orders of magnitude

dynamic range with 2.3% error and 1.1% RSD.

The sample cycle time for DIOS-Tof MS (66 seconds) is significantly faster than that of LC-MS (20 minutes).
 This advantage, combined with simple and rapid method development makes DIOS-TofF MS an attractive alternative technique for high throughput semi-quantitative analysis.

Further reading for complete experimental details:

Wall, D, Finch J, Cohen C. Letter to the Editor: Quantification of codeine by desorption/ionization on silicon (DIOS) time-of-flight mass spectrometry and comparisons to liquid chromatography mass spectrometry. *Rapid Commun. Mass Spectrom.* 2004; in press.

References

- 1. Cohen L, Gusev A. Anal. Bioanal. Chem. 2002; 373: 571-586.
- 2. Hatsis P, Brombacher S, Corr J, Kovarik P, Volmer D. Rapid Commun. Mass Spectrom. 2003; 17: 2303-2309.
- 3. Bucknall M, Fung K, Duncan M. J. Am. Soc. Mass Spectrom. 2002; 13: 1015-1027.
- 4. Go E, Shen Z, Harris K, Siuzdak G. Anal. Chem. 2003; 75: 5475-5479.
- 5. Huikko K, Ostman P, Sauber C, Mandel F, Grigoras K, Franssila S, Kotiaho T, Kostiainen R. *Rapid Commun. Mass Spectrom.* 2003; 17: 1339–1343.
- 6. Lewis W, Shen Z, Finn M, Siuzdak G. Int. J. Mass Spectrom. 2003; 226: 107.
- 7. Thomas J, Shen Z, Crowel J, Finn M, Siuzdak G. PNAS 2001; 98: 4932-4937.

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