

Cover Sheet for

ENVIRONMENTAL CHEMISTRY METHOD

Pesticide Name: Clothianidin

MRID # : 454226-02

Matrix: Soil

Analysis: LC/MS/MS

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Study Title

Analytical Method for the Determination of TI-435 and the Degradates, TZNG, TZMU, MNG and TMG in Soil by Liquid Chromatography with APCI MS/MS-Detection

Data Requirement

USEPA – Subdivision N, 164-1; Soil Dissipation - Supplemental
Canada PMRA – DACO No. 8.3.2 Soil Method

454226-02

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Completion Date

February 6, 2001

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Bayer Report No.

110263



Statement of Data Confidentiality Claims

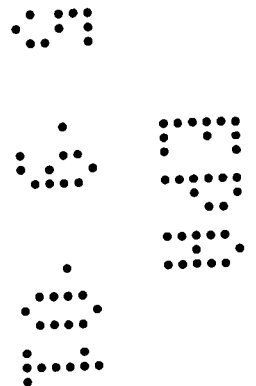
No claim of confidentiality is made for any information contained in this study on the basis of it falling within the scope of FIFRA section 10(d)(1) (A), (B) or (C).

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Good Laboratory Practice Certification

The study was not conducted in compliance with EPA Good Laboratory Practice Standards (40 CFR 160, August 17, 1989). The data presented in the report has been verified by the Sponsor and Ricerca LLC, as being accurate. However, a GLP audit of the data by Ricerca LLC, QAU was not performed.

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Certification of Availability of Raw Data

It is hereby certified that the registrant possesses or has access to all raw data for this report. All raw data or certified copies of raw data which was acquired but has not been included in this report has been archived by Bayer Corporation, Agriculture Division, P.O. Box 4913, Kansas City, Missouri, 64120-0013.

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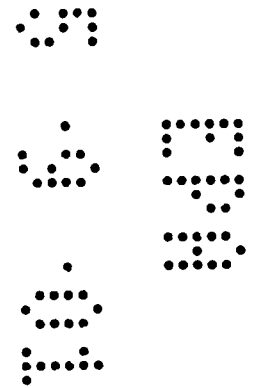
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Quality Assurance Statement

Because this study was not performed under GLP (Good Laboratory Practice regulations of FIFRA, Part 160, August 17, 1989) no QAU audits were performed during the study.



Certification of Authenticity

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Analytical Method for the Determination of TI-435 and the Degradates, TZNG, TZMU, MNG and TMG in Soil by Liquid Chromatography with MS/MS-Detection

1.0 Summary

This report describes the determination of the active ingredient TI-435 and its degradates TZNG, TZMU, MNG and TMG in soil. The method is a modification of the soil residue method described in Bayer report 109586.

Soil samples of 20 g are extracted on an ASE 200 extractor with approximately 45 mL acetonitrile/water/acetic acid/guanidine hydrochloride (25/100/0.1/1; v/v/v/w). Following the extraction, 1 mL of an internal standard solution is added and the volume is adjusted to 50 mL with acetonitrile/water (1:4). Identification and quantification of the active ingredient and the degradates are performed by high performance liquid chromatography using atmospheric pressure chemical ionization (APCI) tandem mass spectroscopy (MS/MS) detection. Isotopically labeled internal standards (d_3 -TI-435, ^{13}C , ^{15}N -TZNG, d_3 -TZMU, d_3 -MNG and d_3 -TMG) are used to compensate for possible matrix effects in the MS/MS-detector.

A loam soil was fortified at two levels (5 and 50 $\mu\text{g}/\text{kg}$) for determining the accuracy of the method. The mean recoveries of the method, determined for the 5.0 $\mu\text{g}/\text{kg}$ fortification ($n=5$) level were 100.4% for TI-435 (relative standard deviation (RSD) = 5.1%), 100.9% for TZNG (RSD = 8.2%), 97.3% for TZMU (RSD = 4.9%), 91.9% for MNG (RSD = 5.5%) and 91.7% for TMG (RSD = 3.0%).

The mean recoveries of the method, determined for the 50.0 $\mu\text{g}/\text{kg}$ fortification level ($n=5$) were 100.3% for TI-435 (RSD = 3.1%), 93.4% for TZNG (RSD = 5.9%), 95.9% for TZMU (RSD = 4.5%), 111.1% for MNG (RSD = 2.0%) and 101.1% for TMG (RSD = 7.9%).

Duplicate control samples and one reagent blank sample were also analyzed and did not show any interferences that would inhibit the analysis of the five compounds in this study.

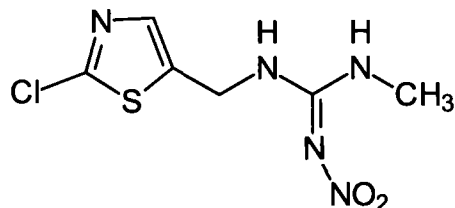
A calibration curve based on nine separate standard concentrations was used to quantify the concurrent recoveries. Standards from 1 $\mu\text{g}/\text{L}$ to 500 $\mu\text{g}/\text{L}$ were used (equivalent to 2.5 $\mu\text{g}/\text{kg}$ to 1250 $\mu\text{g}/\text{kg}$ samples), and the correlation coefficient for each compound ranged from 0.997 to 0.999.

The limit of quantification (LOQ) of the method is 5 $\mu\text{g}/\text{kg}$ for TI-435 and its degradates. The Method Detection Limit (MDL) of the method was 0.95, 1.55, 0.90, 0.96 and 0.51 $\mu\text{g}/\text{kg}$ for TI-435, TZNG, TZMU, MNG and TMG, respectively. The MDL was based upon the standard deviation of five replicate analyses of fortifications made at the LOQ.

2.0 Introduction

The active ingredient TI-435 is an insecticide. A soil residue method is necessary to analyze soil samples collected from soil dissipation studies required for registration of the product. Analysis was targeted for TI-435 and degradates, TZNG, TZMU, MNG and TMG, which were identified at significant concentrations (>10%) in the environmental fate studies conducted with TI-435. The chemical structure of TI-435 and its degradates are:

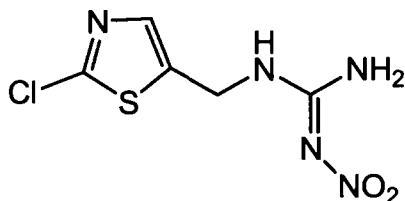
- **TI-435**



Chemical Name (CAS): (*E*)-*N*-[(2-Chloro-5-thiazolyl)methyl]-*N'*-methyl-*N''*-nitroguanidine

CAS Number: 210880-92-5
(previously 205510-53-8)
Formula: C₆H₈ClN₅O₂S
Molar Mass: 249.7 g/mol
Bayer Reference No.: M00343
Purity: 99.8%
Expiration Date: August 2001

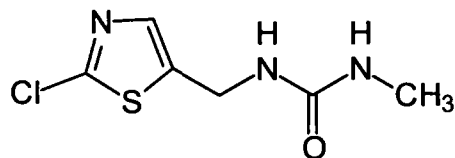
- **TZNG (thiazolyl-nitroguanidine)**



Chemical Name (CAS): *N*-(2-Chlorothiazol-5-ylmethyl)-*N'*-nitroguanidine

Formula: C₅H₆ClN₅O₂S
Molar Mass: 235.7 g/mol
Reference No.: M00674
Purity: 99.7%
Expiration Date: February 2002

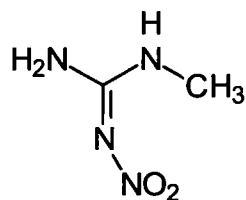
- **TZMU (thiazolyl-methylurea)**



Chemical Name (CAS): *N*-(2-Chlorothiazol-5-ylmethyl)-*N'*-methylurea

Formula: C₆H₈ClN₃OS
Molar Mass: 205.7 g/mol
Reference No.: M00886
Purity: 98.0%
Expiration Date: August 2002

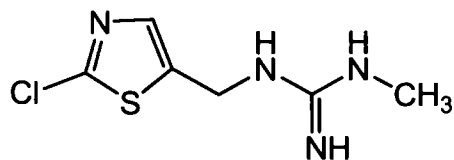
- **MNG (methyl-nitroguanidine)**



Chemical Name (CAS): *N*-Methyl-*N'*-nitroguanidine

Formula: C₂H₆N₄O₂
Molar Mass: 118.1 g/mol
Reference No.: M00500
Purity: 99.0%
Expiration Date: February 2002

- **TMG (thiazolyl-methylguanidine)**

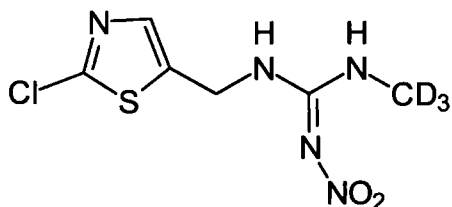


Chemical Name (CAS): *N*-(2-Chlorothiazol-5-ylmethyl)-*N'*-methylguanidine

Formula: C₆H₉ClN₄S
 Molar Mass: 204.7 g/mol
 Reference No.: M01822
 Purity: 91.0%
 Expiration Date: March 2001

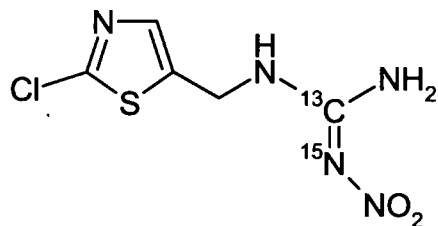
The Reference Substances (Internal Standards) have the following structures:

- **TI-435-d₃**



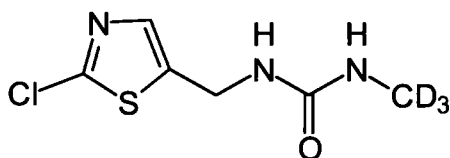
Formula: C₆H₅ClD₃N₅O₂S
 Molar Mass: 252.7 g/mol
 Reference No.: M01288
 Purity: 99.0%
 Expiration Date: October 2002

- **TZNG-¹³C, ¹⁵N**



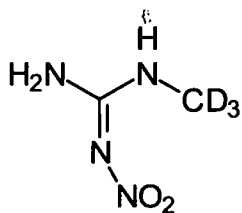
Formula: C₄¹³CH₆ClN₄¹⁵NO₂S
 Molar Mass: 237.7 g/mol
 Reference No.: M01322
 Purity: 99.0%
 Expiration Date: February 2001

- **TZMU-d₃**



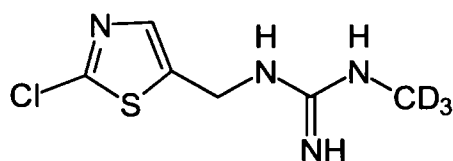
Formula: C₆H₅ClD₃N₃OS
 Molar Mass: 208.7 g/mol
 Reference No.: M01897
 Purity: 99.2%
 Expiration Date: October 2002

- **MNG-d₃**



Formula: C₂H₃D₃N₄O₂
 Molar Mass: 121.1 g/mol
 Reference No.: M01289
 Purity: 99.0%
 Expiration Date: November 2002

• **TMG-d₃**



Formula: C₆H₆ClD₃N₄S
 Molar Mass: 207.7 g/mol
 Reference No.: M01898
 Purity: 99.4%
 Expiration Date: October 2002

2.1 Test Facility and Study Dates

All modifications and validation of the method was performed at Ricerca LLC (7528 Auburn Road, Painesville, Ohio, 44077-1000). Work associated with method development and validation occurred in July of 2000.

3.0 Test System

The method was validated using soil from a dissipation site in Watsonville, California (USA). The soil was selected to represent a “difficult” matrix. The soil samples were classified according to USDA specifications. Soil textural characterization is summarized in Table 1.

Table 1: Soil Characteristics

| Parameter | Value |
|-------------------------------------|-------|
| sand (%) | 38.8 |
| silt (%) | 38.4 |
| clay (%) | 22.8 |
| texture | loam |
| Organic Matter (%) | 2.0 |
| pH | 5.4 |
| Cation Exchange Capacity (meq/100g) | 8.6 |
| Water Holding Capacity-1/3 bar (%) | 21.5 |
| Water Holding Capacity-15 bar (%) | 6.3 |
| Bulk Density (g/mL) | 1.57 |
| Soil Series Classification | Elder |

4.0 Principle of the Method

With the following method, the active ingredient TI-435 and the degradates TZNG, TZMU, MNG and TMG can be determined in soil down to a limit of quantification of 5 µg/kg (LOQ). The method is a modification of the soil residue method described in Bayer Report 109586. Some difficulties encountered with the LC-MS/MS analyses required modification of the existing method, and are described here. All solvent, standard, and extraction conditions are the same as those described in Bayer Report 109586, which also provides additional information regarding mass spectra of the analytes, and linear response curves.

Soil samples of 20 g are extracted on an ASE 200 extractor with approximately 45 mL of a mixture of acetonitrile/water/acetic acid/guanidine hydrochloride (25/100/0.1/1; v/v/v/w). Following the extraction, 1 mL of an internal standard solution is added and the volume is

adjusted to 50 mL with acetonitrile/water. Identification and quantification of the active ingredient and the degradates are performed by high performance liquid chromatography using atmospheric pressure chemical ionization (APCI) tandem mass spectrometry (MS/MS) detection. Isotopically labeled internal standards (d_3 -TI-435, ^{13}C , ^{15}N -TZNG, d_3 -TZMU, d_3 -MNG and d_3 -TMG) are used to compensate for possible matrix effects in the MS/MS-detector.

5.0 Standards and Sample Preparation

Calibration Standards and the samples analyzed by LC-MS/MS were composed of the same ratio of acetonitrile and water (1:4). Additionally, the amount of internal standard was kept constant at 50 $\mu\text{g/L}$ (125 $\mu\text{g/kg}$ soil equivalents) in all samples analyzed. The ratio of native material to internal standard was plotted for each of the calibration standards in a straight line fit, and used to determine the concentration of native material in the recovery samples.

5.1 Standard Preparation

Calibration standards were prepared according to Bayer Report 109586 at the following concentrations: 1, 2.5, 5, 10, 25, 50, 100, 250, and 500 $\mu\text{g/L}$ (native standard). These concentrations correspond to soil equivalents of 2.5 to 1250 $\mu\text{g/kg}$. The amount of internal standard was constant in all standards at 50 $\mu\text{g/L}$ in each of these calibration standards.

5.2 Equipment

Extraction Equipment: Accelerated Solvent Extractor (ASE): ASE 200
Dionex Corporation, USA
Sunnyvale, California

Cellulose Filters: (for ASE 200 extraction cell caps)
Part No. 049458, Dionex Corporation

Extraction Solvent: acetonitrile/water/acetic acid/guanidine HCl (25/100/0.1/1; v/v/v/w)
400 mL Acetonitrile (HPLC Grade, Fisher)
1600 mL Water (HPLC Grade, Fisher)
1.6 mL Acetic Acid (ACS Grade, Fisher)
16 grams Guanidine Hydrochloride (>99%, Fluka)

5.3 Extraction Procedure

1. Weigh 4 g of hydromatrix and 20 g of a soil sample into a weigh dish.
2. Mix the contents thoroughly using a spatula.
3. Place a cellulose filter at the bottom of a 33 mL stainless steel cartridge (ASE sample cell).
4. Place the soil/hydromatrix mixture into the ASE cell using a funnel.
5. Extract the samples on the ASE 200 extractor at 140 °C with approximately 45 mL of extraction solvent (see 5.4 for ASE conditions).

6. After extraction, add 1 mL of the internal standard solution and adjust the volume to 50 mL with 1:4 acetonitrile:water.

5.4 ASE Conditions

| | |
|---------------|--------------|
| Preheat: | 0 min |
| Heat: | 7 min |
| Static: | 10 min |
| Flush volume: | 35 % of cell |
| Purge: | 5 min |
| Cycles: | 3 |
| Pressure: | 150 bar |
| Temperature: | 140 °C |

6.0 LC Conditions

Column: Phenomenex AQUA C₁₈ ; length 25.0 cm, i.d. 4.6 mm

Injection volume: 50 µL

Oven temperature: Column at room temperature

Mobile phase: A: water
B: methanol

Run time: 20 min

Flow rate (column): 1.0 mL/min

Flow rate (interface): 1.0 mL/min

| | | |
|------------------|---------|-------------------|
| Retention times: | TI-435: | approx. 9:82 min. |
| | TZNG: | approx. 9:37 min. |
| | TZMU: | approx. 8.92 min. |
| | TMG: | approx. 8.91 min. |
| | MNG: | approx. 3.87 min. |

Table 2: HPLC-Gradient

| | | | | | | | | |
|------------|----|----|----|----|----|----|------|---------------|
| Time [min] | 0 | 1 | 2 | 7 | 7 | 11 | 11.1 | |
| % A | 90 | 90 | 65 | 40 | 20 | 20 | 90 | |
| % B | 10 | 10 | 35 | 60 | 80 | 80 | 10 | ¹⁾ |

¹⁾The gradient can be changed to optimize the separation.

Table 3: HPLC-Timetable

| Time | Setting | Value | |
|------|------------------------|----------|---------------|
| 14.0 | Column Switching Valve | Column 1 | ¹⁾ |

1) switching eluent stream to waste

7.0 Mass Spectroscopy

Mass Spectrometer: PE Sciex API-III with APCI Interface and heated nebulizer.

7.1 Principle of Measurement

Substances introduced into the mass spectrometer are ionized using an APCI interface. Sample ions are accelerated by an adequate voltage regulation and separated by mass in the first quadrupole (Q1). The most abundant ions of the analyte (parent ions) are impulsed with nitrogen in the collision cell (Q2). Fragments of these ions (daughter ions) are separated by mass in the third quadrupole (Q3) and detected. The mass spectroscopic parameters for the analytes and the selected ions are listed in Table 4 to 8 on the following pages.

7.2 Mass Spectroscopic Parameters

The reported parameters are examples for an optimal adjustment of the mass spectrometer. From time to time these parameters have to be checked and adjusted, if necessary, depending on the instrument used. Abbreviations used in Tables 4 to 8 are as follows:

| | |
|-------------------------------|----------------------------------|
| (DI) Discharge Needle Current | (RE3) Quad 3 Resolution |
| (OR) Orifice Plate | (RX) Lens Element RX |
| (R0) Quad 0 Rod Offset | (R3) Quad 3 Rod Offset |
| (RE1) Q1 Resolution | (L9) Lens Element 9 |
| (DM1) Q1 Delta Mass | (FP) Faraday Plate Voltage |
| (R1) Quad 1 Rod Offset | (MU) Channel Electron Multiplier |
| (L7) Lens Element 7 | (DM3) Quad 3 Delta Mass |
| (R2) Quad 2 Rod Offset | |

Table 4: MS/MS-Parameters of TI-435 and d₃-TI-435

Auxiliary gas (N₂): 4 L/min, temperature: 400 °C
 Curtain Gas (N₂): 1.2 L/min
 Nebulizer Gas (N₂): 80 PSI
 Collision Gas (N₂): 236 (setting)

Experiment Information

Scan Type : Multiple Reaction Monitoring

Mass Range Information

| | | |
|---------------|------------------------|--------------|
| Mass Range 1 | TI-435 | |
| Q1 Mass (amu) | Q3 Mass (amu) | Dwell (msec) |
| 250 | 169 | 300 |
| Mass Range 1 | d ₃ -TI-435 | |
| Q1 Mass (amu) | Q3 Mass (amu) | Dwell (msec) |
| 253 | 172 | 100 |

Positive Ion Mode

| Parameter | Value [volt] |
|-----------|--------------|
| DI | 5μA |
| OR | 54 |
| RO | 30 |
| REI | 120 |
| DMI | 0.2 |
| R1 | 26 |
| L7 | 21 |
| R2 | 16 |
| RE3 | 117 |
| DM3 | 0.14 |
| RX | 1.0 |
| R3 | 11.0 |
| L9 | -50 |
| FP | -50 |
| MU | -4800 |

Table 5: MS/MS-Parameters of TZNG and ¹³C, ¹⁵N-TZNG

Auxiliary gas (N₂): 4 L/min, temperature: 400 °C
 Curtain Gas (N₂): 1.2 L/min
 Nebulizer Gas (N₂): 80 PSI
 Collision Gas (N₂): 236 (setting)

Experiment Information

Scan Type : Multiple Reaction Monitoring

Mass Range Information

| | | |
|---------------|---------------------------------------|--------------|
| Mass Range 1 | TZNG | |
| Q1 Mass (amu) | Q3 Mass (amu) | Dwell (msec) |
| 236 | 132 | 300 |
| Mass Range 1 | ¹³ C, ¹⁵ N-TZNG | |
| Q1 Mass (amu) | Q3 Mass (amu) | Dwell (msec) |
| 240 | 134 | 100 |

Positive Ion Mode

| Parameter | Value [volt] |
|-----------|--------------|
| DI | 5μA |
| OR | 54 |
| RO | 30 |
| REI | 120 |
| DMI | 0.2 |
| R1 | 26 |
| L7 | 21 |
| R2 | 16 |
| RE3 | 117 |
| DM3 | 0.14 |
| RX | 1.0 |
| R3 | 11.0 |
| L9 | -50 |
| FP | -50 |
| MU | -4800 |

Remark: Due to the fact, that the m/z 238 ion derives from both the native and internal standard, due to the chlorine isotopes (³⁷Cl-TZNG and ³⁵Cl-¹³C, ¹⁵N-TZNG), the not superimposed ions m/z 236 (³⁵Cl-TZNG) and m/z 240 (³⁷Cl-¹³C, ¹⁵N-TZNG) were used for the identification and quantification of the two compounds.

Table 6: MS/MS-Parameters of TZMU and d₃-TZMU

Auxiliary gas (N₂): 4 L/min, temperature: 400 °C
 Curtain Gas (N₂): 1.2 L/min
 Nebulizer Gas (N₂): 80 PSI
 Collision Gas (N₂): 236 (setting)

Experiment Information

Scan Type : Multiple Reaction Monitoring

Mass Range Information

| Mass Range 1 | TZMU | Q1 Mass (amu) | Q3 Mass (amu) | Dwell (msec) |
|--------------|------|---------------|---------------|--------------|
| | | 206 | 175 | 600 |

| Mass Range 1 | d ₃ -TZMU | Q1 Mass (amu) | Q3 Mass (amu) | Dwell (msec) |
|--------------|----------------------|---------------|---------------|--------------|
| | | 209 | 175 | 300 |

Positive Ion Mode

| Parameter | Value [volt] |
|-----------|--------------|
| DI | 5μA |
| OR | 54 |
| RO | 30 |
| REI | 120 |
| DMI | 0.2 |
| R1 | 26 |
| L7 | 21 |
| R2 | 16 |
| RE3 | 117 |
| DM3 | 0.14 |
| RX | 1.0 |
| R3 | 11.0 |
| L9 | -50 |
| FP | -50 |
| MU | -4800 |

Table 7: MS/MS-Parameters of MNG and d₃-MNG

Auxiliary gas (N₂): 4 L/min, temperature: 400 °C
 Curtain Gas (N₂): 1.2 L/min
 Nebulizer Gas (N₂): 80 PSI
 Collision Gas (N₂): 236 (setting)

Experiment Information

Scan Type : Multiple Reaction Monitoring

Mass Range Information

| Mass Range 1 | MNG | Q1 Mass (amu) | Q3 Mass (amu) | Dwell (msec) |
|--------------|-----|---------------|---------------|--------------|
| | | 117 | 60.7 | 250 |

| Mass Range 1 | d ₃ -MNG | Q1 Mass (amu) | Q3 Mass (amu) | Dwell (msec) |
|--------------|---------------------|---------------|---------------|--------------|
| | | 120.0 | 60.7 | 250 |

Negative Ion Mode

| Parameter | Value [volt] |
|-----------|--------------|
| DI | -3μA |
| OR | -54 |
| RO | -30 |
| REI | -120 |
| DMI | 0.19 |
| R1 | -20 |
| L7 | -26 |
| R2 | -21 |
| RE3 | 119 |
| DM3 | 0.1 |
| RX | -6.0 |
| R3 | -16.0 |
| L9 | 50 |
| FP | 50 |
| MU | 4800 |

Table 8: MS/MS-Parameters of TMG and d₃-TMG

Auxiliary gas (N₂): 4 L/min, temperature: 400 °C
 Curtain Gas (N₂): 1.2 L/min
 Nebulizer Gas (N₂): 80 PSI
 Collision Gas (N₂): 236 (setting)

Experiment Information

Scan Type : Multiple Reaction Monitoring

Mass Range Information

| Mass Range 1 | TMG | | |
|---------------|---------------|--------------|--|
| Q1 Mass (amu) | Q3 Mass (amu) | Dwell (msec) | |
| 205 | 132 | 300 | |

| Mass Range 1 | d ₃ -TMG | | |
|---------------|---------------------|--------------|--|
| Q1 Mass (amu) | Q3 Mass (amu) | Dwell (msec) | |
| 208 | 132 | 100 | |

Positive Ion Mode

| Parameter | Value [volt] |
|-----------|--------------|
| DI | 5μA |
| OR | 54 |
| RO | 30 |
| REI | 120 |
| DMI | 0.2 |
| R1 | 26 |
| L7 | 21 |
| R2 | 16 |
| RE3 | 117 |
| DM3 | 0.14 |
| RX | 1.0 |
| R3 | 11.0 |
| L9 | -50 |
| FP | -50 |
| MU | -4800 |

Table 9: MS/MS-Timetable

| Detection time | Principle | Dwell time | m/z | m/z | Substance |
|----------------|-------------------|------------|--------|---------|---------------------------------------|
| [min] | | [msec] | Parent | Product | |
| 0 – 5 | APCI ⁻ | 250 | 117 | 60.7 | MNG |
| 0 – 5 | APCI ⁻ | 250 | 120 | 60.7 | d ₃ -MNG |
| 5 – 12.8 | APCI ⁺ | 150 | 205 | 132 | TMG |
| 5 – 12.8 | APCI ⁺ | 150 | 208 | 132 | d ₃ -TMG |
| 5 – 12.8 | APCI ⁺ | 150 | 206 | 175 | TZMU |
| 5 – 12.8 | APCI ⁺ | 150 | 209 | 175 | d ₃ -TZMU |
| 5 – 12.8 | APCI ⁺ | 150 | 236 | 132 | TZNG |
| 5 – 12.8 | APCI ⁺ | 150 | 240 | 134 | ¹³ C, ¹⁵ N-TZNG |
| 5 – 12.8 | APCI ⁺ | 150 | 250 | 169 | TI-435 |
| 5 – 12.8 | APCI ⁺ | 150 | 253 | 172 | d ₃ -TI-435 |

APCI⁺ = positive ion mode, i.e. production of positive ionsAPCI⁻ = negative ion mode, i.e. production of negative ions**8.0 Calculations**

For calculation of the concentrations, nine point calibration curves are used. These curves are calculated using linear regression, automatically after each sequence run, with the Perkin Elmer quantification software *MacQuan Version 1.5*.

The linear equation is expressed as:

$$y = \text{Intercept} + \text{Slope} \cdot x$$

$$y = \text{Area}, x = \text{Concentration}$$

For the calculation of standard/internal standard ratios:

$$y = \frac{\text{Area}_{\text{Standard}}}{\text{Area}_{\text{Internal Standard}}} = \text{Int Ratio} \quad \text{and} \quad x = \frac{\text{Conc}_{\text{Standard}}}{\text{Conc}_{\text{IS}}} = \text{ConcRatio}$$

$$\text{Int. Ratio} = \text{Intensity ratio}$$

$$\text{Conc}_{\text{Standard}} = \text{Concentration of standard solution } [\mu\text{g/L}]$$

$$\text{Conc}_{\text{IS}} = \text{Concentration of internal standard solution } [\mu\text{g/L}]$$

$$\text{Conc}_{\text{ratio}} = \text{Concentration ratio}$$

Because the concentrations of the isotopically labeled internal standards were the same in all solutions that were injected into the HPLC instrument, they do not need to be taken into consideration. However, the concentrations of the internal standard solutions should be in the range of the concentration of the sample solutions.

By means of the linear equation (calibration curve), the compound concentration in soil can be calculated as follows:

$$\text{Dilution}_{\text{Factor}} = \frac{\text{Volume}_{\text{final}}}{\text{Weight}}$$

$$\text{Conc}_{\text{Analyte}} = \frac{A_i}{A_{\text{IS}_i}}, \quad \text{Int. Ratio} = \frac{\text{Area}_{\text{Analyte}}}{\text{Area}_{\text{Internal Standard}}}$$

$$\text{Conc}_{\text{Soil}} = \text{Conc}_{\text{Analyte}} \cdot \text{Dilution}_{\text{Factor}}$$

$$\text{Volume}_{\text{final}} = \text{Final volume of the sample solution [L]}$$

$$\text{Weight} = \text{Weight of the soil sample [kg]}$$

$$\text{Conc}_{\text{Analyte}} = \text{Concentration of the analyte in the sample solution } [\mu\text{g/L}]$$

$$\text{Conc}_{\text{Soil}} = \text{Concentration of the analyte in soil } [\mu\text{g/kg}]$$

The recovery is calculated according to the following equation:

$$\text{Recovery} = \frac{\text{Conc}_{\text{Soil}} \cdot 100\%}{\text{Conc}_{\text{Soil Spiked}}}$$

$$\text{Conc}_{\text{Soil Spiked}} = \text{Concentration of the reference substance spiked } [\mu\text{g/kg}]$$

Example calculation for recovery of TI-435, 50 $\mu\text{g/kg}$ (sample ID 98-0109-10):

$$\text{Dilution}_{\text{Factor}} = \frac{0.05 \text{ L}}{0.02 \text{ kg}} = 2.5 \text{ L/kg}$$

$$\text{Dilution}_{\text{Factor}} = \frac{\text{Volume}_{\text{final}}}{\text{Weight}}$$

$$\text{Conc}_{\text{Analyte}} = \frac{0.391 - (-0.000159)}{0.019450} = 20.1 \mu\text{g/L}$$

$$\text{Conc}_{\text{Soil Wet}} = 20.1 \mu\text{g/L} \cdot 2.5 \text{ L/kg} = 50.3 \mu\text{g/kg}$$

$$\text{Recovery} = \frac{50.3 \frac{\mu\text{g}}{\text{kg}} \cdot 100\%}{50.0 \frac{\mu\text{g}}{\text{kg}}} = 101\%$$

$$\text{Conc}_{\text{Analyte}} = \frac{\text{Int. Ratio} - \text{Intercept}}{\text{Slope}}$$

$$\text{Conc}_{\text{Soil Wet}} = \text{Conc}_{\text{Analyte}} \cdot \text{Dilution}_{\text{Factor}}$$

$$\text{Recovery} = \frac{\text{Conc}_{\text{Soil}} \cdot 100\%}{\text{Conc}_{\text{Soil Spiked}}}$$

9.0 Detector Linearity

Standard solutions of TI-435, TZNG, TZMU, MNG and TMG were measured in a concentration range of 1.0 to 500 $\mu\text{g/L}$ (equivalent to 2.5 to 1250 $\mu\text{g/kg}$ samples). The concentrations of the internal standard solutions were constantly kept at 50 $\mu\text{g/L}$. In this concentration range, the mass spectrometric detector showed a linear correlation between concentration and intensity ratio (Table 10), with correlation coefficients ranging from 0.997 to 0.999.

Table 10: Correlation Coefficients for each Compound

| Substance | Correlation Coefficient |
|-----------|-------------------------|
| TI-435 | 0.998 |
| TZNG | 0.997 |
| TZMU | 0.999 |
| MNG | 0.998 |
| TMG | 0.999 |

10.0 Control and Reagent Blank Samples

The analytical results of duplicate control samples were below 1.5 $\mu\text{g/kg}$ for TI-435, TZNG, TZMU, MNG and TMG. Figure 2 shows a representative chromatogram of the analyte ions of control soil.

11.0 Limit of Quantification and Limit of Detection

The limit of quantification (LOQ) of the method, as determined from this validation, is 5 $\mu\text{g/kg}$ for TI-435 and its degradates.

The method detection limit (MDL) of the method is 0.95, 1.55, 0.90, 0.96, and 0.51 $\mu\text{g}/\text{kg}$ for TI-435, TZNG, TZMU, MNG and TMG, respectively (Tables 11-15). The MDL was calculated by multiplying the standard deviation by the appropriate Student's t value at a 99% confidence level (for $n = 5$, $t = 3.747$), as described in 40CFR Ch. 1, Part 136, App. B, (7-1-94 Edition).

12.0 Determination of the Recoverability

Soil from Watsonville, California was fortified with the active ingredient TI-435 and its degradates. Fortification levels of the different soils of TI-435, TZNG, TZMU, MNG and TMG as well as recoveries and relative standard deviations are presented in the following table (also see Table 11-15).

| Fortification [$\mu\text{g}/\text{kg}$] | N | Mean Values [%] | | | | |
|--|---|-----------------|-------|------|-------|-------|
| | | TI-435 | TZNG | TZMU | MNG | TMG |
| 5.0 | 5 | 100.4 | 100.9 | 97.3 | 91.9 | 91.7 |
| 50.0 | 5 | 100.3 | 93.4 | 95.9 | 111.1 | 101.1 |

Figures 2 to 4 show representative chromatograms of the analysis of control soil, fortifications made at 5 $\mu\text{g}/\text{kg}$ and fortifications made at 50 $\mu\text{g}/\text{kg}$.

13.0 Storage Stability of the Extracts

The stability of the analytes in the extracts was proven by the acceptable recoveries achieved for the sample extracts, after storage in a freezer at <0 $^{\circ}\text{C}$ for the period from extraction on April 4, 2000 to analysis on July 20, 2000 (107 days).

14.0 References

- Schramel, O., "Residue Analytical Method 00521 (MR-343/98) for Determination of TI-435 and the Metabolites TZNG, TZMU, MNG and TMG in Soil by Liquid Chromatography with Electrospray MS/MS-Detection," Bayer Report 109586, 1999.

Table 11: Analysis of Controls and Recovery Samples Fortified with TI-435.

| <u>Sample</u> | <u>Fortified Amount</u> <u>µg/kg</u> | <u>µg/L</u> | <u>µg/kg</u> | <u>% Recovery</u> | <u>Average Recovery</u> | <u>% Relative Std. Dev.</u> | <u>MDL</u> (StdDev*3.747) |
|----------------------|---|------------------|--------------|-------------------|-------------------------|-----------------------------|------------------------------|
| <u>TI-435</u> | | | | | | | |
| 98-0109-1 | 5.0 | 2.054 | 5.14 | 102.7 | | | |
| 98-0109-2 | 5.0 | 2.056 | 5.14 | 102.8 | | | |
| 98-0109-3 | 5.0 | 2.031 | 5.08 | 101.6 | | | |
| 98-0109-4 | 5.0 | 1.828 | 4.57 | 91.4 | | | |
| 98-0109-5 | 5.0 | 2.072 | 5.18 | 103.6 | 100.4 | 5.1 | |
| | | Average = | 5.02 | Std Dev = | 0.254 | | 0.95 |
| <u>TI-435</u> | | | | | | | |
| 98-0109-6 | 50.0 | 20.310 | 50.78 | 101.6 | | | |
| 98-0109-7 | 50.0 | 20.363 | 50.91 | 101.8 | | | |
| 98-0109-8 | 50.0 | 20.558 | 51.40 | 102.8 | | | |
| 98-0109-9 | 50.0 | 18.969 | 47.42 | 94.8 | | | |
| 98-0109-10 | 50.0 | 20.117 | 50.29 | 100.6 | 100.3 | 3.1 | |
| | | Average = | 50.16 | Std Dev = | 1.579 | | |

Table 12: Analysis of Controls and Recovery Samples Fortified with TZNG.

| <u>Sample</u> | <u>Fortified Amount</u> <u>µg/kg</u> | <u>µg/L</u> | <u>µg/kg</u> | <u>% Recovery</u> | <u>Average Recovery</u> | <u>% Relative Std. Dev.</u> | <u>MDL</u> (StdDev*3.747) |
|--------------------|---|------------------|--------------|-------------------|-------------------------|-----------------------------|------------------------------|
| <u>TZNG</u> | | | | | | | |
| 98-0109-1 | 5.0 | 2.237 | 5.59 | 111.9 | | | |
| 98-0109-2 | 5.0 | 1.970 | 4.93 | 98.5 | | | |
| 98-0109-3 | 5.0 | 2.138 | 5.35 | 106.9 | | | |
| 98-0109-4 | 5.0 | 1.893 | 4.73 | 94.7 | | | |
| 98-0109-5 | 5.0 | 1.847 | 4.62 | 92.4 | 100.9 | 8.2 | |
| | | Average = | 5.04 | Std Dev = | 0.414 | | 1.55 |
| <u>TZNG</u> | | | | | | | |
| 98-0109-6 | 50.0 | 17.378 | 43.45 | 86.9 | | | |
| 98-0109-7 | 50.0 | 17.631 | 44.08 | 88.2 | | | |
| 98-0109-8 | 50.0 | 19.809 | 49.52 | 99.0 | | | |
| 98-0109-9 | 50.0 | 19.351 | 48.38 | 96.8 | | | |
| 98-0109-10 | 50.0 | 19.212 | 48.03 | 96.1 | 93.4 | 5.9 | |
| | | Average = | 46.69 | Std Dev = | 2.740 | | |

Table 13: Analysis of Controls and Recovery Samples Fortified with TZMU.

| <u>Sample</u> | <u>Fortified Amount</u> <u>µg/kg</u> | <u>µg/L</u> | <u>µg/kg</u> | <u>% Recovery</u> | <u>Average Recovery</u> | <u>% Relative Std. Dev.</u> | <u>MDL</u> (StdDev*3.747) |
|--------------------|---|------------------|--------------|-------------------|-------------------------|-----------------------------|------------------------------|
| <u>TZMU</u> | | | | | | | |
| 98-0109-1 | 5.0 | 1.914 | 4.79 | 95.7 | | | |
| 98-0109-2 | 5.0 | 2.094 | 5.24 | 104.7 | | | |
| 98-0109-3 | 5.0 | 1.881 | 4.70 | 94.1 | | | |
| 98-0109-4 | 5.0 | 1.987 | 4.97 | 99.4 | | | |
| 98-0109-5 | 5.0 | 1.855 | 4.64 | 92.8 | 97.3 | 4.9 | |
| | | Average = | 4.87 | Std Dev = | 0.241 | | 0.90 |
| <u>TZMU</u> | | | | | | | |
| 98-0109-6 | 50.0 | 19.768 | 49.42 | 98.8 | | | |
| 98-0109-7 | 50.0 | 19.190 | 47.98 | 96.0 | | | |
| 98-0109-8 | 50.0 | 19.905 | 49.76 | 99.5 | | | |
| 98-0109-9 | 50.0 | 17.750 | 44.38 | 88.8 | | | |
| 98-0109-10 | 50.0 | 19.265 | 48.16 | 96.3 | 95.9 | 4.5 | |
| | | Average = | 47.94 | Std Dev = | 2.137 | | |

Table 14: Analysis of Controls and Recovery Samples Fortified with MNG.

| <u>Sample</u> | <u>Fortified Amount</u> <u>µg/kg</u> | <u>µg/L</u> | <u>µg/kg</u> | <u>% Recovery</u> | <u>Average Recovery</u> | <u>% Relative Std. Dev.</u> | <u>MDL</u> (StdDev*3.747) |
|-------------------|---|------------------|--------------|-------------------|-------------------------|-----------------------------|------------------------------|
| <u>MNG</u> | | | | | | | |
| 98-0109-1 | 5.0 | 1.764 | 4.41 | 88.2 | | | |
| 98-0109-2 | 5.0 | 1.828 | 4.57 | 91.4 | | | |
| 98-0109-3 | 5.0 | 2.016 | 5.04 | 100.8 | | | |
| 98-0109-4 | 5.0 | 1.806 | 4.52 | 90.3 | | | |
| 98-0109-5 | 5.0 | 1.780 | 4.45 | 89.0 | 91.9 | 5.5 | |
| | | Average = | 4.60 | Std Dev = | 0.255 | | 0.96 |
| <u>MNG</u> | | | | | | | |
| 98-0109-6 | 50.0 | 22.180 | 55.45 | 110.9 | | | |
| 98-0109-7 | 50.0 | 22.959 | 57.40 | 114.8 | | | |
| 98-0109-8 | 50.0 | 21.945 | 54.86 | 109.7 | | | |
| 98-0109-9 | 50.0 | 22.096 | 55.24 | 110.5 | | | |
| 98-0109-10 | 50.0 | 21.880 | 54.70 | 109.4 | 111.1 | 2.0 | |
| | | Average = | 55.53 | Std Dev = | 1.085 | | |

Table 15: Analysis of Controls and Recovery Samples Fortified with TMG.

| <u>Sample</u> | <u>Fortified Amount</u> <u>µg/kg</u> | <u>µg/L</u> | <u>µg/kg</u> | <u>% Recovery</u> | <u>Average Recovery</u> | <u>% Relative Std. Dev.</u> | <u>MDL</u> (StdDev*3.747) |
|-------------------|---|------------------|--------------|-------------------|-------------------------|-----------------------------|------------------------------|
| <u>TMG</u> | | | | | | | |
| 98-0109-1 | 5.0 | 1.850 | 4.63 | 92.5 | | | |
| 98-0109-2 | 5.0 | 1.757 | 4.39 | 87.9 | | | |
| 98-0109-3 | 5.0 | 1.832 | 4.58 | 91.6 | | | |
| 98-0109-4 | 5.0 | 1.825 | 4.56 | 91.3 | | | |
| 98-0109-5 | 5.0 | 1.910 | 4.78 | 95.5 | 91.7 | 3.0 | |
| | | Average = | 4.59 | Std Dev = | 0.137 | | 0.51 |
| <u>TMG</u> | | | | | | | |
| 98-0109-6 | 50.0 | 22.431 | 56.08 | 112.2 | | | |
| 98-0109-7 | 50.0 | 18.817 | 47.04 | 94.1 | | | |
| 98-0109-8 | 50.0 | 18.873 | 47.18 | 94.4 | | | |
| 98-0109-9 | 50.0 | 19.703 | 49.26 | 98.5 | | | |
| 98-0109-10 | 50.0 | 21.295 | 53.24 | 106.5 | 101.1 | 7.9 | |
| | | Average = | 50.56 | Std Dev = | 3.971 | | |

Figure 1: Flow Diagram of Analysis Procedure

Mix 20 g of soil with 4 g of Hydromatrix in a beaker,
fill into an ASE 200 extraction cell



Extract with approx. 45 mL of acetonitrile:water:acetic acid:guanidine hydrochloride
(25:100:0.1:1; v:v:w), on an ASE Extractor at 140 °C



Add 1 mL of internal standard and adjust the volume
to 50mL with 1:4 acetonitrile:water



Fill into a HPLC vial



HPLC-MS/MS (APCI)

Figure 2: LC-MS/MS Chromatogram of Control sample, 98-0109-C.

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Printed: Thu, Jul 20, 2000 19:44

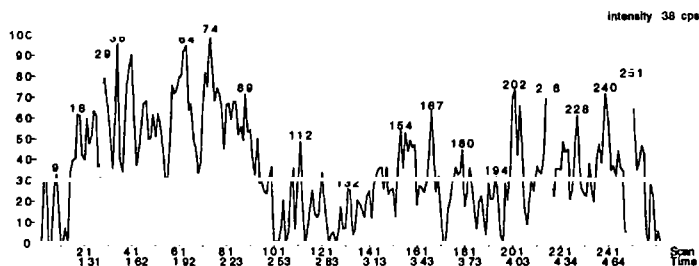
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Calibration File: CFA072000 Path: LS-038 MAC #1 Data Bayer/Takeda.A072000.

Comments: No Comments

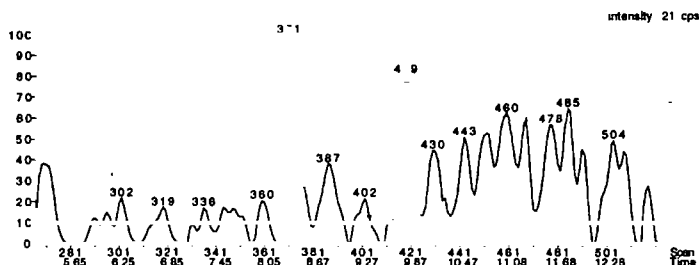
A072000009 A072000009 Thu, Jul 20, 2000 12:29
98-0109-C

13.03 in 2 periods
MNG
Internal Standard MNG-d3
Use Area
Absolute Retention Time
1 4.00 MRM, 266 scans
117 0->80.7
Noise Thres 10.0
Quant Thres 1.0
Min Width 6
Mult Width 6
Base Width 60
RT Win (secs) 20
Smooth 1
Expected RT 3.87
Area 0
Height 0
Start Time -0.00
End Time -0.00
Integration Width -0.00
Retention Time 0.00
Integration Type



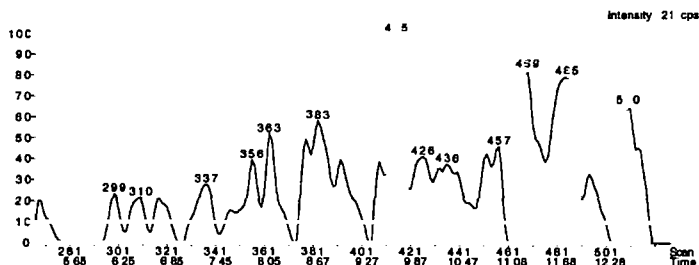
A072000009 A072000009 Thu Jul 20 2000 12:29
98-0109-C

13.03 in 2 periods
TMG
Internal Standard TMG-d3
Use Area
Absolute Retention Time
2 7.80 MRM, 260 scans
206 0->132.0
Noise Thres 5.0
Quant Thres 0.2
Min Width 6
Mult Width 9
Base Width 60
RT Win (secs) 20
Smooth 5
Expected RT 9.91
Area 0
Height 0
Start Time -0.00
End Time -0.00
Integration Width -0.00
Retention Time 0.00
Integration Type



A072000009 A072000009 Thu, Jul 20, 2000 12:29
98-0109-C

13.03 in 2 periods
TZMU
Internal Standard TZMU-d3
Use Area
Absolute Retention Time
2 7.80 MRM, 260 scans
206 0->176.0
Noise Thres 5.0
Quant Thres 0.2
Min Width 6
Mult Width 6
Base Width 60
RT Win (secs) 20
Smooth 5
Expected RT 8.92
Area 0
Height 0
Start Time -0.00
End Time -0.00
Integration Width -0.00
Retention Time -0.00
Integration Type



A072000009 A072000009 Thu, Jul 20, 2000 12:29
98-0109-C

13.03 in 2 periods
TZNG
Internal Standard TZNG-C13 N15
Use Area
Absolute Retention Time
2 7.80 MRM, 260 scans
236 0->132.0
Noise Thres 10.0
Quant Thres 0.2
Min Width 6
Mult Width 6
Base Width 60
RT Win (secs) 20
Smooth 5
Expected RT 9.37
Area 0
Height 0
Start Time 0.00
End Time 0.00
Integration Width 0.00
Retention Time -0.00
Integration Type

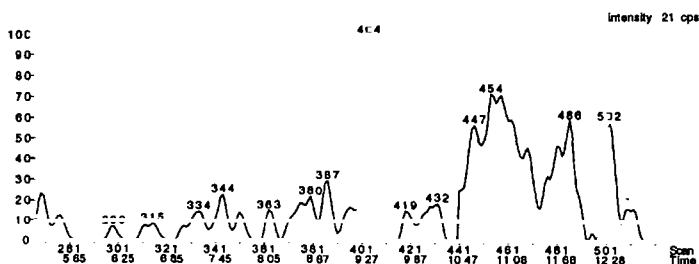


Figure 2: LC-MS/MS Chromatogram of Control sample, 98-0109-C. (Continued)

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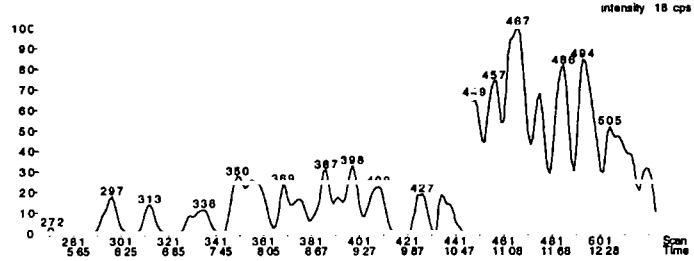
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Calibration File CFA072000 Path LS-038 MAC #1 Data Bayer/Takeda A072000:

Comments. No Comments

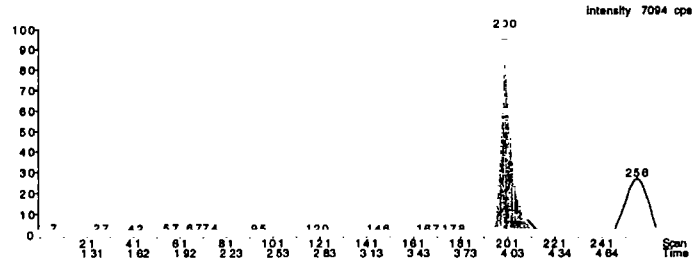
A072000009 A072000009 Thu Jul 20 2000 12 29
98-0109-C

13.03 in 2 periods
TI-435
Internal Standard TI-435-d3
Use Area
Absolute Retention Time
2 7.80 MRM, 260 scans
250 0->169 0
Noise Thres 5.0
Quant Thres 0.5
Min Width 6
Mult Width 6
Base Width 80
RT Win (secs) 20
Smooth 5
Expected RT 9.82
Area 0
Height 0
Start Time 0.00
End Time -0.00
Integration Width -0.00
Retention Time 0.00
Integration Type



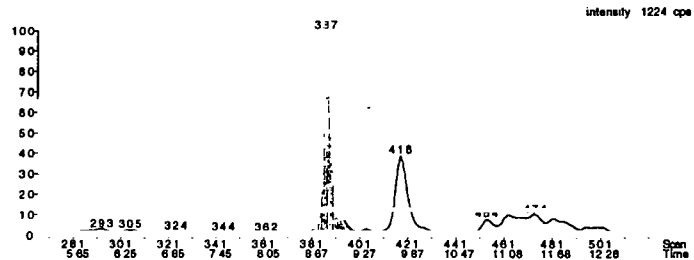
A072000009 A072000009 Thu Jul 20 2000 12 29
98-0109-C

13.03 in 2 periods
MNG-d3
use as Internal Standard
1 4.00 MRM, 266 scans
120 0->80 7
Noise Thres 10.0
Quant Thres 0.5
Min Width 6
Mult Width 8
Base Width 80
RT Win (secs) 20
Smooth 5
Expected RT 3.88
Area 46274
Height 7094
Start Time 3.91
End Time 4.29
Integration Width 0.38
Retention Time 4.02
Integration Type A - BB



A072000009 A072000009 Thu Jul 20 2000 12 29
98-0109-C

13.03 in 2 periods
TMG-d3
use as Internal Standard
2 7.80 MRM, 260 scans
208 0->132 0
Noise Thres 10.0
Quant Thres 1.0
Min Width 6
Mult Width 8
Base Width 80
RT Win (secs) 20
Smooth 5
Expected RT 8.66
Area 12899
Height 1220
Start Time 8.64
End Time 9.45
Integration Width 0.81
Retention Time 8.85
Integration Type A - BB



A072000009 A072000009 Thu Jul 20 2000 12 29
98-0109-C

13.03 in 2 periods
TZNU-d3
use as Internal Standard
2 7.80 MRM, 260 scans
209 0->175 0
Noise Thres 10.0
Quant Thres 1.0
Min Width 6
Mult Width 8
Base Width 80
RT Win (secs) 20
Smooth 5
Expected RT 8.89
Area 26981
Height 2443
Start Time 8.70
End Time 9.38
Integration Width 0.69
Retention Time 8.80
Integration Type A - BV

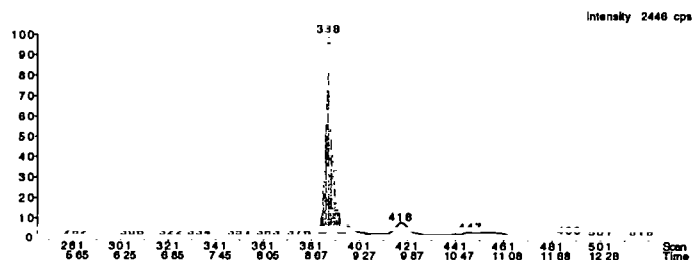


Figure 2: LC-MS/MS Chromatogram of Control sample, 98-0109-C. (Continued)

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Printed: Thu, Jul 20, 2000 19:44

Calibration File CFA072000 Path. LS-038 MAC #1 Data Bayer/Takeda A072000

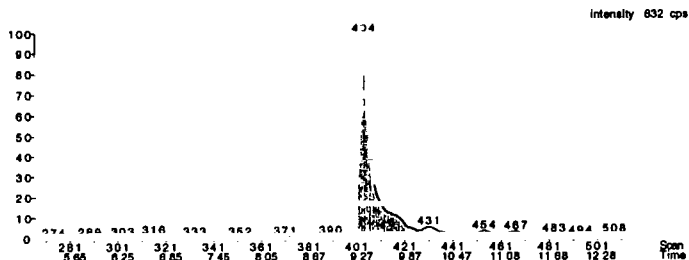
Comments: No Comments

A072000009 A072000009 Thu Jul 20, 2000 12:29
98-0109-C

13.03 in 2 periods
TZNG-C13 N15
use as Internal Standard

2 7.60 MRM, 260 scans
240 0->134 0
Noise Thresh 10.0
Quant Thresh 1.0
Min Width 6
Mult Width 8
Base Width 60
RT Win (secs) 2.0
Smooth 5
Expected RT 9.37

Area 7934
Height 629
Start Time 9.18
End Time 10.02
Integration Width 0.84
Retention Time 9.38
Integration Type A - BV



A072000009 A072000009 Thu, Jul 20, 2000 12:29
98-0109-C

13.03 in 2 periods
T1-435-d3
use as Internal Standard

2 7.60 MRM, 260 scans
253 0->172 0
Noise Thresh 10.0
Quant Thresh 1.0
Min Width 6
Mult Width 8
Base Width 60
RT Win (secs) 2.0
Smooth 5
Expected RT 9.79

Area 53285
Height 4368
Start Time 9.67
End Time 11.14
Integration Width 1.56
Retention Time 9.78
Integration Type A - BV

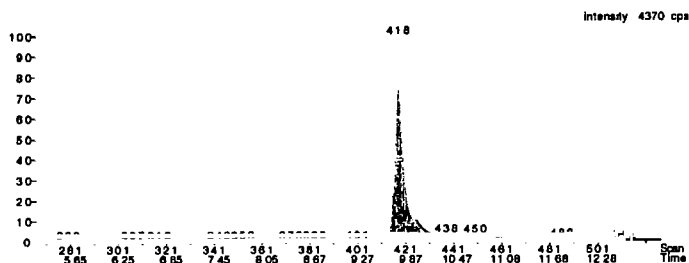


Figure 3: LC-MS/MS Chromatogram of 5 ppb fortified sample, 98-0109-1.

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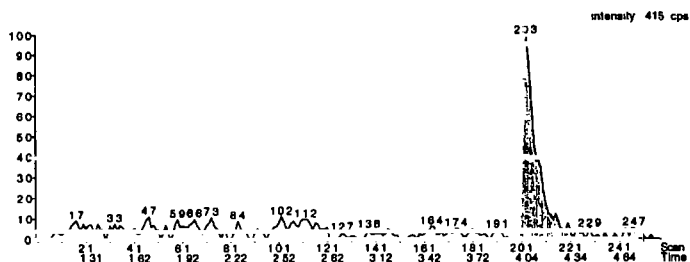
Printed: Thu, Jul 20, 2000 19:44 *TL*

Calibration File CFA072000 Path LS-038 MAC #1 Data Bayer/Takeda A072000

Comments No Comments

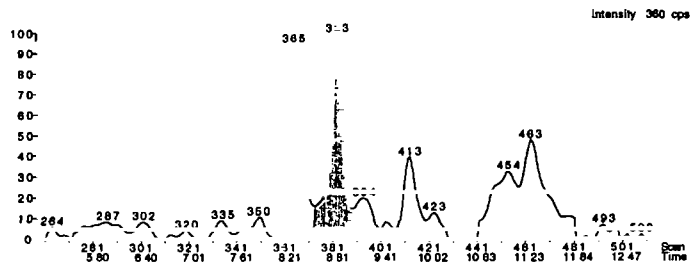
A072000012 A072000012 Thu, Jul 20, 2000 19:27
98-0109-1

12 92 in 2 periods
MNG
Internal Standard MNG-d3
Use Area
Absolute Retention Time
1 3 87 MRM 257 scans
117 0 -> 80 7
Noise Thres 5 0
Quant Thres 0 5
Min Width 6
Mult Width 6
Base Width 80
RT Win (secs) 20
Smooth 1
Expected RT 3 87
Area 2823
Height 415
Start Time 3 99
End Time 4 34
Integration Width 0 35
Retention Time 4 07
Integration Type A - BB



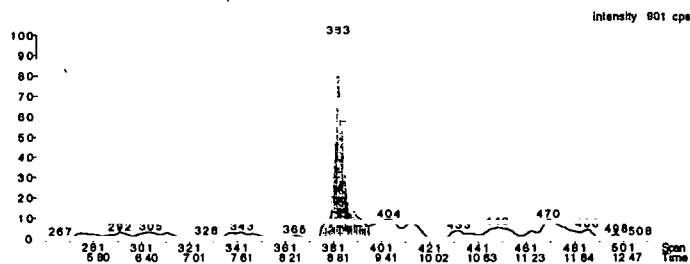
A072000012 A072000012 Thu, Jul 20, 2000 19:27
98-0109 1

12 92 in 2 periods
TMG
Internal Standard TMG-d3
Use Area
Absolute Retention Time
2 7 80 MRM 259 scans
205 0 -> 132 0
Noise Thres 5 0
Quant Thres 1 0
Min Width 6
Mult Width 6
Base Width 40
RT Win (secs) 20
Smooth 5
Expected RT 8 91
Area 4087
Height 343
Start Time 8 57
End Time 9 02
Integration Width 0 45
Retention Time 8 87
Integration Type A - VV



A072000012 A072000012 Thu, Jul 20, 2000 19:27
98-0109 1

12 92 in 2 periods
TZMU
Internal Standard TZMU-d3
Use Area
Absolute Retention Time
2 7 80 MRM 259 scans
206 0 -> 175 0
Noise Thres 5 0
Quant Thres 0 2
Min Width 6
Mult Width 6
Base Width 60
RT Win (secs) 20
Smooth 5
Expected RT 8 92
Area 10511
Height 900
Start Time 8 83
End Time 9 23
Integration Width 0 60
Retention Time 8 87
Integration Type A - BV



A072000012 A072000012 Thu, Jul 20, 2000 19:27
98 0109-1

12 92 in 2 periods
TZNG
Internal Standard TZNG-C13,N15
Use Area
Absolute Retention Time
2 7 80 MRM 259 scans
238 0 -> 132 0
Noise Thres 10 0
Quant Thres 0 2
Min Width 6
Mult Width 6
Base Width 80
RT Win (secs) 20
Smooth 5
Expected RT 9 37
Area 7417
Height 564
Start Time 9 14
End Time 10 24
Integration Width 1 09
Retention Time 9 32
Integration Type A - BB

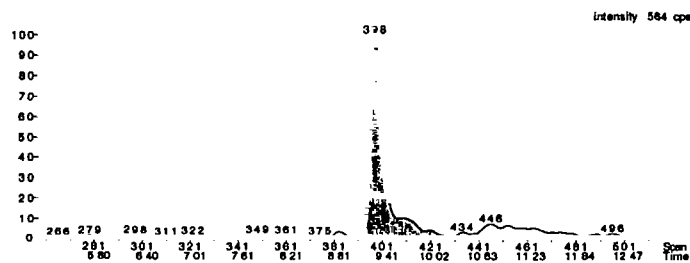
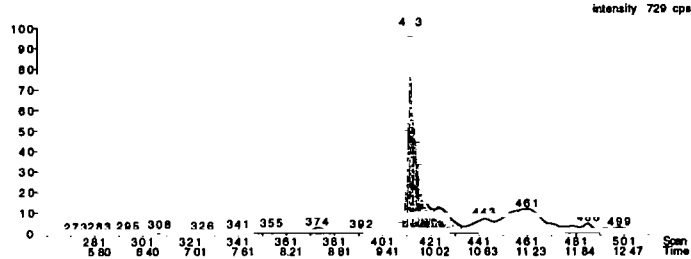


Figure 3: LC-MS/MS Chromatogram of 5 ppb fortified sample, 98-0109-1. (Continued)

MacQuan, version 1.5
 Printed Thu, Jul 20, 2000 19:44 *TK*
 Calibration File CFA072000 Path. LS-038 MAC #1 Data Bayer/Takeda A072000
 Comments No Comments

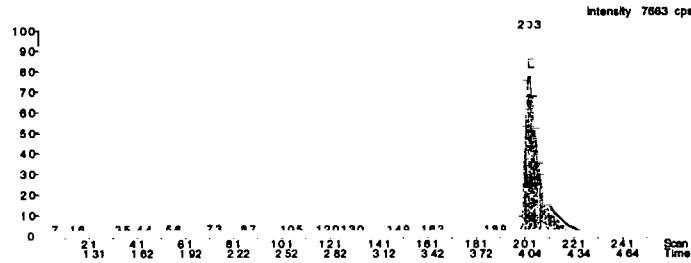
A072000012 A072000012 Thu, Jul 20, 2000 19:27
 98-0109-1

12.92 in 2 periods
 TI-435
 Internal Standard TI-435-d3
 Use Area
 Absolute Retention Time
 2 7.80 MRM 259 scans
 250 0->189 0
 Noise Thres 5 0
 Quant Thres 0 5
 Min Width 6
 Mult Width 6
 Base Width 80
 RT Win (secs) 20
 Smooth 5
 Expected RT 9.82
 Area 9325
 Height 726
 Start Time 9.59
 End Time 10.39
 Integration Width 0.79
 Retention Time 9.78
 Integration Type A BV



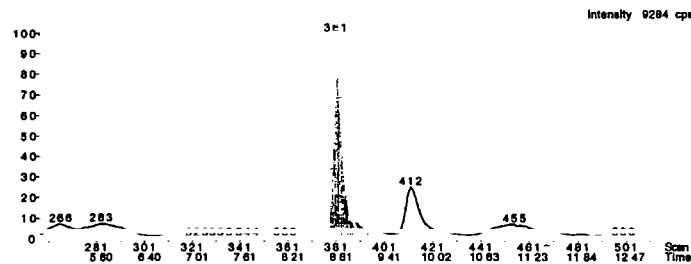
A072000012 A072000012 Thu, Jul 20, 2000 19:27
 98-0109-1

12.92 in 2 periods
 MNG-d3
 use as Internal Standard
 1 3.87 MRM, 257 scans
 120 0->60 7
 Noise Thres 10 0
 Quant Thres 0 5
 Min Width 6
 Mult Width 8
 Base Width 80
 RT Win (secs) 20
 Smooth 5
 Expected RT 3.88
 Area 61503
 Height 7863
 Start Time 3.96
 End Time 4.55
 Integration Width 0.59
 Retention Time 4.07
 Integration Type A BB



A072000012 A072000012 Thu, Jul 20, 2000 19:27
 98-0109-1

12.92 in 2 periods
 TMG-d3
 use as Internal Standard
 2 7.80 MRM, 259 scans
 208 0->132 0
 Noise Thres 10 0
 Quant Thres 1 0
 Min Width 6
 Mult Width 8
 Base Width 80
 RT Win (secs) 20
 Smooth 5
 Expected RT 8.86
 Area 108011
 Height 9277
 Start Time 8.63
 End Time 9.50
 Integration Width 0.87
 Retention Time 8.81
 Integration Type A - BV



A072000012 A072000012 Thu, Jul 20, 2000 19:27
 98-0109-1

12.92 in 2 periods
 TZMU-d3
 use as Internal Standard
 2 7.80 MRM 259 scans
 209 0->175 0
 Noise Thres 10 0
 Quant Thres 1 0
 Min Width 6
 Mult Width 8
 Base Width 80
 RT Win (secs) 20
 Smooth 5
 Expected RT 8.89
 Area 248594
 Height 21564
 Start Time 8.66
 End Time 9.58
 Integration Width 0.90
 Retention Time 8.84
 Integration Type A BV

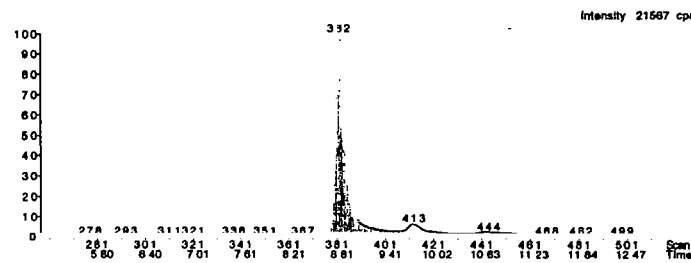


Figure 3: LC-MS/MS Chromatogram of 5 ppb fortified sample, 98-0109-1. (Continued)

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Printed: Thu, Jul 20, 2000 19:44 *tbl*

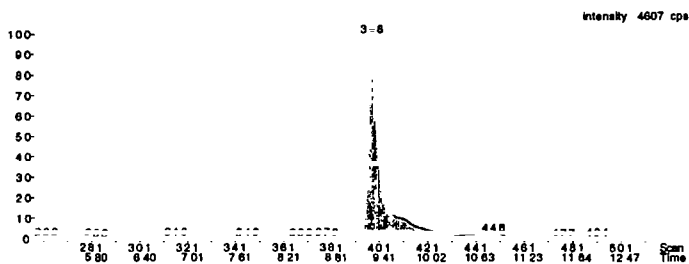
Calibration File CFA072000 Path LS-038 MAC #1 Data Bayer/Takeda A072000

Comments: No Comments

A072000012 A072000012 Thu Jul 20, 2000 19:27
98-0109-1

12.92 in 2 periods
TZNG C13,N16
use as Internal Standard

2.780 MRM, 259 scans
240.0->134.0
Noise Thresh 10.0
Quant Thresh 1.0
Min Width 6
Mul Width 6
Base Width 60
RT Win (secs) 2.0
Smooth 5
Expected RT 9.37
Area 62627
Height 4605
Start Time 9.14
End Time 10.42
Integration Width 1.27
Retention Time 9.32
Integration Type A-BV



A072000012 A072000012 Thu Jul 20, 2000 19:27
98-0109-1

12.92 in 2 periods
TI-436-d3
use as Internal Standard

2.780 MRM, 259 scans
253.0->172.0
Noise Thresh 10.0
Quant Thresh 1.0
Min Width 6
Mul Width 6
Base Width 80
RT Win (secs) 2.0
Smooth 5
Expected RT 9.79
Area 234325
Height 19415
Start Time 9.53
End Time 10.54
Integration Width 1.00
Retention Time 9.75
Integration Type A-BV

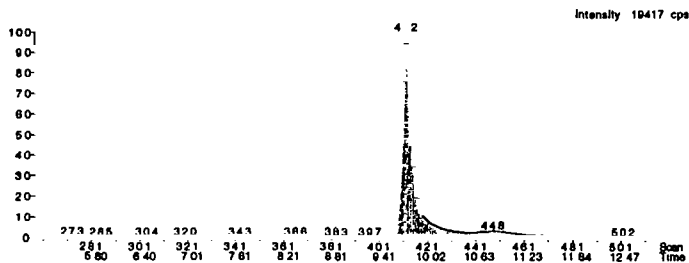


Figure 4: LC-MS/MS Chromatogram of 50 ppb fortified sample, 98-0109-10.

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Printed Thu, Jul 20, 2000 19:44

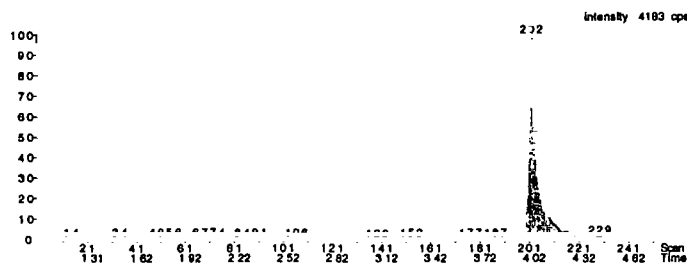
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Calibration File CFA072000 Path. LS-038 MAC #1 Data Bayer/Takeda A072000

Comments: No Comments

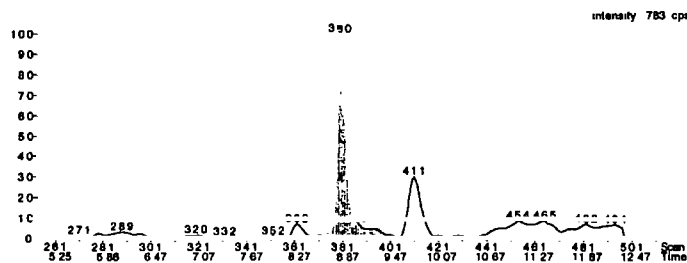
A072000022 A072000022 Thu, Jul 20, 2000 18:19
98-0109-10

12.82 in 2 periods
MNG
Internal Standard MNG-d3
Use Area
Absolute Retention Time
1 3.79 MRM, 253 scans
117 0->80 7
Noise Three 10 0
Quant Three 1 0
Min Width 6
Mult Width 6
Base Width 80
RT Win (secs) 20
Smooth 1
Expected RT 3.87
Area 19909
Height 4183
Start Time 3.96
End Time 4.37
Integration Width 0.41
Retention Time 4.04
Integration Type A - BB



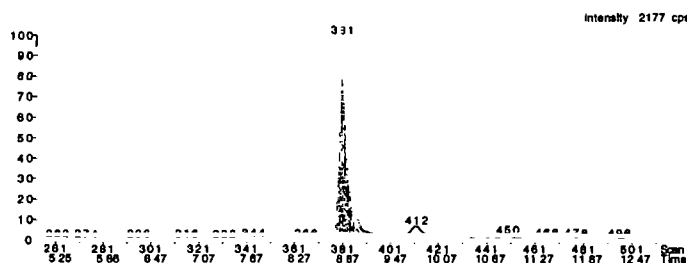
A072000022 A072000022 Thu, Jul 20, 2000 18:19
98-0109-10

12.82 in 2 periods
TMG
Internal Standard TMG-d3
Use Area
Absolute Retention Time
2 7.77 MRM, 259 scans
205 0->132 0
Noise Three 5 0
Quant Three 0 2
Min Width 6
Mult Width 9
Base Width 80
RT Win (secs) 20
Smooth 5
Expected RT 8.91
Area 8622
Height 783
Start Time 8.88
End Time 9.53
Integration Width 0.67
Retention Time 8.84
Integration Type A - BB



A072000022 A072000022 Thu, Jul 20, 2000 18:19
98-0109-10

12.82 in 2 periods
TZMU
Internal Standard TZMU-d3
Use Area
Absolute Retention Time
2 7.77 MRM, 259 scans
205 0->175 0
Noise Three 5 0
Quant Three 0 2
Min Width 6
Mult Width 9
Base Width 80
RT Win (secs) 20
Smooth 5
Expected RT 8.92
Area 23425
Height 2176
Start Time 8.69
End Time 9.44
Integration Width 0.75
Retention Time 8.67
Integration Type A - BV



A072000022 A072000022 Thu, Jul 20, 2000 18:19
98-0109-10

12.82 in 2 periods
TZNG
Internal Standard TZNG-C13,N15
Use Area
Absolute Retention Time
2 7.77 MRM, 259 scans
236 0->132 0
Noise Three 10 0
Quant Three 0 2
Min Width 6
Mult Width 8
Base Width 80
RT Win (secs) 20
Smooth 5
Expected RT 9.37
Area 17153
Height 1270
Start Time 9.17
End Time 10.64
Integration Width 1.47
Retention Time 9.32
Integration Type A - BV

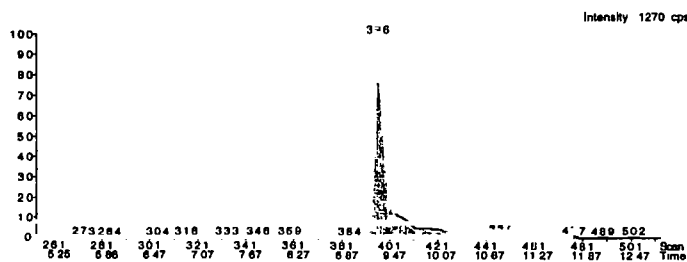


Figure 4: LC-MS/MS Chromatogram of 50 ppb fortified sample, 98-0109-10. (Continued)

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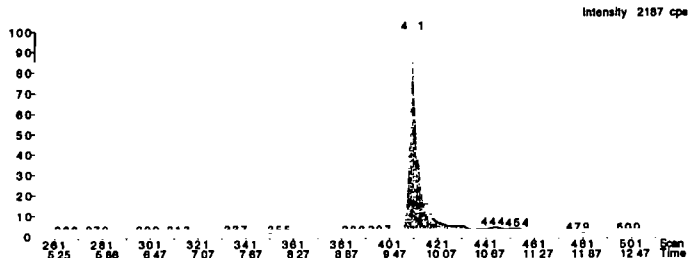
Printed Thu, Jul 20, 2000 19 44 *TK*

Calibration File: CFA072000 Path LS-038 MAC #1.Data Bayer/Takeda A072000

Comments: No Comments

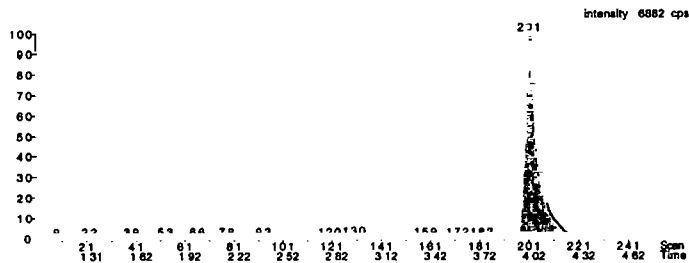
A072000022 A072000022 Thu, Jul 20, 2000 18 19
98-0109-10

12.82 in 2 periods
TI-435
Internal Standard TI-435-d3
Use Area
Absolute Retention Time
2 7.77 MRM 259 scans
250 0 >169 0
Noise Thres 5.0
Quant Thres 0.5
Min Width 8
Mult Width 8
Base Width 80
RT Win (secs) 2.0
Smooth 5
Expected RT 9.82
Area 26528
Height 2188
Start Time 9.58
End Time 10.49
Integration Width 0.93
Retention Time 9.77
Integration Type A BV



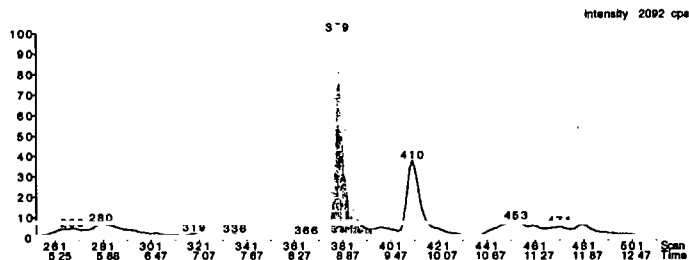
A072000022 A072000022 Thu, Jul 20, 2000 18 19
98 0109-10

12.82 in 2 periods
MNG-d3
use as Internal Standard
1 3.79 MRM, 253 scans
120 0 >80 7
Noise Thres 10.0
Quant Thres 0.5
Min Width 8
Mult Width 8
Base Width 80
RT Win (secs) 2.0
Smooth 5
Expected RT 3.88
Area 44318
Height 6882
Start Time 3.95
End Time 4.32
Integration Width 0.38
Retention Time 4.02
Integration Type A BB



A072000022 A072000022 Thu, Jul 20, 2000 18 19
98-0109-10

12.82 in 2 periods
TMG-d3
use as Internal Standard
2 7.77 MRM 259 scans
208 0 >132 0
Noise Thres 10.0
Quant Thres 1.0
Min Width 8
Mult Width 8
Base Width 80
RT Win (secs) 2.0
Smooth 5
Expected RT 8.86
Area 22829
Height 2087
Start Time 8.63
End Time 9.20
Integration Width 0.57
Retention Time 8.81
Integration Type A BV



A072000022 A072000022 Thu, Jul 20, 2000 18 19
98-0109-10

12.82 in 2 periods
TZMU-d3
use as Internal Standard
2 7.77 MRM, 259 scans
209 0 >175 0
Noise Thres 10.0
Quant Thres 1.0
Min Width 8
Mult Width 8
Base Width 80
RT Win (secs) 2.0
Smooth 5
Expected RT 8.89
Area 54708
Height 4865
Start Time 8.66
End Time 9.47
Integration Width 0.81
Retention Time 8.84
Integration Type A BV

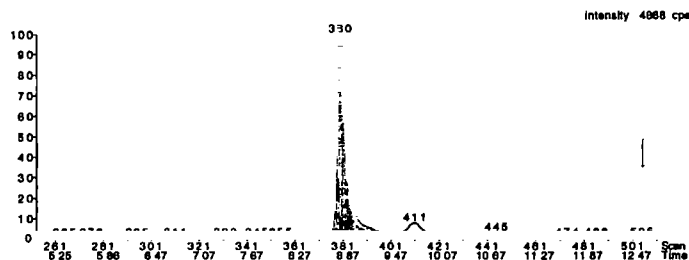


Figure 4: LC-MS/MS Chromatogram of 50 ppb fortified sample, 98-0109-10. (Continued)

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TW

Calibration File: CFA072000 Path: LS-038 MAC #1:Data.Bayer/Takeda:A072000:

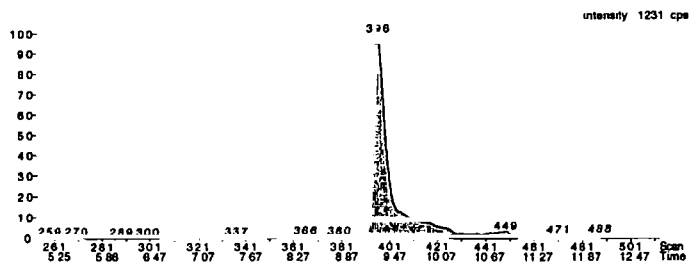
Comments No Comments

A072000022 A072000022 Thu, Jul 20, 2000 18:19
98-0109-10

12.82 in 2 periods
TZNG-C13 N15
use as Internal Standard

2 7.77 MRM, 259 scans
240 0->134 0
Noise Thres 10.0
Quant Thres 1.0
Min Width 6
Mult Width 6
Base Width 60
RT Win (secs) 20
Smooth 5
Expected RT 9.37

Area 17560
Height 1230
Start Time 9.14
End Time 10.43
Integration Width 1.28
Retention Time 9.32
Integration Type A - BV

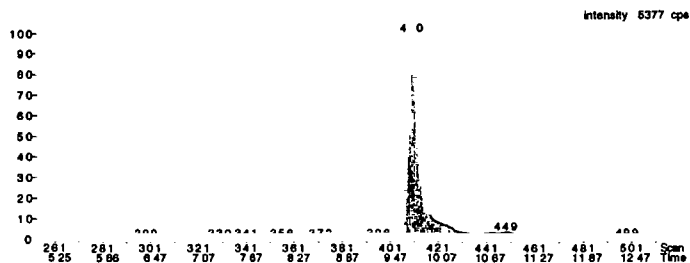


A072000022 A072000022 Thu, Jul 20, 2000 18:19
98-0109-10

12.82 in 2 periods
T1 435-03
use as Internal Standard

2 7.77 MRM, 259 scans
253 0->172 0
Noise Thres 10.0
Quant Thres 1.0
Min Width 6
Mult Width 6
Base Width 80
RT Win (secs) 20
Smooth 5
Expected RT 9.79

Area 67822
Height 5375
Start Time 9.56
End Time 10.55
Integration Width 0.99
Retention Time 9.74
Integration Type A - BV



Appendix 1: Standard Preparation

The following are general working procedures for the preparation of standard solutions. Therefore, given weights and volumes do not necessarily correspond exactly to the weights, volumes and concentrations documented in the raw data.

Solvent mixtures

Solvent A: acetonitrile/water (1/1; v/v); **Solvent B:** acetonitrile/water (1/4; v/v)

Standard Stock Solutions

400 mg/L stock solution of TI-435: Weigh approximately 10 ± 0.01 mg of TI-435 into a 25-mL volumetric flask. Dilute to volume with Solvent A.

400 mg/L stock solution of TZNG: Weigh approximately 10 ± 0.01 mg of TZNG into a 25-mL volumetric flask. Dilute to volume with Solvent A.

400 mg/L stock solution of TZMU: Weigh approximately 10 ± 0.01 mg of TZMU into a 25-mL volumetric flask. Dilute to volume with Solvent A.

400 mg/L stock solution of MNG: Weigh approximately 10 ± 0.01 mg of MNG into a 25-mL volumetric flask. Dilute to volume with Solvent A.

400 mg/L stock solution of TMG: Weigh approximately 10 ± 0.01 mg of TMG into a 25-mL volumetric flask. Dilute to volume with Solvent A.

200 mg/L stock solution of d_3 -TI-435: Weigh approximately 5 ± 0.01 mg of d_3 -TI-435 into a 25-mL volumetric flask. Dilute to volume with Solvent A.

200 mg/L stock solution of ^{13}C , ^{15}N -TZNG: Weigh approximately 5 ± 0.01 mg of ^{13}C , ^{15}N -TZNG into a 25-mL volumetric flask. Dilute to volume with Solvent A.

200 mg/L stock solution of d_3 -TZMU: Weigh approximately 5 ± 0.01 mg of d_3 -TZMU into a 25-mL volumetric flask. Dilute to volume with Solvent A.

200 mg/L stock solution of d_3 -TMG: Aliquot approximately 5 ± 0.01 mg of d_3 -TMG into a 25-mL volumetric flask. Dilute to volume with Solvent A.

200 mg/L stock solution of d_3 -MNG: Weigh approximately 5 ± 0.01 mg of d_3 -MNG into a 25-mL volumetric flask. Dilute to volume with Solvent A.

Remark: Before further use, the standard stock solutions should be ultrasonicated for about one minute to achieve complete dissolution of the compounds.

Internal Standard Solution

(MIX1) 2.5 mg/L of each internal standard:

Pipette 2.5 mL of each of the stock solutions of d_3 -TI-435, ^{13}C , ^{15}N -TZNG, d_3 -TZMU, d_3 -TMG and d_3 -MNG into a 200-mL volumetric flask and dilute to volume with Solvent B.

Appendix 1 (cont.): Standard Preparation

Fortification Standard Solution

(MIX2) 1 mg/L of each compound: Pipette 0.25 mL of the stock solutions of TI-435, TZNG, TZMU, TMG and MNG into a 100-mL volumetric flask and dilute to volume with Solvent B.

Mixed Native Standard Solution

(MIX3) 10 mg/L of each compound: Pipette 2.5 mL of the stock solutions of TI-435, TZNG, TZMU, TMG and MNG into a 100-mL volumetric flask and dilute to volume with Solvent B.

Calibration Standard Solutions

(MIX4) 0.5 mg/L of each compound (1250 µg/kg sample equivalents) and 0.05 mg/L of each internal standard: Pipette 5 mL of the mixed native standard solution MIX3 and 2 mL of the internal standard solution MIX1 into a 100-mL volumetric flask and dilute to volume with Solvent B.

(MIX5) 0.25 mg/L of each compound (625 µg/kg sample equivalents) and 0.05 mg/L of each internal standard: Pipette 2.5 mL of the mixed native standard solutions MIX3 and 2 mL of the internal standard solution MIX1 into a 100-mL volumetric flask and dilute to volume with Solvent B.

(MIX6) 0.1 mg/L of each compound (250 µg/kg sample equivalents) and 0.05 mg/L of each internal standard: Pipette 1 mL of the mixed native standard solutions MIX3 and 2 mL of the internal standard solution MIX1 into a 100-mL volumetric flask and dilute to volume with Solvent B.

(MIX7) 0.05 mg/L of each compound (125 µg/kg sample equivalents) and 0.05 mg/L of each internal standard: Pipette 0.5 mL of the mixed native standard solutions MIX3 and 2 mL of the internal standard solution MIX1 into a 100-mL volumetric flask and dilute to volume with Solvent B.

(MIX8) 0.025 mg/L of each compound (62.5 µg/kg sample equivalents) and 0.05 mg/L of each internal standard: Pipette 0.25 mL of the mixed native standard solutions MIX3 and 2 mL of the internal standard solution MIX1 into a 100-mL volumetric flask and dilute to volume with Solvent B.

(MIX9) 0.01 mg/L of each compound (25 µg/kg sample equivalents) and 0.05 mg/L of each internal standard: Pipette 1 mL of the fortification standard solutions MIX2 and 2 mL of the internal standard solution MIX1 into a 100-mL volumetric flask and dilute to volume with Solvent B.

(MIX10) 0.005 mg/L of each compound (12.5 µg/kg sample equivalents) and 0.05 mg/L of each internal standard: Pipette 0.5 mL of the fortification standard solutions MIX2 and 2 mL of the internal standard solution MIX1 into a 100-mL volumetric flask and dilute to volume with Solvent B.

(MIX11) 0.0025 mg/L of each compound (6.25 µg/kg sample equivalents) and 0.05 mg/L of each internal standard: Pipette 0.25 mL of the fortification standard solutions MIX2 and 2 mL of the internal standard solution MIX1 into a 100-mL volumetric flask and dilute to volume with Solvent B.

(MIX12) 0.001 mg/L of each compound (2.5 µg/kg sample equivalents) and 0.05 mg/L of each internal standard: Pipette 0.1 mL of the fortification standard solutions MIX2 and 2 mL of the internal standard solution MIX1 into a 100-mL volumetric flask and dilute to volume with Solvent B.

All standard solutions should be stored in the freezer.