Analysis of Synthetic Drugs Using Electron and Chemical Ionization High Resolution Time-of-Flight Mass Spectrometry

David E. Alonso, Joe Binkley, and Jeff Patrick | LECO Corporation, Saint Joseph, MI USA

Introduction

The development of unregulated recreational drugs has been increasing at an alarming rate since the mid-2000s. Relatively simple organic transformations produce novel and licit psychotics that can elude detection by standard analytical methods (e.g., GC-MS). This is a major problem for law enforcement agencies; forensic laboratories; and the medical community since the legal status, physiological effects, and long-term toxicity of these synthetic drugs are unclear. Detection and characterization of these substances through routine forensic analyses is challenging due to their novelty and lack of appropriate instrumental methods. Matters are complicated further by the wide range of active ingredients and variety of botanical blends in samples which can be difficult to handle instrumentally.

High performance time-of-flight mass spectrometry (TOFMS) is a practical choice for the analysis of these substances:

- Data acquisition is comprehensive and robust
- Single acquisition, high-resolution accurate mass data can be probed multiple times for traditional and emerging drugs
- High resolving power minimizes background interferences
- Excellent mass accuracy values facilitate confident elemental formula determination for compound characterization
- Complete and high-quality spectral data can be searched against databases

Traditional

Heroin

Cocaine

Δ^9-THC

Synthetic

Phenethylamines

Phenylpiperazines

Benzylpiperazines

Tryptamines

SynCannabinoids (JWH-Type)

Figure 1: Traditional and Synthetic Drug Examples

In this study, two samples (Cases 1 and 2) obtained from a collaborating forensics laboratory were analyzed and found to contain traditional and new psychotropic substances. The combination of electron impact and chemical ionization high resolution TOFMS was vital for unambiguous identification of new drugs in these mixtures.

Experimental

Sample Preparation

Case 1: Suspected Cathinone

White Powder (29.4 mg) → 1 mL NaOH(aq) → 3 x 1.0 mL DCM → Na₂SO₄ (Anhydrous) → Filter, Dry (N₂ g) → Clear Oil (29.4 mg) → 3 mL 2:1 DCM/MeOH

Case 2: Drug Pipe (Cannabinoids)

Dark Mixture (93 mg) → 3 mL 2:1 DCM/MeOH

Figure 2. Sample Preparation Procedures

Instrument Parameters

GC Parameters

GC:	Agilent 7890 and 7693 Auto Sampler
Column Type:	Restek Rxi-5Sil MS (30 m, 0.25 mm ID, 0.25 mm *df*)
Injection:	0.5 mL, Split 150:1 (CI 100:1); Inlet Temp. 250°C
Oven:	60°C (2 min) to 300°C at 20°C/min (1 min)
Carrier Gas:	He, Constant Flow (1.00 mL/min)

MS Parameters

Spectrometer:	LECO Pegasus® GC-HRT
Ion Sources:	LECO EI, CI
Folded Flight Path:	HR (R = 25,000 FWHM)
Spectral Acquisition:	6 spectra/second
Mass Range (m/z):	30–650 (CI 60–650)
Calibration:	PFTBA (Internal)
CI Reagent Gas:	5% Ammonia in Methane

Analysis Workflow

Dilute & Shoot → EI-HRT MS Data → Library Match? → Yes → M⁺ Formula Confirmation

Library Match? → No → CI-HRT MS Data → MH⁺ Formula Confirmation → Database/Literature Search → Propose a Structure (Standard Available?)

Figure 3: Drug Analysis Workflow; EI and CI-HRT Analysis

Results (Case 1: White Powder)

Analysis of the sample produced the analytical ion chromatogram shown in Figure 4. The four major components of the mixture identified through library searches were 2-Fluorophenyl 2-methylbenzoate, *N*-Ethylamphetamine, Ethylone and Dimethylone (Figure 4). Library matches for these compounds were very good (799 to 935/1000); however, quick inspection of the CI-HRT data demonstrated that the number one hit for peak #2, *N*-ethylamphetamine (C₁₁H₁₇N), was incorrect (Figure 5). A formula search on the analyte's protonated molecular ion at m/z = 192.13825 resulted in the formula C₁₂H₁₈NO (-0.21 ppm). An internet search using the formula C₁₂H₁₇NO, led to Cayman Chemical and Scientific Working Group Sites and 4-methylethcathinone as a potential candidate for this compound.

Figure 4: AIC - Major Components, Sample 1

Peak #	Name	Formula	R.T. (s)	Area	LM (1000)
1	2-Methylbenzoic acid, 2-fluorophenyl ester	C14H11FO2	459.7	265777	935
2	N-Ethylamphetamine	C11H17N	551.8	1081942	809
3	Ethylone	C12H15NO3	641.4	731528	799
4	Dimethylone	C12H15NO3	646.2	1882302	802

N-Ethylamphetamine
C₁₁H₁₇N
Expected M⁺ = 163.13555

[MH]⁺ = C₁₂H₁₈NO
-0.21 ppm

Figure 5: EI (top) and CI-HRT Data (bottom) for Analyte 2, Sample 1

EI and CI-HRT data for ethylone and dimethylone is shown in Figs. 6 and 7. Protonated molecular ions (C₁₂H₁₅NO₃) for these structural isomers were found at m/z = 222.11249 (0.08 ppm) and m/z = 222.11247 (-0.015 ppm) respectively.

EI-HRT Data — Ethylone — LM = 799/1000

CI-HRT Data — [MH]⁺ C₁₂H₁₅NO₃ 0.08 ppm

Figure 6: EI (top) and CI-HRT Data (bottom) for Ethylone, Sample 1

EI-HRT Data — Dimethylone — LM = 802/1000

CI-HRT Data — [MH]⁺ C₁₂H₁₅NO₃ -0.015 ppm

Figure 7: EI (top) and CI-HRT Data (bottom) for Dimethylone, Sample 1

Results (Case 2: Drug Pipe)

Sample 2 was a heterogeneous, dark, solid material that was scraped from the inside of a confiscated drug pipe. A representative set of the over 130 compounds in the sample are listed in Table 1. These include nitriles, phenols, sterols and various heterocyclic compounds. An extracted ion chromatogram XIC (Figure 8) of the mixture shows the major component A, which was not present in any of the commercial libraries used during data processing. The XIC also displays an additional unknown (*B), as well as various drugs detected in the sample.

Table 1: Compounds In Sample 2

Name	Similarity	Formula	Expected Ion m/z	Mass Accuracy	Mass Delta
Delta 9-THC	864	C21H30O2	314.22403	-0.12	-0.00004
JWH-018	731	C24H23NO	341.17742	-1.39	-0.00047
JWH-122 (1-pentyl-3-(4-methyl-1-naptho)indole)	779	C25H25NO	355.19307	-0.65	-0.00023
JWH-081 (1-pentyl-3-(4-methoxy-1-naptho)indole)	863	C25H25NO2	371.18798	-0.88	-0.00033

Figure 8: XIC of Sample 2

Ave. = 0.53 ppm

CI-HRT data was critical for the identification of unknowns A and B (Figure 9). EI-HRT Library hits for these unknowns were poor. Confirmation of unidentified compound formulas was accomplished using CI-HRT data. Internet and database searches of the data sets led to AM-2201 and JWH-022 as probable hits, which were then confirmed through the analysis of standard spectra.

Peak True — EI-HRT Data *A — Library Match 546/1000 X

CI-HRT Data — MH⁺ — AM-2201

Name	Formula	Expected Ion m/z	Mass Accuracy (ppm)	Mass Delta (Da)
Unknown A	C24H23FNO	360.17582	1.38	0.00050

Peak True — EI-HRT Data *B — Library Match 687/1000 X

CI-HRT Data — MH⁺ — JWH-022

Name	Formula	Expected Ion m/z	Mass Accuracy (ppm)	Mass Delta (Da)
Unknown B	C24H22NO	340.16959	1.35	0.00046

Figure 9: EI (top) and CI-HRT Data (bottom) for Unknowns A and B

Summary

A combination of EI and CI-HRT resulted in confident identification of traditional and emerging synthetic drugs. The GC-HRT's high resolution, quality spectral data, and excellent mass accuracy values are very useful for the identification of novel compounds in complex matrices. The accurate mass data provided formulas that were searched against forensic databases for identification of unknowns in drug samples.