

Cover Sheet for

ENVIRONMENTAL CHEMISTRY METHOD

Pesticide Name: Terbacil

MRID #: 435855-01

Matrix: Soil

Analysis: GC/ECD

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TRADE SECRET

STUDY TITLE

Long-Term Field Soil Dissipation of Terbacil Herbicide
- Analytical Report -

DATA REQUIREMENT

EPA Pesticide Assessment Guidelines, Subdivision N
Series 164-5: Long Term Soil Dissipation

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STUDY COMPLETION DATE

February 7, 1995

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DuPont Study No. AMR1975-91
PTRL Report No. 314W-1
PTRL Project No. 314W

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Report No. 314W-1
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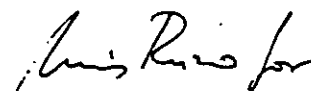
STATEMENT OF GOOD LABORATORY PRACTICE COMPLIANCE

GLP Compliance Statement for Final Report No. 314W "Long-Term Field Soil Dissipation of Terbacil Herbicide", Report no. 314W-1 for E.I. du Pont de Nemours and Company, DuPont Agricultural Products, Wilmington, Delaware.

PTRL's project director confirms that the above study was conducted in compliance with the applicable EPA Good Laboratory Practice Standards (40 CFR 160), except for the following:

- 1) There are a few late entries in the notebook.
- 2) Traces of some GC runs were obscured to preserve confidential information.


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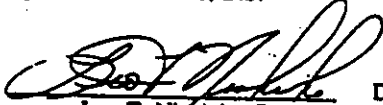
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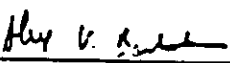
CERTIFICATION OF AUTHENTICITY

TITLE: Long-Term Field Soil Dissipation of Terbacil Herbicide
PROJECT NO.: 314-W
DATE ANALYTICAL WORK INITIATED: November 25, 1991
DATE ANALYTICAL WORK COMPLETED: May 26, 1994
DATE REPORT ISSUED: February 7, 1995

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Storage Location of Records,
Specimens and Final Report:

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PTRL WEST, INC. QUALITY ASSURANCE UNIT STATEMENT

Compound: Terbacil

Title: Long-Term Field Soil Dissipation of Terbacil Herbicide: Analytical Report

GLP QUALITY ASSURANCE INSPECTION:

Date of GLP Inspections:	Date Reported to the Study Director and to Management:	Phases of the Study Which Received a GLP Inspection by the Quality Assurance Unit:
1/10/92	1/10/92	Partitioning/Extraction PHI 30 IL Soil
4/1-2/92	4/6/92	Raw Data
4/20/92	4/21/92	GC Analysis PHI 120 IL/160 DE Soil
6/17/92	6/17/92	Clean-Up PHI 180 IL Soil
10/22/92	11/4/92	GC Analysis PHI 540 CA Soil
3/17-18/93	3/23/93	Raw Data
4/28/93-5/10/93	6/3/93	Raw Data
6/7/93	6/10/93	Extraction PHI 726 CA Soil
12/15/93	1/24/94	GC Analysis PHI 900 DE Soil
4/6-7/94	4/12/94	GC Analysis PHI 1080 CA Soil
5/19-23/94	5/26/94	Raw Data
7/29/94-8/1/93	8/4/94	Raw Data
3/19/94-9/8/94	11/10/94	Draft Final Analytical Report

QUALITY ASSURANCE STATEMENT:

The Quality Assurance Unit has reviewed the analytical study report No. 314W-1 and has determined that the report accurately reflects the raw data generated during the conduct of the analytical portion of the study.

ARCHIVING:

The original project specific raw data files and final report are temporarily located in the archives of PTRL West, Inc., Richmond, California, 94806. The above materials and specimens will be transferred to *E.I. du Pont de Nemours and Company*, DuPont Agricultural Products, Experimental Station Wilmington, Delaware 19880-0402 and/or the DuPont Records Management Center, Wilmington, Delaware 19880-0870 upon the Sponsor's authorization. Facility records, such as Compound Control logs, equipment use, standardization/calibration, and maintenance logs, etc. and a copy of the analytical report will be maintained in the archives of PTRL West, Inc.

Michele Blair
Michele Blair
PTRL West, Inc., Quality Assurance

February 2, 1995
Date

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STATEMENT OF ANALYTICAL STANDARDS

The following standards were used as analytical references:

COMPOUND: Terbacil
(3-tert-butyl-5-chloro-6-methyluracil)
SOURCE: E. I. du Pont de Nemours and Company
REF. NO.: IN-D732-56
PURITY: 99.9%

COMPOUND: Metabolite A
(3-tert-butyl-5-chloro-6-hydroxymethyluracil)
SOURCE: E. I. du Pont de Nemours and Company
REF. NO.: IN-G2449-4
PURITY: 97.3%

COMPOUND: Metabolite B
(6-chloro-2,3-dihydro-7-hydroxymethyl-3,3-dimethyl-5H-oxazolo[3,2-a]-pyrimidin-5-one)
SOURCE: E. I. du Pont de Nemours and Company
REF. NO.: IN-W2207-3
PURITY: 98.0%

COMPOUND: Metabolite C
(6-chloro-2,3-dihydro-7-methyl-3,3-dimethyl-5H-oxazolo[3,2-a]-pyrimidin-5-one)
SOURCE: E. I. du Pont de Nemours and Company
REF. NO.: IN-T2170-5
PURITY: 99.6%

All reference standards, calibration standard solutions, and fortification standard solutions were stored frozen (< 0° C). The above parent and metabolites standards were utilized throughout the study. Traces of gas chromatography chromatograms showing similar retention times and chromatographies for all analytes throughout the study indicated the stability of all reference standards.


Ann Y. Kuo
Project Director

Date

INTRODUCTION

To support the re-registration of terbacil (Sinbar® Herbicide), a registered product of DuPont Agricultural Products, three field residue experiments were conducted to determine the terbacil residue dissipation patterns in/on field soil. Experiments were conducted in the States of California and Delaware over a three year period and in Illinois over a two year period. Soil samples were collected on 1-day prior to application, day of application, designated Day 15, 30, 60, 90, 120, 160 (not collected from CA site), 180, 360, 540, 720, 900, and 1080 after application from both California and Delaware field sites. Soil sampling intervals from Illinois field site was conducted in the same manner as the California and Delaware field sites up to 720 days after application. Since significant levels of terbacil Metabolites A, B and C were not detected from the aerobic and anaerobic soil metabolism study, terbacil metabolites were only monitored at the day of application through Day 60 and at the last sampling interval (Day 1080) for the California site, at the day of application through Day 60 for the Delaware site and at the day of application through Day 30 for the Illinois site. Terbacil was monitored throughout the entire study.

SOIL FIELD TRIAL PROGRAM

Terbacil soil dissipation residue studies were conducted in Madera, California, Newark, Delaware and Geneseo, Illinois. Untreated control and terbacil aged soil samples were collected from the above three locations. All samples were received frozen and unmacerated at PTRL West and remained frozen until processed or sub-sampled for analysis.

STANDARD REFERENCE MATERIALS

Reference standards of terbacil and Metabolites A, B and C were provided by the E. I. du Pont de Nemours and Company, DuPont Agricultural Products, Wilmington, Delaware. Standards were stored at < 0° C until used for stock solution preparation. Stock solutions of each analyte were prepared at 1000 µg/mL in ethyl acetate. Separate dilutions of 100 µg/mL of each analyte were prepared. Then 5 µg/mL combined analyte solutions in ethyl acetate of terbacil (IN-D732), Metabolite A (IN-G2449), Metabolite B (IN-W2207) and Metabolite C (IN-T2170) were prepared for matrix spiking and standard derivatization for the analysis of terbacil and its

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three metabolites. For terbacil soil analysis, terbacil analyte solutions in ethyl acetate at 50 $\mu\text{g/mL}$, 10 $\mu\text{g/mL}$ and 1 $\mu\text{g/mL}$ were prepared for matrix spiking. These were stored at $< 0^\circ\text{C}$ in amber bottles with Teflon lined caps until used for sample fortification. Standard solutions were allowed to return to room temperature prior to use. The primary reference standard, stock and diluted standard solutions were all stored frozen.

MATRICES

Soil samples were collected inside open-ended plastic cores capped with red and black plastic caps and then packed inside canvas bags. Soil samples were shipped frozen via Federal Express or A.C.D.S. to PTRL West. The soil samples were logged in and frozen on the same day upon arrival at PTRL West. All samples were stored frozen ($< 0^\circ\text{C}$).

ANALYTICAL METHOD SUMMARY

Residue analyses were conducted using the published procedures specified in the Pesticide Analytical Manual Vol. II, 180.209 with additional modifications presented in this analytical method summary report. Brief descriptions of the extraction, derivatization, clean up and chromatographic quantitation of terbacil and its metabolites are summarized below:

Reagents:

Solvents

Chloroform, Optima grade, Fisher Scientific, Santa Clara, CA

Ethyl Acetate, Optima grade, Fisher Scientific, Santa Clara, CA

Acetonitrile, Optima grade, Fisher Scientific, Santa Clara, CA

n-Hexane, Optima grade, Fisher Scientific, Santa Clara, CA

Toluene, Optima grade, Fisher Scientific, Santa Clara, CA

Methanol, Optima grade, Fisher Scientific, Santa Clara, CA

Bis (trimethylsilyl) trifluoroacetamide (BSTFA) plus 1% trimethylchlorosilane (TMCS), Pierce, Rockford, IL

Anhydrous Sodium Sulfate, Certified ACS grade, Fisher Scientific, Santa Clara, CA

NaOH, Certified ACS grade, Fisher Scientific, Santa Clara, CA

Glassware and Miscellaneous Equipment:

Disposable culture tubes, 16 x 125 mm, CMS catalogue no. 339-317
Florisil Sep-Paks®, Waters Associates, Milford, MA, part no. 51960
Waring blender and blending cups, stainless steel or glass
1000-mL Round bottom flasks
Cotton batting
250-mL Separatory funnels
60° Long stem filtering funnels
250-mL Round bottom flasks
500-mL Round bottom flasks
10-mL Derivatizing vials, Pierce, Rockford, IL
Volumetric flasks of various sizes
Volumetric pipettes, 1- and 4-mL
Analytichem Bond Elut reservoirs and adaptors, Analytichem International, Harbor City, CA
Vacuum filtration adapter, Aldrich, Milwaukee, WI, catalogue no. Z11, 562-2
Rubber adapters
Vacuum evaporator, Buchi Model RE111, Brinkmann Instruments, Inc., Burlingame, CA,
with temperature controlled waterbath
Nitrogen evaporator, Meyer Model 111, Organomation Associates, Inc., South Berlin, MA,
with temperature controlled waterbath
Glasswool
Pasteur pipettes, 5" and 9"
Food Processor, Regal La Machine I, Model V813, Regalware, Kewaskum, WI
Amber bottles with Teflon lined lids
Drying oven, Fisher Isotemp®, 500 Series, Fisher Scientific, Santa Clara, CA
Deionized Water Generator, Barnstead Four Module NanoPure II

Sample Processing:

Soil cores up to the depth of 0-90 cm were collected. Treated soil cores were sectioned into 0-15 cm, 15-30 cm, 30-45 cm and 45-60 cm soil depths and control soil cores were segmented into 0-15 cm, 15-30 cm and 30-60 cm soil depths. The same depth segments within each of the replicate subplots were combined into a homogeneous mixture by vigorous agitation inside a plastic bag. The remaining 60-90 cm treated soil samples and 60-90 cm control soil samples were left uncomposited. These lower soil depth soil samples were preprocessed, when needed. All samples were stored frozen in labeled plastic bags until sub-sampled for analysis.

Isolation:

1. Weigh 10 g of a representative sample into the blender cup and add 150 mL of chloroform. Blend the sample for 5 minutes.
2. Pass the chloroform extract through a cotton plugged funnel into a 1000-mL round bottom flask.
3. Extract twice more with 100-mL portions of chloroform and filter through the cotton as well. Rinse the blender cup and cap with chloroform until all particulate has been removed from the cup.
4. Add 10 mL of water to the combined extracts and evaporate the chloroform in a vacuum rotary evaporator at -35°C .

Partitioning:

5. Transfer the residue (-5 mL of water) using several volumes of acetonitrile to a 250-mL separatory funnel. (Final volume should be less than 100 mL.)
6. Add 50 mL n-hexane and shake one minute. Allow phases to separate and centrifuge if necessary to ensure separation. Discard hexane and repeat partitioning with two additional portions of hexane.
7. Quantitatively transfer acetonitrile to a 250-mL round bottom flask and evaporate to dryness at -35°C .
8. Dissolve all residues using several rinses of 0.1% NaOH and transfer to a 250-mL separatory funnel (final volume should be less than 80 mL).
9. Add 75 mL of ethyl acetate and shake for two minutes. Allow phases to separate. Filter the ethyl acetate through a 1 1/2 inch bed of anhydrous sodium sulfate into a 500-mL round bottom flask. Repeat the partition with ethyl acetate three more times and combine the ethyl acetate extracts.
10. Concentrate the combined extracts to 5 mL at -35°C by roto-evaporation. (For soil analysis monitoring terbacil compound only, steps 11 through 17 and step 24 were eliminated because no derivatization was required for the terbacil compound.)

Derivatization:

11. Quantitatively transfer the concentrate from the 500-mL round bottom flask to a 10-mL derivatizing vial with additional rinses of ethyl acetate.
12. Concentrate to 1 mL under nitrogen at -35°C .
13. Add 300 μL of BSTFA + 1% TMCS to the derivatizing vial, cap with a Teflon lined lid and shake vigorously for approximately 20 seconds. Allow the derivatization to take place overnight or at least 16 hours at room temperature.

14. Simultaneously with sample derivatization, prepare a 20 µg/mL standard of all analytes by pipetting 4 mL of the 5 µg/mL standard solutions used for fortification into a 10-mL derivatization vial. Concentrate to 1 mL under nitrogen and derivatize with samples.

Final Sample Clean-up:

15. Prepare a sodium sulfate mini-column for each sample and standard by plugging the bottom of a 5-inch Pasteur pipette with a small amount of glass wool and filling with anhydrous sodium sulfate until the pipette is approximately 1/3 full. Place the mini-columns into 16 x 125 mm disposable culture tubes.
16. After at least 16 hours derivatization, add 1 mL of deionized water and shake 20 seconds. Allow phases to separate. Do not add water to the standards until after completion of the final sample clean-up on the samples to ensure standard stability.
17. Remove the upper phase from the derivatizing vial by Pasteur pipette and place the extract on the top of the sodium sulfate mini-column. Rinse the column twice with 2 mL of ethyl acetate and collect in the culture tube.
18. Concentrate the solution to dryness under nitrogen at -35°C and resuspend in 10 mL of ethyl acetate/hexane (20:80). Dissolve all the particulate into the solution using a Pasteur pipette.
19. Attach an Analytichem reservoir to a Waters 900-mg Florisil Sep-Pak, on to a vacuum adaptor joined to a 250-mL round bottom flask and attach to a light vacuum.
20. Pre-rinse the Florisil Sep-Pak with 5 mL of 20:80 ethyl acetate/hexane, then add the 10 mL of sample to the reservoir. Allow the sample to be pulled through the Florisil cartridge at approximately 10 mL/min.
21. Rinse the sample tube with 5 mL of 20:80 ethyl acetate/hexane and transfer it to the reservoir. Rinse the culture tube with 5 mL of 10:5:85 methanol/ethyl acetate/toluene and pull it through the cartridge as well, collecting all fractions into the 250-mL round bottom flask.
22. Concentrate the extract to dryness by roto-evaporation at -35°C and resuspend in 3 mL of ethyl acetate.
23. Transfer the 3 mL of extract to a clean 10-mL centrifuge tube and rinse the flask with several small portions of ethyl acetate. Transfer the rinses to the centrifuge tube as well.
24. Add water to the standard and dry using the mini sodium sulfate column as described in #17 above. A 10-mL centrifuge tube is used for collecting standards instead of a culture tube.
25. Concentrate both samples and standards to 1 mL under nitrogen at -35°C. (For analysis monitoring terbacil compound only, samples were diluted to 10 mL with the exceptions of the 1.0 ppm and 5.0 ppm fortified soil samples which were diluted to 40 mL. To determine the background for the 40 mL diluted fortified samples, the associated control was also diluted to 40 fold.)

26. Transfer all samples and standards to GC vials and prepare the following standard dilutions by means of a 1000- μ L syringe.

Terbacil and Metabolites A, B and C Analysis:

10 μ g/mL standard= 500 μ L of 20 μ g/mL standard + 500 μ L of ethyl acetate

5 μ g/mL standard= 500 μ L of 10 μ g/mL standard + 500 μ L of ethyl acetate

2 μ g/mL standard= 400 μ L of 5 μ g/mL standard + 600 μ L of ethyl acetate

1 μ g/mL standard= 500 μ L of 2 μ g/mL standard + 500 μ L of ethyl acetate

0.5 μ g/mL standard= 500 μ L of 1 μ g/mL standard + 500 μ L of ethyl acetate

Terbacil Analysis Only:

1 mL aliquot of the 1 μ g/mL terbacil reference standard was pipetted into a GC vial and the following serial dilutions were performed. Normally, two sets of the calibration standards were prepared to cover a sample set.

0.5 μ g/mL standard= 500 μ L of 1 μ g/mL standard + 500 μ L of ethyl acetate

0.25 μ g/mL standard= 500 μ L of 0.5 μ g/mL standard + 500 μ L of ethyl acetate

0.125 μ g/mL standard= 500 μ L of 0.25 μ g/mL standard + 500 μ L of ethyl acetate

27. Generally, inject samples on the GC in the following order for sets monitoring terbacil and its three metabolites: 10 ppm standard, sample, sample, 5 ppm standard, sample, sample, 2 ppm standard, etc. Derivatized samples appear to be stable for at least 18 hours at room temperature.

Generally, inject samples on the GC in the following order for terbacil analysis only: 0.125 ppm standard, sample, sample, 0.25 ppm standard, sample, sample, 0.5 ppm standard, sample, sample, 1 ppm standard, sample, sample, and then repeat the sequence again.

Chromatography:

Instrumentation:	5890	Hewlett Packard Gas Chromatograph equipped with Electron Capture Detector, Hewlett Packard Company, Wilmington, DE
	3396A	Hewlett Packard Integrator, Hewlett Packard Company, Wilmington, DE
	7673A	Hewlett Packard Autosampler, Hewlett Packard Company, Wilmington, DE
Column:	HP-5 fused silica column (5% phenyl methyl silicone), 10 m x 0.53 mm id, 2.65 μ m film thickness, Hewlett-Packard, Wilmington, DE	
Injector Temperature:	200°C	

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Detector Temperature: 300°C

Oven Temperature:

Initial Temperature	155°C for 16 min.
Initial Ramp	155°C to 200°C at 10°C/min.
Ramp A	200°C to 280°C at 50°C/min.
Final Hold	280°C for 1 min.

Gases: Carrier Gas= Helium at =12.5 mL/min.
Makeup Gas= 5% Argon/Methane at =29.5 mL/min.

Injection Volume: 1 µL

Representative chromatograms and terbacil calibration curve are shown in Appendix B.

% Moisture Determination for Soil Samples:

Approximately 10 g of each soil sample was placed into an aluminum weighing dish and dried in the oven overnight at approximately 100°C for % moisture determination. The weight of the soil sample before and after oven drying was recorded.

Recovery:

Method recovery was performed with control samples fortified with terbacil at 0.1 ppm (µg/g), 1.0 ppm and 5.0 ppm for each sample set. The diluted standards were fortified on the matrices prior to the initial extraction step. Average and individual recoveries are shown in Table L

Metabolites A, B and C were monitored during the beginning of this soil dissipation study. Concurrent recoveries were performed with untreated soil samples fortified with Metabolites A, B and C at 0.1 ppm (µg/g), 1.0 ppm and 5.0 ppm. Recovery results are presented in Appendix A: Section 4.

Quantitation:

A calibration curve was generated for each analyte with each sample set from the co-injected standards. The equation of the line based on the peak area of the standard versus concentration injected in nanograms was generated by least squares linear regression calculated by the computer program, Cricketgraph™, version 1.2, MacWarehouse, South Norwalk, CT. The correlation coefficient (r^2) calculated for each set of standards could not be less than 0.95 for

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the data to be considered acceptable. The integrated peak area was plotted versus concentration and fitted by linear regression to the formula $y=mx + b$ where y is peak area and x is concentration injected in nanograms. The dry gross residue concentration of the soil sample was then determined from the $y=mx + b$ formula in nanograms, divided by the volume injected in μL , multiplied by the final volume in mL, multiplied by the % moisture factor and divided by sample weight (10 g).

The equation for the calculation of dry gross residues in soil samples was derived from the % moisture factor. Both the % moisture factor and dry gross residue determination equations are shown below:

$$\% \text{ Moisture of Soil} = \frac{(\text{Gross Wet Soil Weight}-\text{Container Weight})-(\text{Gross Dry Soil Weight}-\text{Container Weight})}{(\text{Gross Wet Soil Weight}-\text{Container Weight})} \times 100\%$$

$$\% \text{ Moisture Factor} = \frac{100}{100 - \% \text{ Moisture of Soil}}$$

The % moisture of soil did not apply to the fortified samples, because the ppm ($\mu\text{g/g}$) of analyte fortified was based on the weight of moist soil and the purpose of the fortified samples were to determine the method concurrent recoveries on moist soil samples.

Linear regression formula from calibration curve $y=mx + b$

$$\text{Concentration of dry gross residues in soil (ppm)} = \frac{\text{Integrated Peak Area of analyte (y)} - \text{Calibration Intercept (b)}}{\text{Calibration Slope (m)}} \times \frac{\text{Final Volume (mL)}}{\text{Sample Weight (10 g)}} \times \frac{\% \text{ Moisture Factor}}{\text{Volume Injected (1 } \mu\text{L)}}$$

The peak areas for fortified samples (y) were corrected for background by subtracting the peak response of the control sample.

% Recovery is determined by:

$$\% \text{ Recovery} = \frac{\text{Concentration in Fortified Matrix } (\mu\text{g/g})}{\text{Concentration Fortified } (\mu\text{g/g})} \times 100\%$$

Spreadsheets showing the concentrations of terbacil analyte in soil and the associated concurrent recoveries are displayed in Appendix A: Sections 1 through 3.

RESULTS AND DISCUSSION

Soil core samples from the Sinbar[®] Herbicide field soil dissipation residue have been analyzed for terbacil residues. Metabolites A, B and C were analyzed in selected soil samples, because the three metabolites were not detected at significant levels in the soil metabolism study.

Average recoveries of terbacil on untreated California soil fortified at 0.1, 1.0 and 5.0 ppm were 110 %, 96 % and 89 %, respectively. Untreated Delaware soil gave average recoveries of 109 %, 97 % and 87 % for terbacil fortified at 0.1, 1.0, and 5.0 ppm, respectively. Similar average recoveries were also achieved for untreated Illinois soil fortified with 0.1, 1.0 and 5.0 ppm terbacil. The overall recovery results were 108 %, 96 % and 90 %, respectively at the 0.1, 1.0 and 5.0 ppm fortification levels. Average and individual recoveries for terbacil in soil are shown in Table I.

The terbacil residue levels for Sinbar[®] Herbicide treated soil samples at the California location decreased from 0.94 ppm at day 0 to 0.09 ppm by 1080 days. Similarly, terbacil residue levels found for the Sinbar[®] treated Delaware soil samples declined from 1.39 ppm at day 0 to 0.07 ppm of 1080 days after application. The terbacil dissipation study was conducted in Illinois over a two year period. Terbacil residue levels for Sinbar[®] treated soil samples from the Illinois field site dissipated from 1.58 ppm at the day 0 to 0.05 ppm by 720 days. Individual residue data for California soil, Delaware soil and Illinois soil are presented in Tables II, III and IV, respectively.

Terbacil's three metabolites were investigated on day 0, (day of application) and at a designated PHI of 15, 30, 60 and 1080 days for the California soil samples, on planting day, designated PHI of 15, 30 and 60 days for the Delaware soil samples and on application day, and at a designated

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PHI of 15 and 30 days for Illinois soil samples. Metabolites A and B were reacted with BSTFA + 1% TMCS for 16 hours to form volatile derivatives for chromatography. Recovery results for Metabolites A, B and C in soil were acceptable. Overall, residues found for Metabolites B and C in all soil types were less than the limit of detection of 0.05 ppm. Metabolite A residues were observed in California and Delaware soil only. Average residue levels for Metabolite A in California soil increased from 0.05 ppm at application day to 0.14 ppm at PHI 15 days and then decreased to less than the detection limit at PHI 1080 days. Similarly, maximum average Metabolite A residues were observed in the Delaware soil at PHI 15 days (0.14 ppm) and then dissipated to less than the limit of detection at PHI 60 days. Individual recovery and residue data for Metabolites A, B and C are presented in Appendix A: section 4. Representative chromatograms and calibration curves are included in Appendix B.