Cover Sheet for

ENVIRONMENTAL CHEMISTRY METHOD

Pestcide Name: Prosulfuron

MRID #: 433877-23

Matrix: Water/Soil

Analysis: LC/MS/MS

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ANALYTICAL METHOD FOR THE DETERMINATION OF CGA-152005 AND ITS METABOLITES CGA-159902 AND CGA-300406 (DESMETHYL CGA-152005) IN WATER AND SOIL BY HIGH PERFORMANCE LIQUID CHROMATOGRAPHY WITH MASS SPECTROMETRIC DETECTION INCLUDING VALIDATION DATA

METHOD NO. AG-600

SPONSOR AND TESTING FACILITY:

CIBA-GEIGY CORPORATION AGRICULTURAL DIVISION RESIDUE CHEMISTRY DEPARTMENT 410 SWING ROAD P. O. BOX 18300 GREENSBORO, NC 27419

Project

Number: 168982 Protocol

Number:

75-92 and

Amendment 1

Study Initiation Date: May 21, 1992

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Completion Date: 1/8/93

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IX.

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I. INTRODUCTION/SUMMARY

A. Scope

This method is used for the determination of CGA-152005 [N-[[(4-methoxy-6-methyl-1,3,5-triazin-2-yl)amino]carbonyl]-2-(3,3,3-trifluoropropyl)-benzene sulfonamide,, CAS #94125-34-5] and its metabolites CGA-159902 [2-(3,3,3trifluoropropyl)-benzenesulfonamide, CAS #94125-42-5] and CGA-300406 (desmethyl CGA-152005) in water and soil. The compounds are separated by high performance liquid chromatography (HPLC) and detected by mass spectrometry (LC-MS). An ionspray atmospheric pressure ionization (API) interface is used to introduce the HPLC effluent into the mass spectrometer equipped with a single quadrupole mass analyzer. The analytes are detected by selected ion monitoring (SIM) of their unique negative ions. The structures and chemical names of the analytes are presented in Figure 1.

The limit of detection (smallest standard amount injected during the chromatographic run) is 0.125 ng for all analytes. The limit of determination (the lowest fortification specified by the method which gives adequate recovery according to EPA guidelines) is 0.05 ppb for all analytes in water and 0.5 ppb in soil.

B. Principle

A 100-ml water sample is acidified with phosphoric acid and then passed through a C18 solid phase extraction (SPE) column. CGA-152005, CGA-300406, and CGA-159902 are retained on the C18 column. The column is rinsed with appropriate solvent. analytes are eluted from the C18 column with acetonitrile. The organic solvent in the eluate is removed to dryness using a rotary evaporator. The residue is redissolved in 50% acetonitrile/water and analyzed by LC-MS. A flow diagram for the water method is presented in Figure 2.

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Soil samples (20 g) are extracted at room temperature with a mechanical shaker using 100 ml of 20% (v/v) methanol/phosphate buffer (0.05 M, pH adjusted to 8). A filtered 50-ml aliquot (ca. 10 g) of the sample extract is diluted with water, acidified, and then passed through a C18 SPE cartridge for isolation of the analytes. The remainder of the cleanup procedure is the same as for water samples. The sample is analyzed by LC-MS. A flow diagram for the soil method is presented in Figure 3.

II. MATERIALS AND METHODS

A. Apparatus

- 1.0 Balance, analytical (Sartorius R160P) or equivalent.
- 2.0 Bottle, amber Boston round, with Polyseal-lined cap (Fisher cat. #05-563-2E) or equivalent.
- 3.0 Bottle, polypropylene, wide mouth (Fisher cat. #05-565-19A) or equivalent with cap (Fisher cat. #05-563-2E) or equivalent, appropriate size for soil extractions.
- 4.0 Centrifuge, Sorval Superspeed RC5-B (DuPont Instruments cat. #55228-9) or equivalent, with 6-place GSA rotor head (DuPont, Sorval GSA cat. #08136) or equivalent.
- 5.0 Cylinder, graduated, 100-ml (Fisher cat. #08-562-5C) or equivalent.
- 6.0 Filter paper, Whatman 2V (Fisher cat. #09-832D) or equivalent.
- 7.0 Flask, round bottom, 250-ml (Fisher cat. #10-067E) or equivalent.
- 8.0 Funnel, filter (Fisher cat. #10-326-2C) or equivalent.

- 9.0 Mechanical shaker, orbital (Fisher cat. #15-456-6) or equivalent.
- 10.0 Mixer, Vortex-Genie 2 (Fisher cat.
 #12-812) or equivalent.
- 11.0 Pasteur pipet (Fisher cat. #13-641-536) or equivalent.
- 12.0 pH stick, Corning (Fisher cat. \$13-641-536) or equivalent pH stick or meter.
- 13.0 Rotary evaporator, Buchi (Fisher cat. #09-548-105F) or equivalent.
- 14.0 SPE cleanup cartridge, C18 bonded phase, 1 g size (Varian Sample Preparation Products cat. #1225-6001).
- 15.0 Tube, concentration, 50-ml, with 19/38 ground glass joint (Fisher cat-#05-538-40B) or equivalent, with 24/19 enlarging adapter (Fisher cat. #01-035D) or equivalent.
- 16.0 Vials, amber, 1.5-ml (Sun Brokers, Inc. cat. #200-002) or equivalent, with teflon-lined, crimp-top seals (Sun Brokers, Inc. cat. #200-152) or equivalent.

B. Reagents and Analytical Standards

All reagents and polypropylene glycols are stored at room temperature. Solid analytical standards are stored frozen.

- 1.0 Acetic acid, concentrated, Optima grade (Fisher cat. #A465-250) or equivalent.
- 2.0 Acetonitrile, HPLC grade (Fisher cat. #A998-4) or equivalent.
- 3.0 Ammonium acetate, ACS grade (Fisher cat. #A637-500) or equivalent.
- 4.0 Extraction solvent (soil): 20% (v/v) methanol/phosphate buffer.

- 5.0 Formic acid, ACS grade (Fisher cat. #All8P-500) or equivalent.
- 6.0 HPLC mobile phase A: 0.1% (v/v) acetic acid/acetonitrile. Add 1.0 ml of acetic acid to 999 ml of acetonitrile and thoroughly mix.
- 7.0 HPLC mobile phase B: 0.1/25/74.9% (v/v) acetic acid/acetonitrile/water. Combine 1.0 ml of acetic acid with 250 ml of acetonitrile and 749 ml of water. Thoroughly mix.
- 8.0 Methanol, HPLC grade (Fisher cat. \$A452-4) or equivalent.
- 9.0 Phosphate buffer, 0.05 M, pH=8 ± 0.05. Dissolve 6.7 grams of sodium phosphate dibasic heptahydrate in 500 ml of purified water. Adjust the pH to 8 ± 0.05 with phosphoric acid and sodium hydroxide.
- 10.0 Phosphoric acid, 85% (Conc.), ACS grade (Fisher cat. #A242-1) or equivalent.
- 11.0 Phosphoric acid, 0.1% (v/v). Add 1.0 ml of conc. phosphoric acid to 999 ml of purified water and thoroughly mix.
- 12.0 Polypropylene glycol, M.W. 425 (Aldrich cat. #20,230-4).
- 13.0 Polypropylene glycol, M.W. 1000 (Aldrich cat. #20,232-0).
- 14.0 Polypropylene glycol, M.W. 2000 (Aldrich cat. #20,233-9).
- 15.0 PPG tuning solution. Dissolve 0.007 g PPG 425, 0.017 g PPG 1000, 0.118 g PPG 2000, and 0.021 g of ammonium acetate in 50 ml of methanol, 50 ml water, 0.1 ml acetonitrile, and 0.1 ml formic acid. Mix well. Store at room temperature in an amber bottle.
- 16.0 Sample diluent, 50% (v/v) acetonitrile/water.

- 17.0 Sodium phosphate dibasic, heptahydrate, ACS grade (Fisher cat. \$3373-500) or equivalent.
- 18.0 Water, HPLC grade, purified in-house with a HYDRO^{IM} purification system or equivalent.
- 19.0 CGA-152005, CGA-159902, and CGA-300406 analytical standards, CIBA-GEIGY Corp., P. O. Box 18300, Greensboro, NC 27419.

C. Analytical Procedure

1.0 Water

- 1.1 Measure a 100-ml aliquot of the water sample into a 100-ml graduated cylinder. (Note: A smaller aliquot may be used, but this will increase the limit of determination in ppb for the analyte. A larger aliquot may also be used to increase the sensitivity of the analysis.) Sample fortification, if required, should be done at this time (refer to Section II.J.2.0).
- 1.2 Add 1.0 ml of phosphoric acid. Shake the contents to mix.
- 1.3 Precondition the C18 cleanup cartridge by passing 5 ml each of methanol, acetonitrile, and finally 0.1% phosphoric acid through the cartridge. Discard the rinsate. Either positive pressure or a vacuum system may be used to aid the flow. Do not allow the packing to become dry.
- 1.4 Connect an appropriate reservoir on top of the C18 cartridge.
- 1.5 Load the aqueous sample from Step 1.2 into the reservoir and adjust the flow rate to a fast drip.

- Add 10 ml of 0.1% phosphoric acid to the 100-ml graduated cylinder that previously contained the sample and shake to rinse. (Note: For soil samples, add the solution to the 250-ml round bottom flask.) Add this rinsate to the reservoir just as the reservoir becomes empty. Discard the eluate.
- 1.7 Rinse the cartridge with 5 ml of purified water. Discard the eluate.
- 1.8 Dry the cartridge under vacuum for 5 minutes.
- 1.9 Elute CGA-152005, CGA-300406, and CGA-159902 using 5 ml of acetonitrile. Collect the eluate in a 50-ml concentration tube.
- 1.10 Remove solvent using a rotary evaporator and water bath temperature of approximately 35°C until dry. Use methanol to azeotrope any remaining water.
- 1.11 Dissolve the residue with 50% acetonitrile/water. Mix the contents on a vortex mixer for approximately 15 seconds. Transfer the sample to an appropriate amber vial. Analyze the sample by LC-MS. (Note: Samples that are not immediately analyzed should be stored in a freezer at a temperature of <0°C.)

2.0 Soil

(Note: Soil moisture content for a representative sample for each different soil sample must be determined by appropriate methodology at the time of sample analysis.)

- 2.1 Weigh 20 ± 0.1 g of soil sample and place in an appropriate size centrifugable polypropylene bottle.
- 2.2 Sample fortification, if required for this particular sample, is done at this time (refer to Section II.J.2.0).
- 2.3 Add 100 ml of 20% (v/v) methanol/phosphate buffer. Place the cap on the bottle. Shake the contents vigorously for approximately 15 seconds. Place the bottle in an orbital shaker and shake the sample for approximately one hour at approximately 25°C.
- 2.4 Remove the sample from the shaker. Centrifuge the sample at approximately 10,000 RPM for 10 minutes, or at an alternate speed and time if the results are considered satisfactory.
- 2.5 Decant the extracting solvent through a Whatman 2V filter and collect 50 ml of the filtrate in a 100-ml graduated cylinder for cleanup and analysis.
- 2.6 Add 25 ml of purified water. Shake the contents to mix.
- 2.7 Transfer the contents to a 250-ml round bottom flask. Rinse the 100-ml graduated cylinder with 25 ml of water and add the rinsate to the 250-ml round bottom flask.
- 2.8 Remove methanol (~10 ml) from the sample with a rotary evaporator and water bath of approximately 35°C. Add 1.0 ml of conc. phosphoric acid. Mix thoroughly.

2.9 The remainder of the cleanup procedure is identical to the procedure for water. At this point refer to Step 1.3 above and follow Steps 1.3 through 1.11.

D. <u>Instrumentation</u>

1.0 <u>Description and Operating Conditions</u> - HPLC

See Table I for a description of the HPLC system and chromatographic conditions.

2.0 <u>Description and Operating Conditions</u> - <u>Mass Spectrometer</u>

> CGA-152005, CGA-300406, and CGA-159902 are monitored as negative ions. A single quadrupole mass analyzer is used to separate the masses. Optimum sensitivity is achieved by selected ion monitoring (SIM) of the intense deprotonated molecular ion. See Table II for a description of the mass spectrometer instrumentation and operating conditions.

3.0 <u>Description and Operating Conditions</u>
-Ionspray Interface

The optimized values for the ionspray interface may vary with time and may need to be periodically reoptimized by infusion of an analyte into the mass spectrometer. See Table II for a description of typical ionspray operating conditions used with the analytes in Analytical Method AG-600.

- 4.0 Calibration and Standardization
 - 4.1 Calibrate and tune the mass spectrometer on a daily basis prior to analyzing samples. Check the calibration and tune by infusing a standard solution of polypropylene glycol (PPG)

into the mass spectrometer using the ionspray interface while monitoring positive ions. A typical mass calibration tune with PPG is shown in Figure 4.

- 4.2 Detect analytes at their specific monitoring ions. Determine the specific ion to monitor for each analyte by infusing a solution of that analyte into the mass spectrometer while scanning with the quadrupole mass analyzer to find the optimum ion. Typical monitoring ions for the analytes are listed in Table III. Typical ionspray mass fragmentation spectra are presented in Figure 5.
- 4.3 Determine the retention time of the analytes by injecting a standard solution into the HPLC. During a series of analyses, the analyte retention time should vary no more than 2% from its mean value, on a daily basis.
- Calibrate the instrument by constructing a calibration curve from detector response (chromatographic peak height or area) and the amount of analyte injected, encompassing a range from 0.125 - 1.25 ng (25-µl The response curve injections). can be constructed manually or, preferably, by generation of a linear regression equation by use of a computer or appropriate calculator. Typical standard calculations are presented in Table IV. Typical standard LC-MS chromatograms are shown in Figure 6.

E. <u>Interferences</u>

1.0 There are no known interferences criginating from the sample cleanup procedure. However, interferences can originate from impure chemicals, solvents, contaminated glassware, and the HPLC water supply.

F. Confirmatory Techniques

1.0 Although no confirmation techniques are presented here, this method provides detection based on a highly specific MS technique combined with chromatographic retention time.

G. Time Required

- 1.0 The sample extraction and cleanup procedure can be completed for a set of eight samples in an eight-hour working day.
- 2.0 Each HPLC analysis requires 20 minutes.

H. Modifications and Potential Problems

- 1.0 Contaminants from chemicals, solvents, glassware, and the HPLC water supply can interfere with the analysis. It is recommended that a reagent blank be run with an analysis set to verify that no interferences are originating from the chemicals and reagents used in this procedure. LC-MS is a very sensitive technique. All glassware should be solvent rinsed before use to prevent inadvertent contamination of control or low level samples.
- 2.0 Slow degradation of the parent compound CGA-152005 is observed in acidic solutions. Degradation of other sulfonyl urea compounds has been observed in solutions with high ionic strength, even though the compound exhibited stability at that pH. Therefore, samples and standards should not be dissolved and stored in acidic or buffered solutions. Dissolved samples must be analyzed as soon as possible after dissolution as

trace acidity and salts from the cleanup extractions and from the sample matrix can create a sensitized solution, causing analyte degradation, even though the sample was dissolved in a non-sensitized solvent. Samples should be stored <0°C if they cannot be promptly analyzed after the sample extraction/cleanup procedure.

- 3.0 CGA-152005 is very unstable when stored in methanol. It exhibits good stability in acetonitrile and water.
- 4.0 CGA-152005 exhibits minor photodecomposition. Therefore, all solutions must be stored in amber vials and bottles.
- 5.0 CGA-152005, CGA-159902, and CGA-300406 are weakly acidic. Their retention times in reversed phase HPLC will be affected by changes in the mobile phase pH.
- 6.0 Analytical Method AG-600 was validated only for sandy loam and loamy sand soils from Georgia and Iowa. Other soil types or soil samples from different locations may exhibit binding or interference problems which were not observed with these samples.
- 7.0 Studies have indicated that CGA-300406 exhibits poor stability.
- 8.0 "Bumping" is sometimes observed for soil samples during the solvent removal steps via rotary evaporation. Periodic venting of the vacuum and the use of solvent traps helps minimize inadvertent losses during these steps.

I. Preparation of Standard Solutions

All standards are stored in amber bottles in a freezer (<0°C) when not in use. No analyte stability or solubility problems

have been observed in the standard solutions used in this study. additional comments on CGA-300406.) Sulfonyl urea compounds exhibit varying degrees of stability in aqueous; therefore it is recommended that the analytical standards in acetonitrile/water be made on a weekly basis. CGA-300406 rapidly degrades at high concentrations in acetonitrile. Therefore, the 5 ng/µl mixed solution (Step 2.0) should be prepared immediately after preparation of the 100 ng/µl stock solution. The 100 ng/µl stock solution of CGA-300406 should not be reused. A new 5 ng/µl mixed solution in acetonitrile along with subsequently prepared fortification and analytical solutions should be prepared each week.

- 1.0 Prepare a 100 ng/µl stock solution for each analyte by dissolving 10.0 mg of the compound in 100 ml of acetonitrile.
- 2.0 Prepare a 5 ng/µl mixed solution in acetonitrile by pipetting 10.0 ml of each analyte (from its 100 ng/µl stock solution in Step 1.0) into a 100-ml volumetric flask and then diluting to mark with acetonitrile. This solution is used to prepare all subsequent dilutions.
- Prepare fortification swandards by 3.0 diluting the mixed 5 ng/µl solution (Step 2.0) with 50% acetonitrile/ water. Fortification standards should be prepared such that no more than 1 ml of the fortification solution is added to a sample. (Example: For a 100-ml water sample, the addition of 1.0 ml of a 0.50 ng/µl fortification solution will result in a fortification level of 5.0 ppb.) These solutions also serve as the analytical standards for analysis of CGA-300406, CGA-159902, and CGA-152005.

J. Methods of Calculation

1.0 Determination of Residues in Samples

Inject the sample solution from Step II.C.1.11 into the HPLC. The sample solution may be diluted if the analyte response exceeds the range of the calibration curve. The amount of analyte injected (ng) is determined by entering the value of the chromatographic peak height, or area, in the calibration response curve (Step II.D.4.4) and calculating (by computer, calculator, or manual means) the corresponding value Typical of nanograms injected. chromatograms for control and fortified control water samples are presented in Figure 7. Typical chromatograms for control and fortified control soil samples are presented in Figure 8.

2.0 <u>Determination of Residues in</u> <u>Fortified Samples</u>

Validate the method for each set of samples analyzed by including a control sample and one or more control samples fortified prior to the extraction procedure with 0.05 ppb or more of each analyte in water and with 0.5 ppb or more of each analyte in soil.

2.1 Add an appropriate volume of a fortification solution (from Step II.I.3.0) to the sample prior to any of the cleanup steps. The total volume of the added fortification solution should not exceed 1.0 ml. For soil samples, fortify the sample and allow sufficient time for the fortification solvent to evaporate (approximately 30 min.) before proceeding.

2.2 Proceed with the sample cleanup procedure (Step II.C.1.2 for water and Step II.C.2.3 for soil).

3.0 Calculations

Calculations may be performed by computer program or manually as follows:

3.1 Calculate the analyte concentration (in ppb) for field samples from equation (1):

(1) ppb analyte
$$=$$
 $\frac{\text{n= analyte found}}{\text{g sample injected}} \times \frac{1}{R} \times \frac{1}{R}$

where R is the recovery factor expressed in decimal form (i.e., 0.8 = 80%) and is calculated from equation (4), and P is the chemical purity of the analytical standard expressed in decimal form.

The grams of sample injected for water and soil are calculated from equations (2) and (3).

(2) g sample injected (water) =
$$\frac{g \times V_i}{V_s}$$

where g is the grams of sample used (for water, 1.0 ml = 1.0 g), V_i is the volume (ml) of sample injected onto the HPLC column, and V_i is the final volume (ml) of the sample (from Step II.C.1.11).

(3) g sample injected =
$$\frac{g}{v_a + (m \times g)} \times \frac{v_a \cdot v_1}{v_f}$$

where,

g is the grams of soil (wet weight) used, V_a is the aliquot volume of extracted sample used for analysis (ml), V_a is the volume of extract solvent used (ml), V_i is the volume (ml) injected onto the HPLC column, m is the percent moisture in the sample, expressed in decimal form (ex. 0.1 = 10%), V_f is the final volume (ml) of the cleaned-up sample (from Step II.C.1.11).

The recovery factor, expressed as a percentage (R%), is calculated from fortification experiments and is presented in equation (4).

ppb analyte found - ppb analyte (control) x 1000

The ppb of analyte found is calculated from equation (5).

(5) ppb analyte found =
g sample injected

Residues of metabolites found in test samples may also be expressed as parent equivalents by multiplying the amount found by the ratio of the molecular weight of CGA-152005 to that of the metabolite (equation (6)).

(6) ppb CGA-152005 equiv. = ppb metabolite X HW (m)

where MW(p) is the average molecular weight of CGA-152005 (419.4) and MW(m) is the average molecular weight of the metabolite, 405.4 for CGA-300406 and 253.2 for CGA-159902.

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3.2 The accuracy of the method is determined by the average recovery of the analytes fortified into the test substrate. The precision is estimated by the standard deviation of the determined concentration.

III. RESULTS AND DISCUSSION

This method has been validated under Protocol 75-92 and Amendment 1¹ for the analysis of control and fortified control water and soil. The objective of Protocol 75-92 was to validate "Draft" Analytical Method AG-600 for the determination of CGA-152005, CGA-159902 and CGA-300406 in water and soil at limits of determination of 0.05 ppb and 0.5 ppb, respectively.

Fortification levels ranged from 0.05 - 50 ppb in water and from 0.5 - 50 ppb in soil. limits of determination for CGA-152005, CGA-159902, and CGA-300406 were 0.125 ng, respectively. The recovery data for fortification experiments are presented in Tables V and VI for water and soil, respectively. The average recoveries and standard deviations for CGA-152005, CGA-159902, and CGA-300406 in water were 92 \pm 3.8, 100 \pm 2.3, and 84 \pm 1.56, respectively. No residues >0.025 ppb were observed for any analyte in the water control. and method blank samples. The average recoveries and standard deviations for CGA-152005, CGA-159902, and CGA-300406 in soil were 79 \pm 9.6, 84 \pm 10.5, and 79 \pm 8.2, respectively. residues >0.25 ppb were observed for any analyte in any soil control and method blank samples.

Reference substance ID, test system ID, protocol amendments, protocol deviations, and circumstances affecting the quality and integrity of data are reported in Residue Test Report RI-MV-010-92². All raw data associated with this study and the original final report and protocol are archived in the Metabolism and Residue Chemistry Archives at CIBA-GEIGY Corporation, Greensboro, NC. All non-study specific data (i.e., instrument logbooks, etc.)

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will be stored in the previously mentioned archives when all entry pages are filled or when the logbook is replaced.

IV. CONCLUSION

Analytical Method AG-600 is a valid and accurate method for determining the amounts of CGA-152005, CGA-159902 and CGA-300406 in water and soil.

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V. CERTIFICATION

This report and experimental results included in this study, laboratory Project I.D. AG-600, are certified to be authentic accounts of the experiments.

1/8/93

Date

John D. Vargo, Ph.D. Project Scientist Method Development Residue Chemistry 919-632-7525

Agricultural Division CIBA-GEIGY Corporation Post Office Box 18300 Greensboro, NC 27419

VI. CERTIFICATION OF GOOD LABORATORY PRACTICES

To the best of my knowledge, the analytical work reported in AG-600 was performed in accordance with Good Laboratory Practice Standards, 40 CFR Part 160.

1/8/93

Date

John D. Vargo, Ph.D

Study Director

//r/93 Date

Robert K. Williams

Manager

Method Development Residue Chemistry 919-632-2295

Agricultural Division CIBA-GEIGY Corporation Post Office Box 18300 Greensboro, NC 27419 AG-60**0** Page 23 of 41

CIBA-GEIGY PLANT PROTECTION DIVISION **OUALITY ASSURANCE UNIT**

QUALITY ASSURANCE STATEMENT

Study Title: VALIDATION OF "DRAFT" ANALYTICAL METHOD AG-600 FOR THE DETERMINATION OF CGA-152005 AND

ITS METABOLITES CGA-159902 AND DESMETHYL

CGA-152005 IN WATER AND SOIL BY HIGH

PERFORMANCE LIQUID CHROMATOGRAPHY WITH MASS

SPECTROMETRIC DETECTION

ANALYTICAL METHOD NO. AG-600

Study Director: J. D. Vargo

Project Number: 168982

Protocol No.: 75-92 & Amendments

Report No.: AG-600

Pursuant to Good Laboratory Practice Regulations, this statement verifies that the aforementioned study was inspected and/or audited and the findings reported to Management and to the Study Director by the Quality Assurance Unit on the dates listed below.

INSPECTION/AUDIT TYPE	INSPECTION/AUDIT DATES	REPORTING DATE
Protocol Audit	5/18/92	5/20/92
In-Progress Inspection	9/21/92	9/21/92
AG-600 Report Audit	12/14/92 - 12/15/92	12/18/92

Prepared By:

Hackworth

12/18/92

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VIII. LIST OF TABLES AND FIGURES

TABLE I. HPLC OPERATING CONDITIONS

Instrumentation:

Perkin-Elmer Model 410 Gradient Pump Perkin-Elmer Model ISS 200 Autosampler FIAtron Model CH-30 Column Heater

Operating Conditions:

Column Heater: 30°C Injection Volume: 25 µl

Mobile Phase Flow Rate: 1.0 ml/min Column: Zorbax Rx C8, 15 cm x 4.6 mm,

equipped with an Upchurch pre-column filter

(0.5 jum)

Mobile Phase A: 0.1% (v/v) acetic acid/

acetonitrile

Mobile Phase B: 0.1/25/74.9% (v/v) acetic

acid/acetonitrile/water

Mobile Phase Gradient Program:

Time (min.)	3 A	₹B		
0	0	100		
10	100	0		
4	100	0		
1	0	100		
. 5	0	100		

Total Run Time: 20 min.

Analyte Retention Times:

CGA-300406	. 7.8	min
CGA-159902	8.8	min
CGA-152005	10.1	min

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TABLE II. MASS SPECTROMETRY OPERATING CONDITIONS

Instrumentation:

PE Sciex API 1 Single Quadrupole Mass Spectrometer Ionspray Liquid Introduction Interface

Instrument Control and Data Collection:

MacIntosh IIfx Computer

Software:

Calibration and Mass Tuning: Tune 2.1.2 Acquisition: RAD 2.15 β

Quantitation: MacQuan 1.1.2 Display: MacSpec 3.2

All software programs written and provided by PE Sciex.

Operating Condition:

Interface Heater: 70°C
Mobile Phase Split Ratio: ~20-25:1
Curtain Gas Flow: 0.9 on gauge
Nebulizer Gas Flow: 0.9 on gauge

Typical State File Values:

Parameter	PPG Positive Ion Mass Calibration	Negative Io Analytes		
DĪ	Linked	100		
ISV	5500	-4200		
·IN	650	- 650		
OR	35	~ 55		
RO	35	- 30		
MI	10	10		
RE1	109	109		
DM1.	08	- 0.04		
R	32	- 28		
Ľ9	- 150	100		
FP	- 150 ·	100		
MU	-3600	3800		

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TABLE III. TYPICAL ANALYTE MONITORING IONS

Analyte	Exact Molecular Weight		Mass For Negative Ion Monitoring
CGA-152005	419.09		418.0
CGA-300406	405.07	•	404.0
CGA-159902	253.04		252.0

Scan Rate: 1 scan/sec

Dwell Time: 249.98 msec

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TABLE IV. TYPICAL STANDARDIZATION DATA

	FEAK AREA						
Amount Injected (ng)	CGA-300406	CGA-159902	CGA-152005				
0.125	23208	50119	20982				
0.25	40278	99568	45127				
0.5	83314	199043	88559				
1.25	197083	491848	214861				
Slope (area/ng)	155298	392443	171255				
Y-Intercept (area)	3469	1659	1403				
Correlation	0.9997	1.0000	0.9998				

Peak area obtained with PE Sciex MacQuan 1.1.2 software. Linear regression data obtained with CIBA-GEIGY Multichrom 2.0 computer system. Software by VG Laboratory Systems Ltd. AG-600 Page 28 of 41

TABLE V. RECOVERY DATA FOR FORTIFIED WATER

Sample Code	Fortification Amount (ppb)	<u>CGA</u> ppb <u>Found</u>	-300406 % Recovery	ppb	159902 % Recovery	ppb	152005 % Recovery
152VAL1 152VAL2 152VAL3 152VAL4 152VAL5 152VAL6 152VAL7	0 (Blank) 0 (Control) 0.05 0.05 0.5 5	<0.025 <0.025 0.042 0.043 0.42 4.1 42	NA NA 83 86 83 82 85	<0.025 <0.025 0.048 0.050 0.51 5.1	NA NA 96 101 101 102 100	<0.025 <0.625 0.044 0.047 0.48 4.3 46	NA NA 89 94 96 87 92
Average Recov Std. Deviation:	ery:		84 1.56		100 2.3	•	92 3.8

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TABLE VI. RECOVERY DATA FOR FORTIFIED SOIL

		1	CGA-300406		CGA-159902		CGA-152005	
•		Fortification	ppb	Æ	ppb	%	ppb	9E
Sample Code	Description	Amount (ppb)	Found	Recovery		Recovery		Recovery
	<u>-</u>							
152VAL8	Blank	0 (Blank)	<0.25	NA	<0.25 ⋅	NA NA	<0.25	NA
152VAL9	GA. 0-24"	0 (Control)	<0.25	NA .	< 0.25	NA	<0.25	NA
152VAL10	GA. 0-24"	ر کا	0.44	88	0.42	83	0.40	75
152VAL11	GA. 0-24"	0.50	0.38	76	0.35	69	0.33	61
152VAL12	GA 0-24"	0.98	0.85	87	0.78	80	0.81	81
152VAL13	GA. 0-24"	9.7	8.1	83	9.2	95	9.2	94
152VAL14	GA. 0-24"	51	41	80	52	101	47	91
152VAL15B	GA, 24-48"	0 (Control)	<0.25	NA	<0.25	NA	<0.25	NA
152VAL16B	GA, 24-48"	0.51	0.40	78	0.43	84	0.40	79
152VAL17B	GA, 24-48	0.49	0.40	82	0.43	87	0.39	79
152VAL18B	GA, 24-48"	1.0	0.82	82	0.84	84	0.73	73
152VAL19B	GA. 24-48"	10	6.6	65	9.2	91	8.6	85
152VAL20B	GA. 24-48"	48	38	80	46	95	44	92
152VAL21E	IA, 0-24"	0 (Control)	<0.25	NA	<0.25	NA	<0.25	NA
152VAL22E	IA. 0-24"	0.30	0.40	68	0.40	69	0.36	72
152VAL23E	IA. 0-24"	0.50	0.42	73	0.38	63	0.36	71
152VAL24E	IA, 0-24"	1.0	0.78	72	0.74	68	0.71	71
152VAL25E	IA. 0-24"	10	6.7	66	9.8	97	7.7	77
152VAL26E	IA, 0-24"	50	34	. 69	46	92	38	76
								, ,
152VAL27	IA, 24-48"	0 (Control)	<0.25	NA	<0.25	NA	<0.25	NA
152VAL28	IA, 24-48"	0.30	0.42	84	. 0.41	82	0.38	- 66
152VAL29	IA. 24-48"	0.50	0.40	80	0.44	89	0.43	76
152VAL30	IA. 24-48"	1.0	0.91	91	0.80	80	0.80	75
152VAL31	LA. 24-48"	10 .	8.9	89	8.9	89	9.3	93
152VAL32	IA, 24-48"	50	45	91	46	91	47	94
	, == •, = • • •		_			,		
•	•			,	., .			•
Average Recov	erv:			79	•	` 84		79
Treate mount	··· .			• •		-	1.	.,
est married				8.2		10.5		9.6
Std. Deviation:			. ,	۰.۲	. ,	10.3		7.0

Note: Calculated finite amounts of residues less than 0.25 ppb are reported in the table as "<0.25 ppb." The Multichrom Worksheet subtracts the finite residues found in control samples from the ppb found value of the associated fortified sample to obtain the corrected percent recovery value.

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FIGURE 1. CHEMICAL NAMES AND STRUCTURES

CGA-152005 N-[[(4-methoxy-6-methyl-1,3,5-triazin-2-yl)amino]carbonyl]-2-(3,3,3-trifluoropropyl)-benzene-sulfonamide CAS 494125-34-5 Chemical Purity: 97.1

CGA-300406 Chemical Purity: 97.8

CGA-159902 2-(3,3,3-tr1fluoropropyl)-benzenesulfonamide CAS #94125-42-5 Chemical Purity: 99.6 AG-600 Page 31 of 41

FIGURE 2. AG-600 FLOW DIAGRAM FOR WATER

Aliquot 100 ml of water. Fortify, if necessary.

Acidify with concentrated phosphoric acid.

Isolate analytes on a pre-conditioned C18 SPE cartridge.

Elute analytes with acetomitrile.

Analyze by LC-MS.

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FIGURE 3. AG-600 FLOW DIAGRAM FOR SOIL

Weigh 20 g sample of soil. Fortify, if necessary

Add 100 ml of 20% methanol/0.05 M phosphate buffer, pH = 8. Extract for one hour at room temperature with mechanical shaking.

Centrifuge and filter extract.

Aliquot 50 ml of the extract. Add 50 ml of water. Acidify with concentrated phosphoric acid.

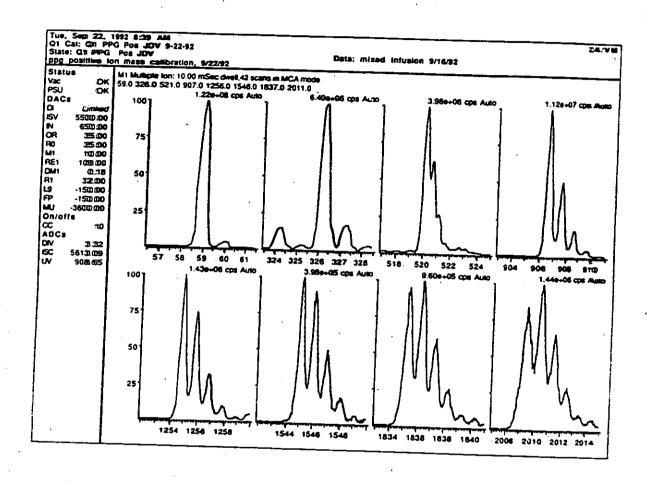
Isolate analytes on a pre-conditioned C18 SPE cartridge.

Elute analytes with acetonitrile.

Analyze by LC-MS.

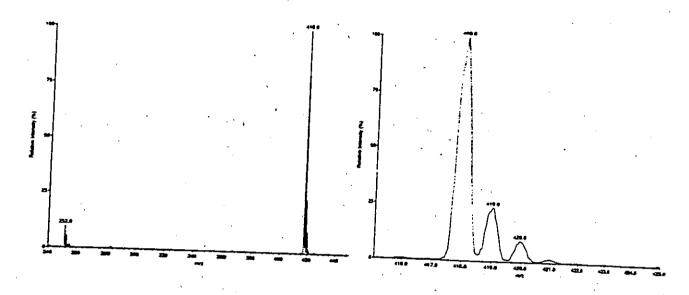
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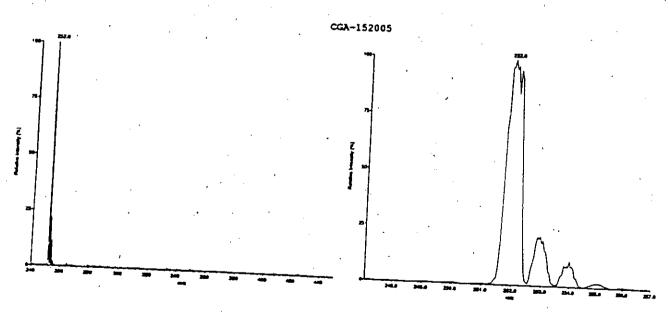
FIGURE 4. TYPICAL PPG MASS CALIBRATION TUNE



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FIGURE 5. TYPICAL NEGATIVE ION MASS FRAGMENTATION SPECTRA

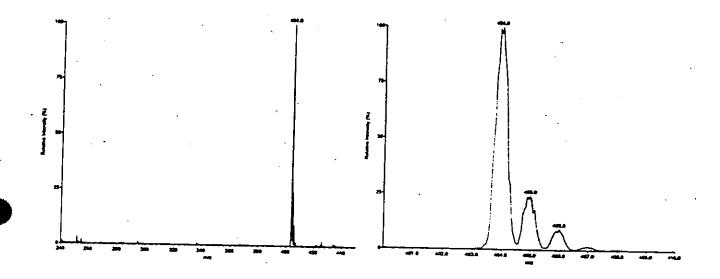




CGA-159902

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FIGURE 5. TYPICAL NEGATIVE ION MASS FRAGMENTATION SPECTRA (Continued)

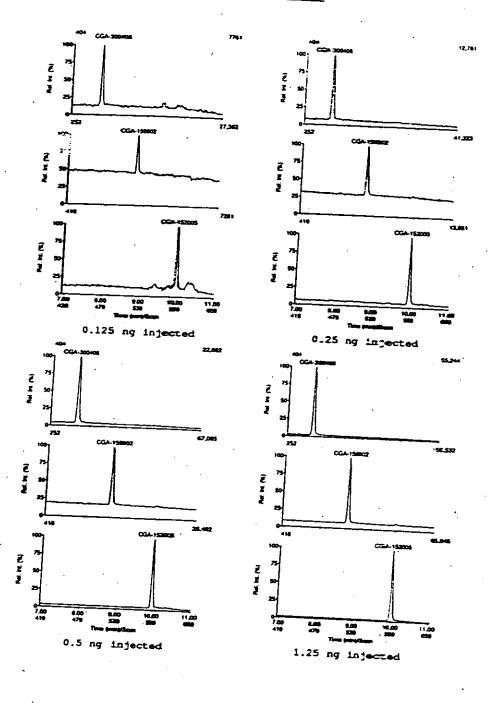


CGA-300406

All infusion spectra obtained in 0.1/39.9/60% (v/v) acetic acid/water/acetonitrile.

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FIGURE 6. TYPICAL LC-MS CHROMATOGRAMS OF ANALYTICAL STANDARDS



BEST AVAILABLE COPY

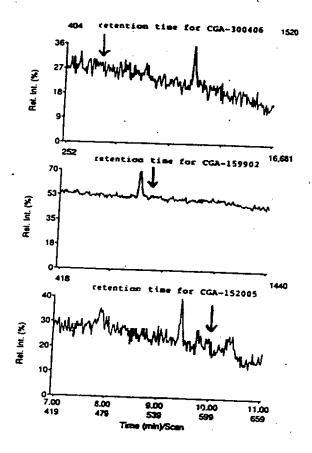
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PAGE

112

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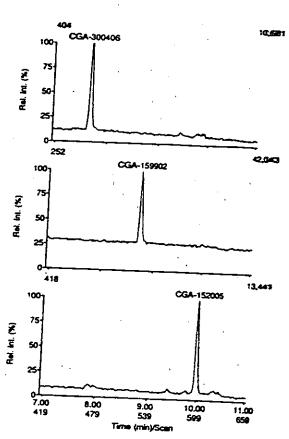
FIGURE 7. TYPICAL LC-MS CHROMATOGRAMS FOR CONTROL AND FORTIFIED WATER



Sample: 152VAL2, Control 5.0 g Injected

Amount Found:

CGA-300406: <0.125 ng, <0.025 ppb CGA-159902: <0.125 ng, <0.025 ppb CGA-152005: <0.125 ng, <0.025 ppb



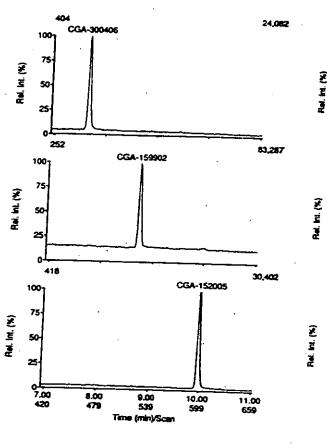
Sample: 152VAL4, Control + 0.05 ppb 5.0 g Injected

Amount Found:

CGA-159902: 0.22 ng, 0.043 ppb, 86% CGA-159902: 0.25 ng, 0.050 ppb, 101% CGA-152005: 0.24 ng, 0.047 ppb, 94%

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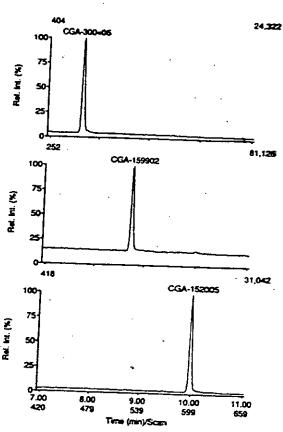
FIGURE 7. TYPICAL LC-MS CHROMATOGRAMS FOR CONTROL AND FORTIFIED WATER (Continued)



Sample: 152VAL5, Control + 0.5 ppb 1.25 g Injected

Amount Found:

CGA-300406: 0.52 ng, 0.42 ppb, 83% CGA-159902: 0.63 ng, 0.51 ppb, 101% CGA-152005: 0.60 ng, 0.48 ppb, 96%



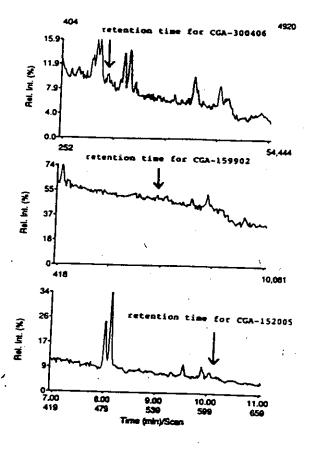
Sample: 152VAL7, Control + 50 ppb 0.013 g Injected

Amount Found:

CGA-30G406: 0.53 ng, 42 ppb, 85% CGA-159902: 0.63 ng, 50 ppb, 100% CGA-152005: 0.58 ng, 46 ppb, 92%

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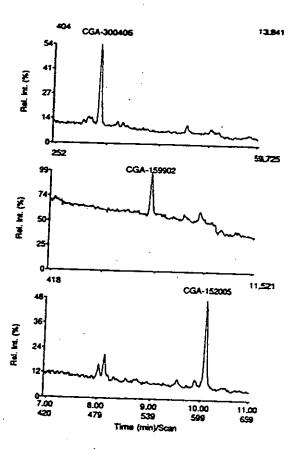
FIGUPE 8. TYPICAL LC-MS CHROMATOGRAMS FOR CONTROL AND FORTIFIED SOIL



Sample: 152VaL9, Control 0.51 g Injected

Amount Found:

CGA-300406: <0.125 ng, <0.25 ppb CGA-159902: <0.125 ng, <0.25 ppb CGA-152005: <0.125 ng, <0.25 ppb



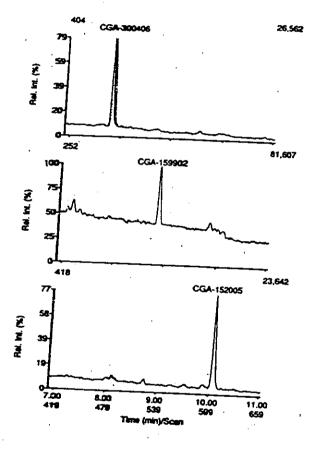
Sample: 152VAL10, Control + 0.51 ppb 0.49 g Injected

Amount Found:

CGA-300406: 0.22 ng, 0.44 ppb, 88% CGA-159902: 0.21 ng, 0.42 ppb, 83% CGA-152005: 0.20 ng, 0.40 ppb, 75%

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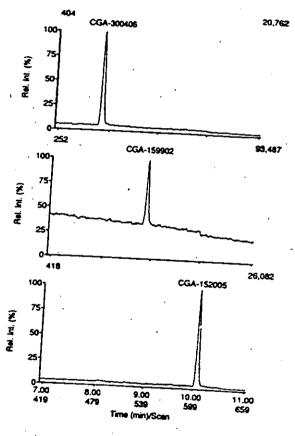
FIGURE 8. TYPICAL LC-MS CHROMATOGRAMS FOR CONTROL AND FORTIFIED SOIL (Continued)



Sample: 152VAL12, Control + 0.98 ppb 0.51 g Injected

Amount Found:

CGA-152005: 0.43 ng, 0.85 ppb, 87% CGA-159902: 0.40 ng, 0.78 ppb, 80% CGA-152005: 0.41 ng, 0.81 ppb, 81%



Sample: 152VAL14, Control + 51 ppb 0.010 g Injected

Amount Found:

CGA-300406: 0.40 ng, 41 pph, 80% CGA-159902: 0.51 ng, 52 pph, 101% CGA-152005: 0.45 ng, 47 pph, 91%

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IX. REFERENCES

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