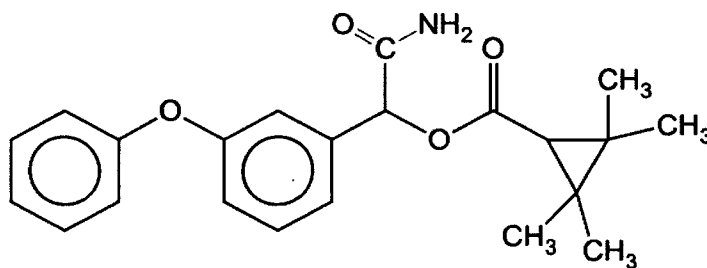
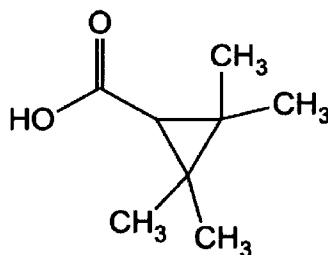


**Analytical Reference Standards**

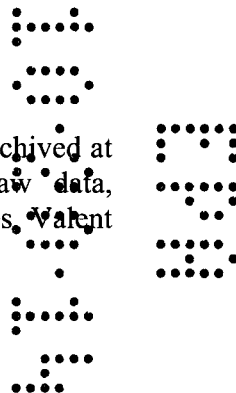
Standard Name: CONH<sub>2</sub>-Fenpropathrin  
 Lot Number: AS 2129b  
 CPS ID: 14-CPS-Aug28-01  
 Valent Sample Archive No.: V-Arc-2312  
 Source: Valent USA  
 Purity: 99.6%  
 Molecular Formula: C<sub>22</sub>H<sub>25</sub>NO<sub>4</sub>  
 Average Mass: 367.2  
 Molecular Structure:



Standard Name: TMPA  
 Lot Number: AS 2357a  
 CPS ID: 14-CPS-Aug28-03  
 Valent Sample Archive No.: V-Arc-2317  
 Source: Valent USA  
 Purity: 99.7%  
 Molecular Formula: C<sub>8</sub>H<sub>14</sub>O<sub>2</sub>  
 Average Mass: 142.1  
 Molecular Structure:

**Other**

Upon completion of the study, a copy of the protocol and the final report will be archived at Critical Path Services, LLC (CPS). The original protocol, final report, raw data, correspondence, and other documentation will be transferred to the Valent Archives, Valent U.S.A. Corporation, 6560 Trinity Court, Dublin, California, 94568.



## 2.0 INTRODUCTION

The objective of this study was to validate Valent method (Golden Pacific Laboratories method GPL-MTH-084) “Analytical Method for the Determination of Fenpropathrin Metabolites CONH<sub>2</sub>-Fenpropathrin and TMPA in Soil by LC-MS/MS” [1]. This method passed the ILV for both analytes in soil on the first attempt with no major modifications.

This study was designed to fulfill the requirements of the US EPA Ecological Effects Test Guidelines OCSPP 850.6100: Environmental Chemistry Methods and Associated Independent Laboratory Validation [2]. In addition, this study was conducted in compliance with US EPA FIFRA (40 CFR Part 160) GLP standards [3].

## 3.0 MATERIALS AND METHODS

### 3.1 Test Substances

**Standard name:** CONH<sub>2</sub>-Fenpropathrin  
**Lot number:** AS 2129b  
**CPS ID:** 14-CPS-Aug28-01  
**Sample archive no.:** V-Arc-2312  
**Manufacturer’s ID:** Sumitomo Lot 08SC8046993  
**Purity:** 99.6%  
**Date of analysis:** April 07, 2014  
**Expiration date:** April 07, 2015  
**Storage conditions:** Frozen

**Standard name:** TMPA  
**Lot number:** AS 2357a  
**CPS ID:** 14-CPS-Aug28-03  
**Sample archive no.:** V-Arc-2317  
**Manufacturer’s ID:** Sumitomo Lot F-16  
**Purity:** 99.7%  
**Date of analysis:** April 22, 2014  
**Expiration date:** April 22, 2016  
**Storage conditions:** Frozen

### 3.2 Test System

The test system used for the validation was a control soil sample collected at Tift County, Georgia. The sample was stored in a freezer when not in use.

### 3.3 Equipment and Reagents

The equipment and reagents used for the method validation were as outlined in pages 5 through 7 of the Valent method (Golden Pacific Laboratories method GPL-MTH-084;

included in the protocol in Appendix 1) and precisely documented in the study records (Notebook 201, pages 4–8). Identical or equivalent equipment and materials were used.

### 3.3.1 Equipment and Apparatuses

Analytical balance (Mettler Toledo)  
Manual pipettor of multiple volumes  
Electronic pipettor of multiple volumes  
Refrigerator/freezer (Nor-Lake<sup>®</sup> Scientific)  
Top-loading balance (Mettler Toledo)  
Wrist-Action<sup>®</sup> Shaker Model 75 (Burrell Scientific, LLC)  
Allegra<sup>®</sup> X-22R centrifuge (Beckman Coulter, Inc.)  
Ultrasonic cleaner 5210 (Branson)  
Polypropylene conical tubes (15-mL and 50-mL)  
Nalgene<sup>™</sup> bottle (125-mL)  
50-mL plastic syringes  
N-EVAP with water bath and thermometer (Organomation Associates, Inc.)  
Vortexer (VWR International)  
Walk-in -20°C freezer (Imperial Brown, Inc.)  
PTFE syringe filter, 0.45 µm, 25 cm (Pall Life Sciences)  
Oasis<sup>®</sup> Max cartridges, 60 mg/3cc (Waters)  
SPE manifold (Supelco)  
Various sizes of volumetric flasks  
Various sizes of glass graduated cylinders  
LC-MS/MS - Agilent 1200 binary pump HPLC system and autosampler, coupled to an Applied Biosystems<sup>®</sup> API 4000<sup>™</sup> mass spectrometer with an electrospray ionization interface

### 3.3.2 Reagents

Acetone (Pharmco-AAPER)  
Acetonitrile (EMD)  
Methanol (EMD)  
Formic acid (Sigma-Aldrich<sup>®</sup>)  
Ammonium hydroxide (Sigma-Aldrich<sup>®</sup>)  
Monobasic sodium phosphate, monohydrate (Sigma-Aldrich<sup>®</sup>)  
Dibasic sodium phosphate, anhydrate (EMD)  
Milli-Q water

## 3.4 Experimental Design

### 3.4.1 Establishment of the Method

Prior to performing the ILV, the analyte retention times, instrument detection limits, and linearity of instrument responses to a range of analyte concentrations were determined. The SPE cartridge recovery was also checked before the validation.

### 3.4.2 Standard Solutions Preparation

The primary stock solution for each reference standard was prepared by weighing approximately 50.0 mg of compound onto a tared piece of weigh paper and transferring to a 50-mL glass volumetric flask. Acetone was added up to volume, and the solution was sonicated appropriately.

Fortification solution 015-FS1A, containing 10.0 µg/mL of CONH<sub>2</sub>-fenpropathrin, was prepared by transferring 100 µL of the primary stock solution into a 10-mL volumetric flask and diluting to volume with methanol. A second fortification solution, 015-FS2A, was prepared at a concentration of 1.00 µg/mL by measuring 1.00 mL of 015-FS1A into a 10-mL volumetric flask and diluting to volume with methanol.

TMPA fortification standard 015-FS3A (10.0 µg/mL) was prepared by pipetting 99.8 µL of TMPA stock standard (1.00 mg/mL) into a 10-mL volumetric flask and bringing to volume with methanol. The TMPA fortification standard 015-FS4A (1.00 µg/mL) was prepared by measuring 1.00 of 015-FS3A into a 10-mL volumetric flask and bringing to volume with methanol.

The CONH<sub>2</sub>-fenpropathrin calibration standard solutions were prepared from an intermediate standard, IM2A (100 µg/L), which was prepared from FS2A in methanol/water (50:50, v/v) solution. The calibration standards were prepared by adding an appropriate amount of IM2A standard to 10-mL volumetric flasks and diluting to total volume with methanol/water (50:50; v/v) solution. Calibration standards for CONH<sub>2</sub>-fenpropathrin ranged from 0.250 µg/L to 10.0 µg/L.

TMPA calibration standard solutions were prepared from an intermediate standard 015-IM1A containing TMPA at 1.00 µg/mL, which was prepared from 015-FS3A in methanol/water/formic acid (50:50:1, v/v/v) solution. The TMPA calibration standards were prepared by adding an appropriate amount of IM1A standard to 10-mL volumetric flasks and diluting to total volume with methanol/water/formic acid (50:50:1, v/v/v) solution. Calibration standards for TMPA ranged from 2.50 µg/L to 100 µg/L.

All standard solutions were refrigerated (~4°C) when not in use.

### 3.4.3 Sample Validation Sets, Fortification, and Extraction Procedure

#### Sample Validation Sets

The analytical set consisted of 13 samples: one reagent blank, two untreated controls, five untreated controls fortified at the LOQ (0.0100 ppm), and five untreated controls fortified at 10× LOQ (0.100 ppm).

### Sample Preparation for Analysis

1. Weighed  $10 \pm 0.1$  grams of soil sample into a 125-mL Nalgene™ bottle.
2. Added the appropriate amount of fortification solution to the sample.
  - a. For the reagent blank and control samples, added nothing.
  - b. For the LOQ samples, added 100  $\mu\text{L}$  of 015-FS1A (1.00  $\mu\text{g}/\text{mL}$  of  $\text{CONH}_2$ -fenpropathrin) and 100  $\mu\text{L}$  of 015-FS3A (1.00  $\mu\text{g}/\text{mL}$  TMPA) fortification solution.
  - c. For the 10 $\times$  LOQ samples, 100  $\mu\text{L}$  of 015-FS2A (10.0  $\mu\text{g}/\text{mL}$  of  $\text{CONH}_2$ -fenpropathrin) and 100  $\mu\text{L}$  of 015-FS4A (10.0  $\mu\text{g}/\text{mL}$  TMPA) fortification solution.
3. Added 50-mL extraction solvent methanol:water (90:10, v/v) to sample.
4. Placed tubes horizontally on a Wrist-Action® shaker, and shook for ~45 minutes.
5. Centrifuged samples at 3000 rpm for 3 minutes.
6. Filtered approximately 30 mL of sample extract through a 0.45  $\mu\text{m}$  filter using a syringe.
7. Pipetted 20.0 mL sample into a 50-mL centrifuge tube.
8. Evaporated sample until approximately 2 mL remained using an N-EVAP set at 40°C.
9. Added 10.0 mL of 100 mM phosphate buffer to each sample and mix.
10. Placed Waters Oasis® Max cartridge (60mg/3cc) onto an SPE manifold. Conditioned the cartridge with 3.00 mL methanol followed by 3.00 mL water.
11. Loaded the sample to the cartridge.
12. Washed the cartridge with 3.00 mL water, 3.00 mL of 0.15 mM ammonium hydroxide solution and 3.00 mL of methanol/water (50:50, v/v) solution. Discarded all washes.
13. Placed a 15-mL centrifuge tube under the cartridge, eluted  $\text{CONH}_2$ -fenpropathrin with 5.00 mL of methanol and collected into a 15-mL tube (eluent 1).
14. Placed a clean 15-mL centrifuge tube under the cartridge, eluted TMPA with 2.00 mL of 2% formic acid in methanol solution, and collected into a 15-mL tube (eluent 2). Vacuum was applied to pull the liquid completely.
15. Brought the total volume of eluent 1 to 10.0 mL with water and diluted five times for the  $\text{CONH}_2$ -fenpropathrin analysis. Brought eluent 2 to 4.00 mL with water and diluted two times for TMPA analysis.

#### 3.4.4 Sample Processing and Analysis

The samples were analyzed as described by the Valent method (Golden Pacific Laboratories method GPL-MTH-084). The samples were analyzed with six calibration standards interspersed with the samples in a sequence. The continuing calibration standards (2.00  $\mu\text{g}/\text{L}$  for  $\text{CONH}_2$ -fenpropathrin and 20.0  $\mu\text{g}/\text{L}$  for TMPA) were injected at the beginning, middle, and end of the sequence. The coefficient of determination of the continuing calibration standards was acceptable (<15%) for the  $\text{CONH}_2$ -fenpropathrin and TMPA analyses.

### 3.5 LC-MS/MS Instrumentation

#### Instrumentation

Agilent 1200 HPLC System (Agilent Technologies)

API 4000™ Tandem Mass Spectrometer, MS/MS (Applied Biosystems®)

HPLC Column: Phenomenex Luna® C18, 30 × 2 mm, 3 μm

Software: Applied Biosystems® Analyst® 1.6.2

Refer to Table 2 for the details of the instrument conditions.

### 3.6 Data Acquisition and Reporting

Peak integration was performed by Analyst® software version 1.6.2. The MS detector responses (peak area) for various injected standard concentrations were used to generate an external calibration curve for the analytes of interest. The overall purpose for the external calibration curve was to display acceptable linearity ( $r^2 \geq 0.99$ ) of the assigned calibration range. The recoveries of the analytes from the fortified samples were calculated by multi-point calibration.

Recovery results of each analyte were computed for each sample. The equations used for quantification are presented in Appendix 2. A statistical treatment of the data includes the calculation of means, standard deviations (SD), RSDs as percentages (%), and the 95% confidence intervals. All statistics were calculated using Microsoft® Office Excel 2003.

### 4.1 Method Establishment

Prior to performing the ILV, the analyte retention times, instrument detection limits, and linearity of instrument responses to a range of analyte concentrations were determined, and the test system was verified as free of interferences at appropriate retention times. The SPE cartridges were checked for TMPA recoveries. Due to the instrument sensitivity for TMPA, CPS requested and obtained Sponsor approval to double the sample size for the cartridge load (20 mL instead of 10 mL) for the TMPA analysis.

**Table 2 LC-MS/MS System Operating Parameters**

HPLC System: Agilent Model 1200  
 Software: Applied Biosystems® Analyst® 1.6.2

HPLC Column: Phenomenex Luna® C18, 30 × 2 mm, 3 µm  
 Needle Wash: Water/methanol (50:50, v/v)  
 Wash Time: Flush port, 15 seconds

**CONH<sub>2</sub>-Fenpropathrin Analysis**

Mobile Phase: (A - Aqueous): 0.2% formic acid in HPLC-grade water  
 (B - Organic): 0.2% formic acid in acetonitrile

Injection Volume: 10.0 µL  
 Run Time: 6.5 minutes  
 Gradient:

Time (min)	A (%)	B (%)	Flow
0.0	60.0	40.0	500
2.0	30.0	70.0	500
3.5	30.0	70.0	500
3.6	10.0	90.0	500
4.6	10.0	90.0	500
4.7	60.0	40.0	500
6.5	60.0	40.0	500

**Mass Spectrometer Conditions:**

Parameter	Setting	
Ion Source:	TurboSpray	
Scan Type:	MRM	
Curtain Gas (CUR):	25	
Temperature (TEM):	100	
Ion Source Gas 1 (GS1):	60	
Ion Source Gas 2 (GS2):	50	
Interface Heater (ihe):	ON	
Polarity	Positive (0–3.75 min)	
Declustering Potential (DP):	66.00	
Entrance Potential (EP):	10.00	
Dwell Time (msec)	150	
Collision Gas (CAD):	6.00	
Ion Spray Voltage (IS):	5500	
Transitions Monitored:	Quant.	Conf.
	368.2→125.2	368.2→97.1
Collision Cell Exit Potential(CXP):	8.00	6.00
Collision Energy (CE):	15.0	43.0

**TMPA Analysis**

Mobile Phase: (A - Aqueous): Water  
 (B - Organic): Acetonitrile  
 Injection Volume: 50.0  $\mu$ L  
 Run Time: 6.5 minutes  
 Gradient:

Time (min)	A (%)	B (%)	Flow
0.0	90.0	10.0	500
3.0	40.0	60.0	500
4.0	40.0	60.0	500
4.1	10.0	90.0	500
4.5	10.0	90.0	500
4.6	90.0	10.0	500
6.5	90.0	10.0	500

**Mass Spectrometer Conditions:**

Parameter	Setting	
Ion Source:	TurboSpray	
Scan Type:	MRM	
Curtain Gas (CUR):	30	
Temperature (TEM):	500	
Ion Source Gas 1 (GS1):	70	
Ion Source Gas 2 (GS2):	50	
Interface Heater (ihe):	ON	
Polarity	Negative	
Declustering Potential (DP):	-65.00	
Entrance Potential (EP):	-5.00	
Dwell Time (msec)	150	
Collision Gas (CAD):	6.00	
Ion Spray Voltage (IS):	-4500	
Transitions Monitored:	Quant.	Conf.
	141.0→106.9	141.0→97.0
Collision Cell Exit Potential(CXP):	-5.00	-10
Collision Energy (CE):	-26	-18



## APPENDIX 2 CALCULATIONS

For calculation of the concentrations, calibration curves were used. These curves were calculated automatically after each sequence run with the Applied Biosystems<sup>®</sup> Analyst<sup>®</sup> software version 1.6.2 using a linear regression with 1/concentration weighting. Further calculations were performed using Microsoft<sup>®</sup> Office Excel 2003.

The linear equation is expressed as:

$$y = mx + b$$

where

y = peak area

x = concentration (ng/mL)

The concentration of analyte in the final sample solution can be calculated as follows:

$$\text{Final Sample Concentration } C \text{ (ng/mL)} = (y-b)/m$$

The residue of analytes in test samples is calculated as follows:

$$\text{Residue (ppb)} = \frac{C \times (1/1000) \times FV \times DF}{W \times AL}$$

where

C = Concentration in final sample (ng/mL)

FV = Final sample volume (10.0 mL for CONH<sub>2</sub>-fenpropathrin and 4.00 mL for TMPA)

W = Sample weight

DF = Additional dilution factor

AL = Aliquot factor 0.4

Recoveries are calculated using the following equation:

$$\text{Recovery (\%)} = \frac{\text{Analyzed Final Sample Concentration}}{\text{Theoretical Final Sample Concentration}} \times 100$$