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Determination of Residues of 1,2-Dichloropropane and *cis*- and *trans*-1,3-Dichloropropene in Soil by Purge and Trap Extraction, Capillary Gas Chromatography and Mass Selective Detection

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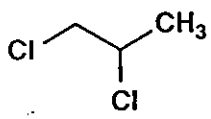
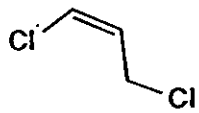
A. Scope

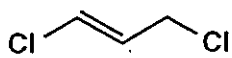
This method is applicable for the quantitative determination of residues of 1,2-dichloropropane (1,2-D) and *cis*- and *trans*-1,3-dichloropropene (1,3-D) in soil over the concentration range of 0.200-160,000 µg/kg with a validated limit of quantitation of 0.200 µg/kg for each compound.



1,2-D
CAS No. 78-87-5



cis-1,3-D
CAS No. 10061-01-5



trans-1,3-D
CAS No. 10061-02-6

B. Principle

This analytical method is based on established EPA purge-and-trap methodology for volatile organic analytes (VOAs), such as Method 8260 (1). Initial extraction of the analytes from soil is by one of two related methods depending on the levels at which the analytes are present in the sample. For analysis in the low-level range (0.20-200 µg/kg), a slurry of soil and water is heated and stirred. The volatile chlorinated hydrocarbons are purged from the sample by sparging with helium, and captured on a sorbent-containing trap. For analysis in the high-level range (200-160,000 µg/kg), the soil sample is first extracted with methanol. An aliquot of the methanol is diluted with water and the resulting water sample is then sparged with helium. The analytes purged from the sample are captured on a sorbent-containing trap. Subsequent steps for both analyses are identical. When purging is

complete, the trap is heated and backflushed with helium, and the VOAs are desorbed and transferred to a gas chromatograph (GC). The analytes are then separated on a capillary column and quantitated using mass selective detection (MSD).

C. Safety Precautions

1. Each analyst must be acquainted with the potential hazards of the reagents, products, and solvents used in this method before commencing laboratory work. SOURCES OF INFORMATION INCLUDE: MATERIAL SAFETY DATA SHEETS, LITERATURE, AND OTHER RELATED DATA. Safety information on non-DowElanco products should be obtained from the container label or from the supplier. Disposal of reagents, reactants, and solvents must be in compliance with local, state, and federal laws and regulations.
2. The analytes covered by this method can cause severe and possibly fatal respiratory distress at air concentrations in the ppm range. All operations involving the neat analytes, or concentrated solutions of these compounds, must be carried out in a fume hood. In addition, the effluent from the GC splitter should be routed into an exhaust vent or through a carbon trap to prevent the release of the analytes into the laboratory air.

D. Equipment (Note R.1.)

1. Balance, analytical, Model AE200, Mettler Instrument Corporation, Hightstown, NJ 08520.
2. Balance, pan, Model PM600, Mettler Instrument Corporation.
3. Gas chromatograph, Model 5890 Series II, Hewlett-Packard, Wilmington, DE 19808.
4. Mass selective detector, Model 5971, Hewlett-Packard, Palo Alto, CA 94304.
5. Mass selective detector data system, Model G1034B, Hewlett-Packard.
6. Purge and trap autosampler, Model 2016, Tekmar Company, Cincinnati, OH 45249.
7. Purge and trap autosampler/concentrator, Dynatrap, Dynatech Precision Sampling Corp., Baton Rouge, LA 70895.
8. Purge and trap concentrator, Model 3000, Tekmar Company.
9. Water purification system, Model Milli-Q UV Plus, Millipore Corporation, Milford, MA 01757.

E. Glassware and Materials (Note R.1.)

1. Column, capillary gas chromatography, DB-VRX, 30 m x 0.25 mm i.d., 1.4 µm film thickness, catalog number 122-1534, J&W Scientific, Folsom, CA 95630.
2. Column inlet liner, deactivated, catalog number 5181-8818, Hewlett-Packard, Kennett Square, PA 19348.
3. Filter, charcoal, catalog number 7972, Chrompack, Inc., Raritan, NJ 08869. (Note R.2.)
4. Filter, moisture, catalog number 7971, Chrompack, Inc. (Note R.2.)
5. Filter, oxygen, catalog number 7970, Chrompack, Inc. (Note R.2.)

6. Fritted sparge glassware, 25 mL, catalog number 14-3022000, Tekmar Company.
7. Gas, helium, 99.995% purity, Airco, Murray Hill, NJ 07974.
8. Gas-tight syringes, fixed needle, 10, 100 and 500 µL, catalog numbers 1701, 1710, and 1750, Hamilton Company, Reno, NV 89520.
9. Gas-tight syringe, Leur Lock, 25 mL, catalog number 1025, Hamilton Company.
10. Magnetic stir bars, 3 mm x 13 mm, catalog number 14-511-61, Fisher Scientific, Pittsburgh, PA 15219.
11. Syringe valve, catalog number 2-0940M, Supelco, Inc., Bellefonte, PA 16823.
12. Trap, Tenax, catalog number 2-1075, Supleco, Inc.
13. Trap, Tenax, catalog number 12-0083-003, Tekmar Company.
14. Vials, 11-dram, catalog number 033395D, with polyethylene-lined screw caps, catalog number ALP5026, Fisher Scientific.
15. Vials, 40 mL pre-cleaned amber for volatile analysis, with Teflon-lined septa and screw caps, catalog number 2V42MHSE2BS, Fisher Scientific, Pittsburgh, PA 15219. (Note R.3.)

F. Reagents and Chemicals (Note R.1.)

1. Reagents

a. Internal standard, 2-bromo-1-chloropropane, 95%, compound number 23,127-4, Aldrich Chemical Company, Milwaukee, WI 53233.

b. Methyl alcohol, purge and trap grade, catalog number 41,481-6, Sigma-Aldrich, St. Louis, MO 63178.

c. Standards

(1) 1,2-dichloropropane

The 1,2-D standard used for generating the validation data contained in this method was Lot Number AGR277102, with a purity of 99.2% (2).

(2) *cis*-1,3-dichloropropene

The *cis*-1,3-D standard used for generating the validation data contained in this method was Lot Number AGR164301, with a purity of 97.1% (3).

(3) *trans*-1,3-dichloropropene

The *trans*-1,3-D standard used for generating the validation data contained in this method was Lot Number TSN100232, with a purity of 97.2% (4).

Obtain standards from Test Substance Coordinator, DowElanco, Indianapolis, IN 46268.

d. Water, distilled/deionized, purified using a Milli-Q UV Plus purification system (Section D.9.).

G. Preparation of Standards

1. Preparation of VOA Standard Solutions

NOTE: CARRY THIS PROCEDURE OUT IN A FUME HOOD. When mixing standard solutions do not shake them excessively, as loss of analytes may occur. Mix

by capping and gently inverting the solutions approximately five times. Store all standard solutions under frozen conditions in vials with Teflon lined screw cap lids. Allow standard solutions to warm to room temperature prior to use.

a. Tare a 100-mL volumetric flask containing approximately 80 mL of methanol. Add 0.1100-0.1300 g of 1,2-D dropwise via Pasteur pipette (approximately eight to ten drops). Make sure that the liquid falls directly into the methanol and does not run down the inside walls of the volumetric flask. Stopper the flask. Reweigh the flask and calculate the exact weight of the analyte added (Note R.4.). Bring the solution to volume with methanol to yield a stock solution of approximately 1.2 g/L 1,2-D. Calculate the exact concentration of the stock solution; if the standard is less than 97% pure, make the correction for percent purity as follows (Assume a purity of 96% for the 1,2-D standard for the purpose of this example.):

0.1100 g 1,2-D/0.1 L x 96/100 = 1.056 g/L

b. Repeat step G.1.a. for *cis*-1,3-D and *trans*-1,3-D, preparing a stock solution of approximately 1.2 g/L for each analyte.

c. Using gas-tight syringes, transfer 1/C mL of each stock solution (where C = the concentration of the stock solution in g/L) into a single 10 mL volumetric flask containing approximately 8 mL methanol. Dilute to volume with methanol to obtain a standard solution containing all three analytes, each at a concentration of 100.0 mg/L.

d. Using gas-tight syringes, transfer 1/C mL of each stock solution (where C = the concentration of the stock solution in g/L) into a single 100 mL volumetric flask containing approximately 80 mL methanol. Dilute to volume with methanol to obtain a standard solution containing all three analytes each at a concentration of 10.0 mg/L. For example, a stock solution of 1.056 g/L 1,2-D would be diluted as follows:

$$1/1.056 = 0.947 \text{ mL}$$
$$(947\ \mu L)(1\times10^{-6} L/\mu L)(1.056\ g/L)(1/0.100\ L) = 0.0100\ g/L = 10.0\ mg/L$$

e. Transfer 10.0 mL of the 10.0 mg/L standard solution into a 100 mL volumetric flask. Dilute to volume with methanol to obtain a standard solution containing all three analytes each at a concentration of 1.00 mg/L.

f. Transfer 10.0 mL of the 1.00 mg/L standard solution into a 100 mL volumetric flask. Dilute to volume with methanol to obtain a standard solution containing all three analytes each at a concentration of 0.100 mg/L.

g. Tare a 100-mL volumetric flask containing approximately 80 mL of methanol. Add 11.0-13.0 g of 1,2-D dropwise via Pasteur pipette (approximately eighty drops). Make sure that the liquid falls directly into the methanol and does not run down the inside walls of the volumetric flask. Stopper the flask. Reweigh the flask and calculate the exact weight of the analyte added (Note R.4.). Bring the solution to volume with methanol to yield a stock solution of approximately 120 g/L 1,2-D. Calculate the exact concentration of the stock solution; if the standard is less than 97% pure, make the correction for % purity as follows (Assume a purity of 96% for the 1,2-D standard for the purpose of this example.):

11.0 g 1,2-D/0.100 L x 96/100 = 105.6 g/L

h. Repeat step G.1.g. for *cis*-1,3-D and *trans*-1,3-D, preparing a stock solution of approximately 120 g/L for each analyte.

i. Using gas-tight syringes, transfer 1/(C/100) mL of each stock solution from Section G.1.g. and G.1.h. (where C = the concentration of the stock solution in g/L) into a single 10 mL volumetric flask containing approximately 6 mL methanol. Dilute to volume with methanol to obtain a standard solution containing all three analytes each at a concentration of 10,000 mg/L. For example, a stock solution of 105.6 g/L 1,2-D would be diluted as follows:

$$1/(105.6/100) = 0.947 \text{ mL}$$
$$(947\ \mu L)(1x10^{-6} L/\mu L)(105.6\ g/L)(1/0.010\ L) = 10.0\ g/L = 10000\ mg/L$$

j. Using gas-tight syringes, transfer 1/(C/100) mL of each stock solution G.1.g. and G.1.h. (where C = the concentration of the stock solution in g/L) into a single 100 mL volumetric flask containing approximately 80 mL methanol. Dilute to volume with methanol to obtain a standard solution containing all three analytes each at a concentration of 1000 mg/L.

2. Preparation of the Internal Standard

Following the procedure outlined above in Sections G.1.a. and d., prepare a 10 mg/L solution of the internal standard (IS), 2-bromo-1-chloropropane. (It is not necessary to correct for percent purity of the IS.)

3. Preparation of the Calibration Standard Solutions for the Low-level Dynatrap Analysis

Prepare aqueous calibration standards of the VOAs by transferring aliquots of the standard solutions as outlined below into empty 40-mL volatile analysis vials (Section E.15.). Use an appropriately-sized gas-tight syringe to transfer the standard solution into the vial, add a magnetic stir bar (Section E.10.) to each vial, and cap immediately. The Dynatrap purge and trap will add 10 mL of distilled/deionized water to each standard prior to analysis. **Calibration standards should be prepared just prior to loading onto the Dynatrap purge and trap for analysis.**

a. Low range calibration standards (Section K.1.):

Initial Solution Concentration mg/L[a]	Aliquot of Initial Solution µL	Calibration Std. Final Conc. µg/L	Equivalent Soil Conc. µg/kg[b]
0.100	5.0	0.050	0.10
0.100	10	0.10	0.20
0.100	25	0.25	0.50
0.100	50	0.50	1.0
1.00	10	1.0	2.0
1.00	25	2.5	5.0

[a] Section G.1.e.-f.

[b] Equivalent soil concentration based upon a 5.0 g sample size.

A 10.0 µL aliquot of the 10.0 mg/L IS solution (Section G.2.) must be added to each calibration standard manually using a gas-tight syringe and transferring the IS to the vial containing the standard just prior to sealing, to obtain an IS concentration of 10.0 µg/L. IS addition can be done automatically using the Dynatrap IS addition

feature, in which case a 1.0 µL aliquot of a 100 mg/L IS solution (prepared as described in G.1.c.) must be added to the sample.

b. High range calibration standards (Section K.1.):

Initial Solution Concentration	Aliquot of Initial Solution	Calibration Std. Final Conc.	Equivalent Soil Conc.
mg/L[a]	µL	µg/L	µg/kg[b]
1.00	25	2.5	5.0
1.00	50	5.0	10
1.00	100	10	20
10.0	20	20	40
10.0	40	40	80
10.0	100	100	200

[a] Section G.1.d.-e.
[b] Equivalent soil concentration based upon a 5.0 g sample size.

IS must be added to each calibration standard as described in G.3.a.

4. Preparation of the Calibration Standard Solutions for the High-level Tekmar Analysis

Prepare aqueous calibration standards of the VOAs by transferring aliquots of the standard solutions into 100-mL volumetric flasks containing distilled/deionized water, as outlined below. Use an appropriately-sized gas-tight syringe to transfer the standard solution and inject it into the water below the narrow neck of the flask. Because **aqueous solutions of VOAs are not stable in any container with headspace, each calibration standard should be prepared just prior to purge and trap analysis.**

Calibration standards (Section L.1.):

Initial Solution Concentration	Aliquot of Initial Solution	Calibration Std. Final Conc.	Equivalent Soil Conc.
mg/L[a]	µL	µg/L	µg/kg[b]
1.00	5	0.05	100
1.00	10	0.10	200
1.00	20	0.20	400
1.00	40	0.40	800
1.00	80	0.80	1600
10.0	10	1.0	2000
10.0	50	5.0	10000
100	10	10	20000
100	20	20	40000
100	40	40	80000
100	80	80	160000
100	100	100	200000

[a] Section G.1.c.-e.
[b] Equivalent soil concentration based upon a 5.0 g sample size.

Transfer a 10.0 µL aliquot of the 10.0 mg/L IS solution (Section G.2.) to each calibration standard, to obtain an IS concentration of 1.0 µg/L.

H. Low-level Dynatrap Purge and Trap

1. Purge and Trap Concentrator

Install the Tenax trap (Section E.12.) on the Dynatrap purge and trap concentrator (Section D.7.) following the manufacturer's recommended procedure.

2. Typical Operating Conditions

Instrumentation:	Dynatrap Autosampler/Concentrator
Purge/Carrier Gas:	helium
Purge Flow	40 mL/min
Desorb Flow	20 mL/min
Dynatrap Controller Menu Settings:	
GC cycle time:	23 min
Trap purge ready temperature:	42 °C
Prepurge	0 min
Soil Preheat	70 °C
Preheat stir	Enabled
Preheat time	2 min
Purge time	11 min
Dry purge time	2 min
Desorb preheat	160 °C
Desorb temperature	180 °C
Desorb time	2.0 min
Bake temperature	220 °C
Bake time	6 min
Transfer line temperature	130 °C
Water trap	Enabled
WTKO Cool	45 °C
WTKO Bake	150 °C
Valve oven	110 °C
GC Start	Desorb start
Data start	Desorb start
GC ready	Closed

I. High-level Tekmar Purge and Trap

1. Purge and Trap Concentrator

Install the Tenax trap (Section E.13.) and sparge glassware (Section E.6.) on the Tekmar purge and trap concentrator (Section D.8.) and autosampler (Section D.6.) following the manufacturer's recommended procedure.

2. Typical Operating Conditions

Instrumentation:	Tekmar 3000 Concentrator Tekmar 2016 Autosampler
Purge/Carrier Gas:	helium
Purge Flow	40 mL/min
Desorb Flow	20 mL/min
Trap Pressure	5 psi
Tekmar Controller Menu Settings:	
3000 Transfer Line	140 °C
3000 Valve	140 °C
2016 Transfer Line	130 °C
2016 Valve	130 °C
Moisture control system temp.	130 °C
Trap purge ready temp.	30 °C
Trap purge temp. setting	20 °C (i.e. not heated above ambient temp. during purge)
Sample heater	Off
Prepurge	0.00 min
Preheat time	0.00 min
Purge time	11.00 min
Dry purge time	2.00 min
Moisture control system desorb temp.	45 °C
GC Start	Desorb start
Cryo Focuser	Not applicable
GC cycle time	23 min
Desorb preheat	170 °C
Desorb time	2.00 min
Desorb temp.	180 °C
Sample drain	off
Bake time	10 min
Bake temp.	185 °C
Bake gas bypass	On
Bake gas bypass delay	2.00 min
Moisture control system bake	180 °C

J. Gas Chromatography/Mass Spectrometry (High and Low-level)

1. Column (Note R.5.)

Install the column inlet liner (Section E.2.) and the capillary column (Section E.1.) in the split/splitless injection port of the gas chromatography/mass spectrometer (GC/MSD) following the manufacturer's recommended procedure.

2. Typical Operating Conditions

Instrumentation:	Hewlett-Packard Model 5890 Series II GC Hewlett-Packard Model 5971 Mass Selective Detector Hewlett-Packard Model G1034B Data System Software
Column:	J&W Scientific fused silica capillary DB-VRX liquid phase 30 m x 0.25 mm i.d. 1.4 µm film thickness
Temperatures:	
Column	35 °C for 1.0 min 35 °C to 140 °C at 9 °C/min 140 °C for 0.10 min 140 °C to 210 °C at 20 °C/min 210 °C for 2.0 min
Injector Interface	200 °C 230 °C
Carrier Gas:	helium
Head Pressure Linear Velocity	6 psi approximately 30 cm/sec at 35 °C
Injection Mode:	Split (Note R.5.)
Splitter Flow Septum Purge	20 mL/min Off (capped)
Detector:	electron impact selected ion monitoring
Calibration Program Electron Multiplier	midmass autotune (Note R.6.) 1600 volts

Ions Monitored:

Compound	*m/z*, Quantitation	*m/z*, Confirmation
1,2-D	63	76
cis-1,3-D	75	112
trans-1,3-D	75	112
2-bromo-1-chloropropane	77	--

Dwell Time 75 msec

Full scan mass spectra of the above analytes and IS are shown in Figures 1-4.

3. Typical Chromatograms

Typical chromatograms of a standard, control sample, and a 0.2 µg/kg recovery sample for soil are illustrated in Figures 5-13, respectively.

K. Calibration - Low-level Dynatrap

1. General Approach

The Dynatrap purge and trap which has the capacity to analyze soil samples directly, eliminating the need for an initial solvent extraction, is used for the low-level analysis. Because of the wide range of calibration standard concentrations (0.05-100 µg/L) used in this analysis, it is unlikely that any single calibration curve will provide accurate quantitation over the entire range. In general, deviations from the calculated curve will be most severe at the low end of the curve, which affects measurements near the limit of quantitation. To improve quantitation, it is typically useful to divide the calibration range into two subranges and to produce a separate calibration curve for each. Thus, the low range will encompass VOA standard concentrations of 0.05-2.50 µg/L (equivalent to 0.10-5.0 µg/kg soil sample concentrations), and the high range 2.5-100 µg/L (equivalent to 5.0-200 µg/kg soil sample concentrations).

A calibration check (Section K.4.) must be carried out at the beginning of each 12-hr period during which samples are analyzed in order to confirm calibration of the instrumentation.

2. Initial Calibration

a. Prior to analyzing any standards, bake out the trap at approximately 185 °C and set the GC oven temperature to 220 °C for 10 minutes.

b. Select either the low or high standard range for calibration based upon anticipated levels of the VOAs in the samples being analyzed. After the system has been returned to its starting conditions, load the vials containing the calibration standards for the selected range (Section G.3.) into the Dynatrap autosampler tray. Set the autosampler to add 1.0 µL of a 100 mg/L IS solution (prepared as described in G.1.c.) to each vial (if not added manually during standard preparation) and 10.0 mL of distilled/deionized water. Analyze the calibration standards by purge and trap using GC/MSD as described in Sections H. and J.

3. Calibration Curve

a. Following analysis of the range of calibration standards described in G.3.a. (low range) or G.3.b. (high range), determine the peak areas for 1,2-D (*m/z* 63), *cis*-1,3-D (*m/z* 75), *trans*-1,3-D (*m/z* 75), and 2-bromo-1-chloropropane (IS) (*m/z* 77).

b. For each analyte, prepare a standard curve by plotting the standard concentration (µg/L) on the abscissa (x-axis) and the standard/IS peak area ratio (Quantitation Ratio) on the ordinate (y-axis) as shown in Figures 14-16. Using regression analysis, determine the equation for the curve with respect to the abscissa.

For example, using power regression (5) with the 1,2-D data from Figure 14:

$$Y = \text{constant} \times X^{(\text{exponent})}$$

$$X = \left(\frac{Y}{\text{constant}}\right)^{1/\text{exponent}}$$

where: Y = Quant. Ratio; X = Conc. (µg/L)

$$\frac{\text{1,2 - D Conc.}}{(\mu g / L)} = \left(\frac{\text{Quant. Ratio}}{\text{constant}}\right)^{(1/\text{exponent})}$$

$$\frac{\text{1,2 - D Conc.}}{(\mu g/L)} = \left(\frac{\text{Quant. Ratio}}{0.06498}\right)^{(1/1.02715)}$$

c. Typical calibration curves for the determination of each of the analytes in soil for the low-level, low range analysis are shown in Figures 14-16.

4. Calibration Check

a. For the low calibration range, prepare a 0.25 µg/L calibration standard (Section G.3.a.); for the high range, prepare a 10.0 µg/L calibration standard (Section G.3.b.).

b. Analyze the calibration check standard and calculate the amount of each analyte present using the equation for the current calibration curve for each analyte. (Section K.3.)

c. Calculate the absolute value of the percent difference in the calculated and the theoretical standard concentration as follows:

|(1 - (Calculated/Theoretical))| x 100

If the difference in the calculated and theoretical value for each analyte in the calibration check standard is less than 10%, the existing curves are considered to be valid. Sample analysis may proceed.

d. If the difference in the calculated and theoretical value for any analyte exceeds 10%, the original calibration is no longer considered valid, and a new initial calibration (Section K.2.) must be carried out prior to continuing with sample analysis. A new initial calibration must always be carried out prior to sample analysis whenever major maintenance (column change, source cleaning, filament or multiplier replacement, trap replacement, etc.) is performed on the purge and trap GC/MSD system. Minor maintenance (replacing GC septum or inlet liner, removing a portion of the upper end of the column, etc.) does not automatically necessitate recalibration; however, the system should be evaluated on a case by case basis.

L. Calibration - High-level Tekmar

1. General Approach

The Tekmar autosampler/concentrator (Sections D.6. and 8.), fitted with fritted U-shaped sparge tubes (Section E.6.), is used for the high-level analysis. Soil samples are subjected to an initial solvent extraction, and the resulting extract is diluted with water and analyzed on the Tekmar instrument. A single calibration curve is used in the high-level analysis, covering the range of standard concentrations from 0.05-160 µg/mL (equivalent to 100-320000 µg/kg sample concentration). If problems are encountered in obtaining good correlation of the calibration curve over this range, splitting the curve into two ranges, as done for the low-level method, (Section K.1.) should improve the fit.

A calibration check must be carried out at the beginning of each twelve-hour period during which, samples are analyzed in order to confirm calibration of the instrumentation.

2. Initial Calibration

   a. Prior to analyzing any standards, bake out the trap at approximately 185 °C and set the GC oven temperature to 220 °C for 10 minutes.

   b. After the system has been returned to its starting conditions, load the first aqueous calibration standard (Section G.4.) by removing the plunger from the 25-mL gas-tight Luer Lock syringe (Section E.9.), attaching the closed syringe valve (Section E.11.) to the Luer lock, and pouring the standard into the open end of the syringe until it is nearly full. (Do not draw the standard up into the syringe.) Replace the syringe plunger in the barrel, open the valve, expel any air from the syringe, and adjust the volume to exactly 25.0 mL. Open the loading valve on the Tekmar purge and trap device and load the standard into the sparge tube. Analyze the standard by purge and trap using gas chromatography/mass spectrometry, as described in Sections I. and J.

   c. Repeat L.2.b. for each standard in the calibration range. After each standard has been analyzed, rinse the sparging tube with two approximatley 25-mL aliquots of distilled/deionized water before loading the next standard. Standards should always be analyzed from lowest to highest concentration to minimize carryover. (Note R.7.)

3. Calibration Curve

   a. Following analysis of the range of calibration standards described in G.4.a., determine the peak areas for 1,2-D (*m/z* 63), *cis*-1,3-D (*m/z* 75), *trans*-1,3-D (*m/z* 75), and 2-bromo-1-chloropropane (IS) (*m/z* 77).

   b. For each analyte, prepare a standard curve by plotting the concentration (µg/L) on the abscissa (x-axis) and the standard/IS peak area Quantitation Ratio on the ordinate (y-axis) as shown in Figures 17-19. Using regression analysis, determine the equation for the curve with respect to the abscissa. (See Section K.3.b. for an example calculation.)

   c. Typical calibration curves for the determination of each of the analytes in soil for the high-level analysis are shown in Figures 17-19.

4. Calibration Check

   a. Prepare 0.20 and 20 µg/L calibration check standards (Section G.4.).

   b. Analyze the calibration check standards (Section L.2.b.) and calculate the amount of each analyte present, using the equation for the current calibration curve for each analyte (Section L.3.).

   c. If the difference in the calculated and theoretical value for each analyte in the calibration check standards is less than 10%, the existing curves are considered to be valid. Sample analysis may proceed. (See Section K.4.c. for an example calculation.)

   d. If the difference in the calculated and theoretical value for any analyte exceeds 10%, the original calibration is no longer considered valid, and a new initial

calibration (Section L.2.) must be carried out prior to continuing with sample analysis. A new initial calibration must always be carried out prior to sample analysis whenever major maintenance (column change, source cleaning, filament or multiplier replacement, trap replacement, etc.) is performed on the purge and trap GC/MSD system. Minor maintenance (replacing GC septum or inlet liner, removing a portion of the upper end of the column, etc.) does not automatically necessitate recalibration; however, the system should be evaluated on a case by case basis.

M. Determination of Recovery of 1,2-D, *cis*-1,3-D and *trans*-1,3-D in Soil Using the Low-level Dynatrap Analysis

1. Preparation of Recovery Samples

a. Following analysis of the last calibration standard or check standard, analyze a blank (empty) volatile analysis vial containing a magnetic stir bar. To each vial, add 10 mL distilled/deionized water (automatically added to the vial by the Dynatrap) and 10.0 µL of the 10.0 mg/L IS solution (G.2.) (added manually prior to sealing the vial; if added automatically using the Dynatrap, add 1.0 µL of a 100 mg/L IS solution). Analyze the blank by purge and trap using gas chromatography/mass spectrometry as described in Sections H. and J. Calculate the levels of analytes detected using the current calibration curve for each analyte (Section K.3.). If the concentration of any analyte exceeds the limit of detection or 30 % of the targeted limit of quantitation (Section Q.1.c.), additional blanks should be analyzed until a clean blank is obtained.

b. Obtain a control soil sample from the field sampling location. Weigh a 5.0-g aliquot of the control into an amber volatile analysis vial. Add a magnetic stir bar, IS and 10 mL distilled, deionized water to the sample and analyze it as described for the blank in Section M.1.a., to demonstrate that none of the analytes of interest are detectable in the control sample.

c. Prepare fortified samples at appropriate concentrations by weighing 5.0± 0.05 g aliquots of control soil into amber volatile analysis vials (Note R.8.). Fortify the soils over the range of 0.20-200 µg/kg using the standard solutions, as indicated below:

Low range calibration

Standard Solution Concentration mg/L[a]	Aliquot of Standard Solution µL	Soil Sample Conc. µg/kg	Equivalent Calibration Std. Conc. µg/L
0.10	10	0.2	0.10
0.10	25	0.5	0.25
0.10	50	1.0	0.50
1.0	10	2.0	1.0
1.0	25	5.0	2.5

[a]Section G.1.e.-f.

IS must be added to each calibration standard as described in G.3.a.

High range calibration

Standard Solution Concentration mg/L[a]	Aliquot of Standard Solution µL	Soil Sample Conc. µg/kg	Equivalent Calibration Std. Conc. µg/L
1.0	25	5.0	2.5
1.0	50	10.0	5.0
10.0	10	20.0	10.0
10.0	20	40.0	20.0
10.0	40	80.0	40.0
100.0	10	200.0	100.0

[a]Section G.1.c.-e.

IS must be added to each calibration standard as described in G.3.a.

d. Analyze the fortified recovery samples by purge and trap using gas chromatography/mass spectrometry as described in Sections H. and J.

2. Calculation of Percent Recovery

a. Determine the peak areas for 1,2-D (*m/z* 63, 76), *cis*-1,3-D (*m/z* 75, 112), *trans*-1,3-D (*m/z* 75, 112), and IS (*m/z* 77) for each calibration standard analyzed as part of the current calibration curve (Section K.2.).

b. For each standard, calculate the confirmation ratio for each of the three analytes. The average standard confirmation ratio for each analyte will be used to confirm the presence of that analyte in the soil samples.

For example, using the data for 1,2-D from Figure 5:

Confirmation Ratio = peak area of quantitation ion/peak area confirmation ion

Confirmation Ratio = peak area *m/z* 63 / peak area *m/z* 76

Confirmation Ratio = 1023/538

Confirmation Ratio = 1.90

Positive confirmation of the presence of each analyte in a soil sample is indicated when the confirmation ratio for the sample is in the range of ± 15% of the average found for the standards in the current calibration curve.

c. Prepare a standard curve for each of the analytes and determine the equations for the calibration curves using regression analysis, as described in K.3.

d. Calculate the net concentration (µg/kg) of 1,2-D in each recovery sample by first subtracting the quantitation ratio in the control sample (average ratio if more than one control was analyzed) from that of the recovery sample. Substitute the net quantitation ratio obtained into the calibration curve equation and solve for the concentration (µg/L). Multiply the µg/L purged by the method factor to determine the soil concentration (µg/kg).

For example, using power regression and the 1,2-D data from Figures 8, 11 and 14:

$$\frac{\text{1,2-D Conc.}}{(\mu g / L)} = \left(\frac{(\text{(Sample Quant. Ratio) - (Control Quant. Ratio)})}{\text{constant}}\right)^{1/\text{exponent}}$$

$$\frac{\text{1,2-D Conc.}}{(\mu g / L)} = \left(\frac{(0.00615 - 0)}{0.06498}\right)^{1/1.02715}$$

1,2-D Conc. = 0.10073 µg/L

1,2-D Conc. (µg/kg) = 0.10073 µg/L x Method Factor (L/kg)

where: Method Factor = extraction volume (L) /sample weight (kg)
=0.01 L / 0.00502 kg
=1.99 L/kg

1,2-D Conc. (µg/kg) = 0.10073 µg/L x 1.99 L/kg

1,2-D Conc. = 0.201 µg/kg

e. Determine the percent recovery by dividing the net concentration found for each recovery sample by the theoretical concentration added.

$$\text{Recovery} = \frac{\text{Concentration Found}}{\text{Concentration Added}} \times 100\%$$

$$\text{Recovery} = \frac{0.201\ \mu g / kg}{0.199\ \mu g / kg} \times 100$$

Recovery = 101%

f. For each analyte, determine the net concentration and corresponding percent recovery as described for 1,2-D in Section M.2.d.-e.

The average recovery for each analyte in a given sample set will be used to correct for daily method efficiency.

N. Determination of VOAs in Soil using the Low-level Dynatrap Analysis

1. Analysis of samples

a. Determine the desired calibration range for sample analysis based upon anticipated sample residue levels. Calibrate the instrument/check calibration over the appropriate range for the sample set as described in Section K.

b. Prepare and analyze blank, control, and recovery samples as described in Section M.1. Prior to commencing with analysis of soil samples, analyze a blank volatile analysis vial (Section M.1.a.) to determine that the system is uncontaminated.

c. Weigh 5.0 ± 0.05-g portions of soil samples into amber volatile analysis vials (Note R.8.). Add a magnetic stir bar to each vial. Add 10.0 µL of the 10.0 mg/L IS solution (G.2.) to each sample. (The IS can also be added automatically using the Dynatrap as described in Section M.1.a.) Seal the vials and load them into the carousel of the Dynatrap.

d. Analyze the samples by purge and trap using gas chromatography/mass spectrometry, as described in Sections H. and J.

2. Determination of Soil Moisture

a. Accurately weigh an approximately 10-g portion of soil into a tared weighing dish and record the weight to the nearest 0.01 g.

b. Place the sample in an oven at approximately 110 °C and allow to dry for a minimum of 16 hours.

c. Remove the sample from the oven, place in a desiccator until the sample has cooled to room temperature, and then reweigh.

d. Calculate the percent moisture (dry weight basis) as follows:

$$\text{Percent Moisture (dry weight basis)} = \frac{\text{water(g)}}{\text{dry soil (g)}} \times 100$$

$$= \frac{\text{(sample weight before drying - weight after drying)}}{\text{sample weight after drying}} \times 100$$

3. Calculation of analyte concentration in soil

a. Determine the soil concentration of each analyte by substituting the quantitation ratio into the equation for the corresponding standard calibration curve (Section K.3.), and calculating the µg/L purged. Calculate the uncorrected (gross) µg/kg in the soil by multiplying the µg/L by the method factor.

For example, using the 1,2-D data from Figures 11 and 14, the uncorrected concentration is calculated as follows:

$$\frac{\text{1,2 - D Conc.}}{(\mu g / L)} = \left(\frac{\text{(Quant. Ratio)}}{0.06498}\right)^{1/1.02715}$$

$$\frac{\text{1,2 - D Conc.}}{(\mu g / L)} = \left(\frac{0.00615}{0.06498}\right)^{1/1.02715}$$

1,2-D Conc. = 0.10073 µg/L

1,2-D Conc. (µg/kg) = 0.10073 µg/L x Method Factor (L/kg)

1,2-D Conc. (µg/kg) = 0.10073 x (0.01 L/0.00502 kg)

1,2-D Conc. = 0.201 µg/kg

Correct the amount found for percent moisture of the soil sample as follows:

$$\mu g/kg = (\mu g/kg) \times (1 + (\% \text{ Moisture}/100)$$

b. The uncorrected results as determined in Section N.3.a. must be reported. In addition, the results can be corrected for method recovery using the following procedure:

Calculate the mean % recovery of each analyte for the recovery samples analyzed with the treated samples on the same day.

Determine the corrected analyte concentration in the soil samples as follows (Assume a 1,2-D average percent recovery of 115 for purpose of the example.):

$$\frac{\text{1,2 - D Conc.}}{(\text{corrected } \mu g / L)} = \left(\frac{\text{uncorrected 1,2 - D}(\mu g / kg) \times 100}{\% \text{ Recovery}} \right)$$

$$\frac{\text{1,2 - D Conc.}}{(\text{corrected } \mu g / L)} = \left(\frac{0.201 \times 100}{115} \right)$$

1,2-D Conc. $= 0.175$ µg/kg

c. Any sample giving a quantitation ratio response for one or more of the analytes, which is greater than 10 percent above that of the highest standard in the current calibration range, must be reanalyzed following recalibration of the instrument in the appropriate concentration range. Samples with estimated residues greater than 200 µg/kg should be analyzed using the high-level analysis (Section O.).

d. When analyzing samples using the high range calibration of the low-level analysis, any sample giving a quantitation ratio response for one or more of the analytes, which is greater than 10 percent below that of the lowest standard in the current calibration range, must be reanalyzed following recalibration of the instrument in the appropriate concentration range.

O. Determination of Recovery of 1,2-D, *cis*-1,3-D and *trans*-1,3-D in Soil Using the High-level Tekmar Analysis

1. Preparation of Recovery Samples

a. Following analysis of the last calibration standard/check standard, rinse the sample sparge tube(s) twice with distilled/deionized water. Analyze a blank distilled/deionized water sample by removing the plunger from the 25 mL gas-tight Luer Lock syringe, attaching the closed syringe valve to the Leur Lock, and pouring the water into the open barrel of the syringe until it is almost full. (Do not draw the water up into the syringe.) Replace the syringe plunger in the barrel, open the valve, expel any air from the syringe and adjust the volume to exactly 25.0 mL. Remove the syringe valve, pull the plunger back slightly, and inject 2.5 µL of the 10.0 mg/L IS (Section G.2.) through the Luer Lock tip into the sample. Open the loading valve on the Tekmar purge and trap device, load the sample into the sparge tube, and begin the analysis. (Sections I. and J.) Calculate the levels of analytes detected using the current calibration curve for each analyte (Section L.3.). If the concentration of any analyte exceeds the limit of detection or 30% of the targeted

limit of quantitation, additional blanks should be run until a clean blank is obtained.

b. Obtain a control soil sample from the field sampling location. Weigh 5.0-g portions of the control into 11-dram vials with poly-seal caps (Section E.14.) (Note R.8.). Use one sample as a control and fortify the remaining samples at appropriate levels according to the table below:

Standard Solution Concentration mg/L[a]	Aliquot of Standard Solution µL	Soil Sample Conc. µg/kg	Equivalent Calibration Std. Conc. µg/L
100	10	200	0.1
100	50	1000	0.5
1000	25	5000	2.5
10000	10	20000	10
10000	40	80000	40
10000	80	160000	80

[a]Section G.1.c., i.-j.

c. Add 10.0 mL methanol to each soil sample, seal the vial, vortex for approximately 15 seconds, and centrifuge at 2500 rpm for 3 minutes.

d. Analyze each control and fortified recovery sample by loading the Luer Lock syringe with distilled/deionized water and adjusting the volume to exactly 25.0 mL, as described in section O.1.a. Remove the syringe valve and using a gas-tight syringe transfer 25.0 µL of the soil methanol extract (Section O.c.) into the water through the Luer Lock tip. Inject 2.5 µL of the 10.0 mg/L IS (Section G.2.) through the Luer Lock tip into the water sample. Open the loading valve on the purge and trap device, load the sample into the sparge tube, and begin the analysis. (Sections I. and J.)

2. Calculation of Percent Recovery

a. Calculate the average standard confirmation ratio and standard curves for each analyte as described in M.2.a.-c.

b. Determine the net concentration (µg/kg) of 1,2-D in each recovery sample by first subtracting the quantitation ratio for the control sample (average ratio if more than one control was analyzed) from that of the recovery sample. Substitute the net quantitation ratio obtained into the calibration curve equation and solve for the concentration (µg/L), as shown in Section M.2.d.

Multiply the µg/L purged by the method factor to determine the soil concentration.

$$\text{1,2-D Conc. (µg/kg)} = \text{µg/L} \times \text{Method Factor (L/kg)}$$

where:

$$\text{Method Factor} = \left(\frac{\text{extraction vol. (L)}}{\text{Sample wt. (kg)}}\right) \times \left(\frac{\text{final aqueous vol. (L)}}{\text{methanol aliquot vol. (L)}}\right)$$

$$= \left(\frac{0.01\ L}{0.005\ kg}\right) x \left(\frac{0.025\ L}{0.000025\ L}\right)$$

$$= 2000\ L/kg$$

c. Determine the percent recovery by dividing the net concentration of each recovery sample by the theoretical concentration added, as shown in Section M.2.e.

$$\%\ Recovery = \frac{Concentration\ Found}{Concentration\ Added} x\ 100\%$$

d. For each analyte, determine the net concentration and corresponding percent recovery as described for 1,2-D in Section O.2.b.-c.

The average percent recovery for each analyte in a given sample set will be used to correct for daily method efficiency.

P. Determination of VOAs in Soil using the High-level Tekmar Method

1. Calibrate the instrument/check calibration over the range indicated in section G.4.
2. Prepare and analyze blank, control, and recovery samples as described in Section O.1. Rinse the sample sparge tube(s) with distilled/deionized water. If standards or recovery samples fortified at 10 µg/L or higher were analyzed in any sparge tubes prior to initiating the analysis of a set of samples, analyze a blank as described in section O.1.a. in order to ensure that carryover in each of the tubes is not a problem.
3. Weigh 5.0-g portions of soil samples into 11-dram vials and analyze, as described in Sections O.1. c.-d. (Note R.8.)
4. Calculate the soil concentration (µg/kg) of each analyte by substituting the quantitation ratio into the equation for the corresponding standard calibration curve (Section L.3.), calculating the µg/L purged as shown on Section N.3.a. Calculate the uncorrected (gross) µg/kg in the soil by multiplying the µg/L by the method factor, as follows:

(µg/kg) = µg/L x method factor (L/kg)

where:

$$Method\ Factor = \left(\frac{extraction\ vol.\ (L)}{Sample\ wt.\ (kg)}\right) x \left(\frac{final\ aqueous\ vol.\ (L)}{methanol\ aliquot\ vol.\ (L)}\right)$$

$$= \left(\frac{0.01\ L}{0.005\ kg}\right) x \left(\frac{0.025\ L}{0.000025\ L}\right)$$

$$= 2000\ L/kg$$

5. The uncorrected results as determined in section P.4. must be reported. In addition, the results can be corrected for method recovery using the following procedure (See Section N.3.b. for example.):

a. Calculate the mean % recovery of each analyte for the recovery samples analyzed with the treated samples.

b. Determine the corrected analyte concentration in the soil samples as follows:

$$\frac{\text{1,2 - D Conc.}}{(\text{corrected } \mu g / L)} = \left(\frac{\text{uncorrected 1,2 - D}(\mu g / kg) \times 100}{\text{\% Recovery}} \right)$$

6. Any sample giving a quantitation ratio response, for one or more of the analytes, which is greater than 10 percent below that of the lowest standard in the current calibration range must be reanalyzed using the low-level analysis (Section N.). Any sample giving a quantitation ratio response greater than 10 percent above that of the highest standard in the calibration range must be reanalyzed. A fresh soil aliquot must be extracted with methanol, and the resulting extract diluted appropriately with water to bring the sample response into the calibration range. The additional dilution must then be accounted for in the calculation of the result (Section P.4.) as follows:

1,2-D Conc. ($\mu g/kg$) = $\mu g/L$ x method factor (L/kg) x additional dilution factor

2. Confirmation of Residue Identity

Confirmation of the presence of residues is described in Section M.2.b. For each of the three analytes, confirmation is by comparison of the retention time (gas chromatography) as well as the confirmation ratios resulting from selected ion monitoring (mass spectrometry). Positive confirmation of the presence of each analyte is indicated when the confirmation ratio for the sample is in the range of ± 15% of the average found for the corresponding standards. If additional confirmation is required beyond that discussed in this method, the mass spectrum of each of the three analytes contain additional ions that may be used for confirmation.

3. Assay Time

A typical analytical run would consist of a minimum of six calibration standards for the low-level analysis or 12 for the high-level analysis, a reagent blank, a control (a non-fortified sample), a minimum of two fortified controls (one of which must be at the LOQ when analyzing a sample set in the low range of the low-level analysis), and sixteen samples. This typical analytical run could be prepared in approximately 8 hours, with the purge and trap/chromatographic analysis continuing into the same evening.

As indicated in Sections K.1. and L.1., once the instrument has been calibrated, a calibration check can be done every twelve hours by analyzing one or two standards to verify that the existing calibration curve is still valid. This reduces the amount of time required for daily calibration, and allows the analysis of a greater number of treated samples per day. In addition, the Dynatech can be calibrated in both the low and high range prior to analysis of samples. The sample results can then be calculated using the appropriate curve, increasing the acceptable range for sample responses and ultimately reducing the number of reassays required of samples which fall out of the standard curve range.

R. Notes

1. Equipment, glassware, materials, reagents, and chemicals considered to be equivalent to those specified may be substituted with the understanding that their performance

must be confirmed by appropriate tests. Common laboratory supplies are assumed to be readily available and are, therefore, not listed.

2. The filters are used in the carrier gas supply lines to purify the helium entering the gas chromatograph and sparging apparatus.

3. Vials (40 mL) appropriate for volatile organic analysis are available from a variety of vendors. Regardless of which vendor the vials are purchased from, a check should be done to evaluate the vials and septa for potential interferences in the purge and trap analysis. This is easily done by analyzing three empty vials using the Dynatrap/GC/MSD conditions described in this method and evaluating the resulting chromatograms for interferences at the retention times of the analytes of interest.

4. An alternate method for transferring the neat standards to the methanol contained in the volumetric flask is to tare a gas-tight syringe, draw an amount of the neat standard into the syringe, reweigh the syringe determining the weight of the standard, and transfer the standard from the syringe to the flask by injecting it below the surface of the methanol contained in the flask. Rinse the syringe into the flask to be certain that the analyte contained in the syringe is completely transferred to the flask.

5. The purge and trap concentrator is interfaced with the gas chromatograph by cutting the GC helium inlet line approximately 4 cm from its entry into the injection port. The cut end of the helium line is routed to the concentrator, and the concentrator's transfer line is connected to the remaining short section of tubing leading to the injector. The flow from the purge and trap concentrator is 20 mL/min and the flow through the column is approximately 1 mL/min, therefore a 19:1 split will occur. The injector should be run in the split mode at all times. While this technique appears to sacrifice sensitivity by splitting off 95% of the analytes, it permits the use of normal-dimension capillary columns (0.18 or 0.25 mm i.d.), rather than megabore columns which necessitate the use of jet separators or cryogenic interfaces. By virtue of the superior resolution possible with the 0.25 mm i.d. capillary column, the desired limits of quantitation (0.20 µg/kg) are readily attained for all analytes.

6. The mass spectrometer should be tuned prior to analysis of a set of calibration standards and generation of a calibration curve. Once the instrument has been calibrated it should not be tuned again until just prior to analysis of a fresh set of calibration standards for generation of a new calibration curve.

7. It is possible to load all standards in a calibration curve onto the Tekmar 2016 autosampler prior to beginning the analysis; however, this then requires rinsing of all sparge tubes used and confirming the absence of carryover prior to analyzing samples. Using a single sparge tube to analyze consecutively loaded standards confines the potential for contamination and need for cleaning to a single tube.

8. If soil aliquots are weighed to 5.0 ± 0.05 g, a value of 5.0 may be used to calculate the soil sample concentration. If less precision is used in weighing the soil aliquot, the exact weight must be recorded and used in the calculation of soil concentration (fortified and/or actual).

S. References

1. United States Environmental Protection Agency, "Volatile Organic Compounds by Gas Chromatography/Mass Spectrometry (GC/MS): Capillary Column Technique", Method 8260, Revision 0, July 1992.

2. Stolz, W. L., DowElanco Laboratory Notebook, B065, p 23, 1993, unpublished data of DowElanco.