ENVIRONMENTAL CHEMISTRY METHOD EVALUATION REPORT

NUMBER: ECM 0033W1

AN ELISA IMMUNOASSAY METHOD FOR THE DETERMINATION OF RESIDUES OF METHOMYL IN WATER.

ENVIRONMENTAL CHEMISTRY SECTION (ECS)

ANALYTICAL CHEMISTRY BRANCH

BIOLOGICAL AND ECONOMIC ANALYSIS DIVISION

02/06/95

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PART I

SUMMARY AND CONCLUSION

We have completed an Environmental Chemistry Method Evaluation on methomyl in pond water. This method, Dupont number AMR-2396-92, is an enzyme-linked immunoassay (ELISA) for measurement of methomyl over a range of 0.05 to 5.0 parts per billion (ppb).

Following the suggestion of EFGWB we fortified a water matrix with methomyl at 0.5, 1.0, and 5.0 ppb. In order to evaluate the method at a level near the limit of detection, we also fortified water at 0.125 and 0.25 ppb. All samples were done in replicates of four or more at each level. The Dupont method limit of detection (LOD) of 0.1 ppb and limit of quantitation (LOQ) of 0.2 ppb were validated by our data. The recoveries of methomyl in eighteen samples, at or above the (LOQ), ranged from 76% to 112% with relative standard deviation (RSD) of 10.6%. Although this method involves no extraction or clean-up, we found the recoveries and precision to be good at or above the Limit of Quantitation.

We feel that the method could be used for low-cost monitoring of water for methomyl. However, since there can be some cross-reactivity from other compounds, we emphasize the necessity of confirmatory analysis.

We encountered no problems with the method. However, the precoated microplates and some reagents were obtained from the registrant since they are not readily available from commercial sources. EFED should make sure that Dupont understands that they are responsible for making the precoated microplates commercially available or for licensing their technology to one of the kit manufacturers. They are responsible for making the test kits commercially available to all potential users.

PART II

ANALYTICAL RESULTS FOR METHOMYL

EPA RECOVERIES IN POND WATER

01 02 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	Sample no.	Added (ppb)	Found (ppb)	Recovery Data
03 04 0 0 0 17 0.125 0.09 mean(ppb)=.105 18 0.125 0.10 sd =.0152 19 0.125 0.11 rsd = 14.4% 20 0.125 0.09 mean recovery= 84% 21 0.125 0.11 22 0.125 0.13 23 0.25 0.28 mean(ppb)=.235 24 0.25 0.20 sd =.029 25 0.25 0.21 rsd = 12.3% 26 0.25 0.25 0.21 rsd = 12.3% 27 0.25 0.25 0.24 28 0.25 0.24 28 0.25 0.25 0.24 05 0.5 0.46 mean(ppb) = 0.50 06 0.5 0.49 sd =.029 07 0.5 0.52 rsd = 5.7% 08 0.5 0.52 mean recovery = 100% 09 1.0 0.99 mean(ppb) = 1.02 10 1.0 0.99 mean(ppb) = 1.02 10 1.0 0.99 rsd = 4.7%		0		
04 0 0 0 17				
17				
18	04	0	0	•
18	4=	0.405	0.00	monn(nnh) - 105
19				
20				
21				
0.125 0.13 0.25 0.28 mean(ppb)= .235 24 0.25 0.20 sd = .029 rsd = 12.3% 26 0.25 0.25 0.25 0.25 0.24 28 0.25 0.25 0.24 28 0.25 0.25 0.24 28 0.25 0.25 0.24 28 0.25 0.23 mean(ppb) = 0.50 sd = .029 rsd = 5.7% nean recovery = 100% 0.5 0.5 0.5 0.52 mean recovery = 100% 0.5 0.52 mean recovery = 100%				mean recovery= 84%
23				
24 0.25 0.20 sd = .029 25 0.25 0.21 rsd = 12.3% 26 0.25 0.25 mean recovery = 94% 27 0.25 0.24 28 0.25 0.23 05 0.5 0.46 mean(ppb) = 0.50 06 0.5 0.49 sd = .029 07 0.5 0.52 rsd = 5.7% 08 0.5 0.52 mean recovery = 100% 09 1.0 0.99 mean(ppb) = 1.02 10 1.0 1.01 sd = .048 11 0.99 rsd = 4.7%	22	0.125	0.13	
24 0.25 0.20 sd = .029 25 0.25 0.21 rsd = 12.3% 26 0.25 0.25 mean recovery = 94% 27 0.25 0.24 28 0.25 0.23 05 0.5 0.46 mean(ppb) = 0.50 06 0.5 0.49 sd = .029 07 0.5 0.52 rsd = 5.7% 08 0.5 0.52 mean recovery = 100% 09 1.0 0.99 mean(ppb) = 1.02 10 1.0 1.01 sd = .048 11 0.99 rsd = 4.7%		0.05	0.20	maam/nnh) = 225
25				
0.25		0.25		
0.25				
0.25 0.23 0.5 0.46 mean(ppb) = 0.50 0.5 0.49 sd = .029 0.5 0.52 rsd = 5.7% 0.5 0.52 mean recovery = 100% 0.5 0.99 mean(ppb) = 1.02 1.0 1.0 sd = .048 1.0 0.99 rsd = 4.7%				mean recovery = 94%
05 0.5 0.46 mean(ppb) = 0.50 06 0.5 0.49 sd = .029 07 0.5 0.52 rsd = 5.7% 08 0.5 0.52 mean recovery = 100% 09 1.0 0.99 mean(ppb) = 1.02 10 1.0 1.01 sd = .048 11 1.0 0.99 rsd = 4.7%				
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06 0.5 0.49 sd = .029 07 0.5 0.52 rsd = 5.7% 08 0.5 0.52 mean recovery = 100% 09 1.0 0.99 mean(ppb) = 1.02 10 1.0 1.01 sd = .048 11 1.0 0.99 rsd = 4.7%		0.5	0.46	(
0.5 0.52 rsd = 5.7% 0.52 mean recovery = 100% 0.5 0.52 mean recovery = 100% 0.5 0.52 mean recovery = 100% 0.5 0.52 mean recovery = 100% 1.0 0.99 mean(ppb) = 1.02 sd = .048 sd = .048 sd = .048 sd = 4.7%				
08 0.5 0.52 mean recovery = 100% 09 1.0 0.99 mean(ppb) = 1.02 10 1.0 1.01 sd = .048 11 1.0 0.99 rsd = 4.7%				
09 1.0 0.99 mean(ppb) = 1.02 10 1.0 1.01 sd = .048 11 1.0 0.99 rsd = 4.7%				
10 1.0 1.01 sd = .048 11 1.0 0.99 rsd = 4.7%	08	0.5	0.52	mean recovery = 100%
10 1.0 1.01 sd = .048 11 1.0 0.99 rsd = 4.7%	00	1 0	0 99	mean(nnh) = 1.02
11 1.0 0.99 $rsd = 4.7\%$				
12 1.0 1.09 mean recovery - 102%				
	12	1.0	1.09	mean recovery - 102%
13 5.0 4.20 mean(ppb) = 4.25	13	5.0	4.20	mean(ppb)= 4.25
14 5.0 3.82 sd = .417				
15 5.0 4.15 rsd = 9.8%				
16 5.0 4.82 mean recovery = 85%				

PART III

EXPERIMENTAL SUMMARY

(a) Principle of Method

Polyclonal anti-methomyl antibodies(Ab) and buffer are added to a sample containing an unknown amount of methomyl and incubated. The antibodies bind to methomyl molecules present in the sample (if any).

Aliquots of the solution are added to wells on a 96-well microplate, which have been coated with a methomyl derivative-Ovalbumin conjugate. Any excess (Ab) not bound to methomyl in the sample will bind to methomyl immobilized on the microwell plate. The plate is then washed to remove any (Ab) not bound to the plate. The amount of (Ab) bound to the microwell is an inverse measure of the amount of methomyl in the sample.

To detect the (Ab) bound to each microwell, an anti-rabbit antibody conjugated to an enzyme, alkaline phosphatase(Ab-E), is added to each well and incubated. This (Ab-E) will bind to any anti-methomyl antibodies bound to the microwell. The microwell plate is then washed to remove any unbound (Ab-E).

The alkaline phosphatase substrate, para-nitrophenyl-phosphate, is added to the microwells. The enzyme-substrate reaction produces a yellow color which is inversely proportional to the concentration of methomyl in the sample. The Microplate Reader quantitates absorbance in each well at 405 nm. Computer software is used to construct a standard curve from standards run on the plate, and to calculate the methomyl concentrations of each unknown on the plate.

(b) Source of Analytical Reference Standard

Methomyl standard was supplied and certified at 99.77% by E.I.Dupont de Nemours & company, Wilmington, Del.

(c) Source of Sample Matrix

Pond water was obtained locally from a pond at Stennis Space Center. The water was collected in pre-cleaned one gallon dark glass bottles. The water was kept refrigerated and brought to ambient temperature before aliquots were taken for fortification and analysis.

(d) Instrumentation for Quantitation

- 1. Vmax Kinetic Microplate Reader with 405 nm optical filter, Molecular Devices Corporation.
- 2. ULTRAWASH PLUS, automatic microplate washer/aspirator, Dynatech Laboratories, Inc.

(e) Modification of Method

We used microplates supplied by Dupont and already coated with Coating Antigen Reagent. The method includes a procedure for coating blank microplates.

(f) <u>Calculations</u>

The absorbance of each of the 96 wells of the microwell plate is read on a microplate reader equipped with a 405 nm filter and processed by a computer program which generates a standard curve based upon a four-parameter logit function. The standard four-parameter logit curve generated is sigmoidal in shape with optical density (OD) on the Y-axis and the log of the concentration of methomyl on the X-axis. Each calibration standard and sample is pipetted into three wells of the microplate and analyzed in triplicate. The average of the optical density (OD) readings for these three wells is then used to interact with the standard curve to calculate the concentration of methomyl in each sample. For comparison, a semi-log curve fit would provide reasonably good data and can be used to check the calculations.

(g) Graphs and Data

The following pages contain a print-out of the standard curve and data generated by the methomyl calibration standards and selected samples.

MOLECULAR DEVICES **Analyzed Curve**

DATA FILE: DESCRIPTION: 11089405

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PROTOCOL: DESCRIPTION:

WAVELENGTH: CALIBRATION:

MODE:

methornyl

On

Endpoint 405

AUTOMIX:

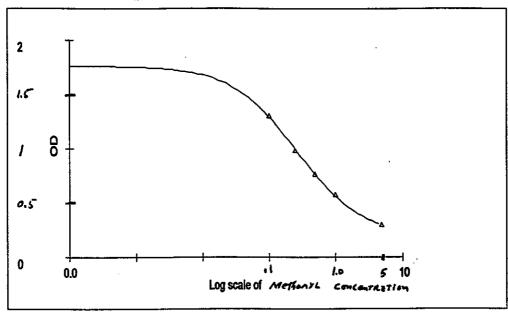
On

Curve Fit: 4-Parameter

Corr. Coeff: 1.00

 $y = (A-D)/(1 + (x/C)^B) + D$

A= 1.77 B= 0.886 C= 0.268 D= 0.191



STANDARD STD01	Std. Value 0	Well B2 C2 D2	OD 1.851 1.761 1.687	Mean 1.767	Std Dev 0.082	CV 4.649	Calc. Value <<<< 5.68e-4 0.010	Sample ID
STD02	0.1	B3 C3 D3	1.370 1.283 1.283	1.312	0.050	3.827	0.079 0.107 0.107	
STD03	0.25	B4 C4 D4	1.039 0.964 0.965	0.990	0.043	4.346	0.226 0.280 0.279	
STD04	0.5	85 C5 D5	0.791 0.769 0.743	0.768	0.024	3.128	0.464 0.496 0.538	
STD05	1	B6 C6 D6	0.579 0.569 0.573	0.574	0.005	0.877	0.947 0.984 0.969	
STD06	5	B7 C7 D7	0.309 0.280 0.302	0.297	0.015	5.088	4.567 6.418 4.920	•••

MOLECULAR DEVICES Raw Data (Report)

DATA FILE: DESCRIPTION: PROTOCOL: 11089405

methornyl

PROTOCOL:
DESCRIPTION:
MODE:
WAVELENGTH:
CALIBRATION:

Endpoint 405 On

AUTOMIX:

On

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PLATE BLANK BL	Mean OD 0.113	Std Dev 0.002	CV 1.578	Well A2 A3 A4 A5 A6 A7 A8 A9 A10	OD 0.114 0.111 0.113 0.113 0.111 0.115 0.109 0.113 0.114	Substrate Blanks
STANDARDS STD01	Mean OD 1.767	Std Dev 0.082	CV 4.649	Well B2 C2 D2	OD 1.851 1.761 1.687	Sample ID O.O ns/ml
STD02	1.312	0.050	3.827	B3 C3 D3	1.370 1.283 1.283	0.1 mg/ml
STD03	0.990	0.043	4.346	B4 C4 D4	1.039 0.964 0.965	0.25 ns/ml
STD04	0.768	0.024	3.128	85 C5 D5	0.791 0.769 0.743	0.50 ms/ml 1.0 ms/ml 5.0 ms/ml
STD05	0.574	0.005	0.877	B6 C6 D6	0.579 0.569 0.573	1.0 ms/ml
STD06	0.297	0.015	5.088	B7 C7 D7	0.309 0.280 0.302	5.0 ns/ml
UNKNOWNS UNK01	Mean OD 1.802	Std Dev 0.079	CV 4.372	Well B8 C8 D8	OD 1.876 1.809 1.719	Sample ID UMFORT I FIED WATER
UNKO2	1.794	0.092	5.145	C9 D9	1.876 1.812 1.694	1.7
UNK03	1.754	0.055	3.108	B10 C10 D10	1.798 1.771 1.693	1+
UNK04	1.851	0,065	3.531	B11 C11 D11	1.920 1.843 1.790	<i>f</i> 1

MOLECULAR DEVICES

UNKNOWNS UNK05	Mean OD 1.037	Std Dev 0.019	CV 1.821	Well E2 F2 G2	OD 1.032 1.020 1.057	Dil.Factor 2.000	. <u>PP b</u> 0.455	Sam Foktifiel A	ple ID TO.5 pp b
UNK06	1.016	0.043	4.243	E3 F3 G3	1.054 1.024 0.969	2.000	0.485	"	
UNK07	0.988	0.037	3.715	E4 F4 G4	1.026 0.953 0.983	2.000	0.525	17	
UNK08	0.993	0.011	1.086	E5 F5 G5	0.985 1.005 0.988	2.000	0.516	. * 1	
 UNK13	0.411	0.019	4.541	E10 F10 G10	0.408 0.394 0.431	2.000	4.196	FONTIFIEL AT	5.0 ppb
UNK14	0.427	0.005	1.057	E11 F11 G11	0.426 0.422 0.431	2.000	3,815		
UNK15	0.413	0.020	4.759	H2 H3 H4	0.416 0.392 0.431	2.000	4.150	, 1	
UNK16	0.390	0.021	5,335	H5 H6 H7	0.374 0.413 0.381	2.000	4.816	_f 1	·