

Cover Sheet for

Environmental Chemistry Method

Pesticide Name: Iodosulfuron + Metsulfuron methyl

MRID#: 451085-22

Matrix: Water

Analysis: LC/MS/MS

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C006395

Title

Enforcement Method of Iodosulfuron-methyl-sodium and its metabolite Metsulfuron-methyl in Surface Water by HPLC incl. Validation

Extension of the enforcement method EM F 01/98 - 0 for Iodosulfuron-methyl-sodium in Drinking Water to its metabolite Metsulfuron-methyl incl. Validation

451087-22

Iodosulfuron-methyl-sodium (AE F115008) Metsulfuron-methyl (AE F075736)

Guideline Référence

EPA OPPTS 860.1340

OECD IIA 4.55

Author

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Completed On
08 Feb 2000

Test Facility

Hoechst Schering AgrEvo GmbH
Entwicklung
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Federal Republic of Germany

Study Identification

VOL. 66 of 229

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Date: March 20, 2000



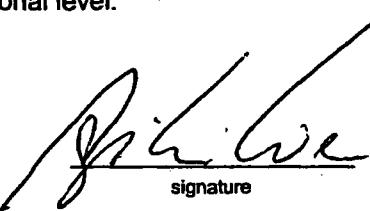
Method EM F12/99-0
Page 3

GOOD LABORATORY PRACTICE COMPLIANCE STATEMENT

The validation of this analytical method was conducted in compliance with the *Principles of Good Laboratory Practice* as adopted by the Council of OECD on 12 May 1981 [C(81)30 (Final)] and implemented at the national level.

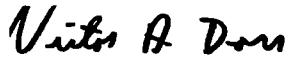
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Sponsor/Submitter

March 20, 2000

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Page: 4 (86)

GLP Quality Assurance

8.2.2000

Quality Assurance Statement

Title: **Enforcement Method of Iodosulfuron-methyl-sodium and its metabolite Metsulfuron-methyl in Surface Water by HPLC incl. Validation**

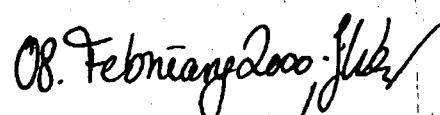
Extension of the enforcement method EM F 01/98 - 0 for Iodosulfuron-methyl-sodium in Drinking Water to its metabolite Metsulfuron-methyl incl. Validation

**Iodosulfuron-methyl-sodium (AE F115008)
Metsulfuron-methyl (AE F075736)**

Study: CR99/029

This study was periodically inspected and properly signed records of these inspections were submitted to testing facility management and the study director as shown below. This report has been audited by the Quality Assurance unit. The reported results accurately reflect the original data of the study.

Inspection	Phase of Study	Reported
18.11.1999	study plan	18.11.1999
22.11.1999	study conduct	22.11.1999
24.11.1999	study conduct	24.11.1999
17.1.2000	study conduct	17.1.2000
4.2.2000	draft report	7.2.2000
8.2.2000	final report	8.2.2000



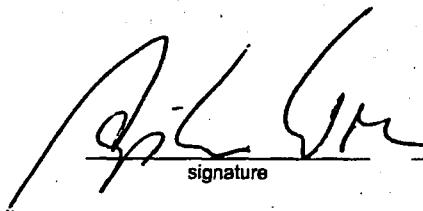
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GLP Quality Assurance

APPROVALS PAGE

Study Director
and Author

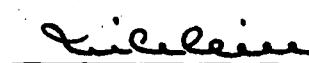
Dr. A. Wrede


signature

08.02.2000
date (d/m/y)

Head of
Test Facility

Dr. M. Uihlein


signature

09.02.2000
date (d/m/y)

Audited by
Quality Assurance Unit

08. Februar 2000; lk

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SUMMARY

Enforcement Method of Iodosulfuron-methyl-sodium and its metabolite Metsulfuron-methyl in Surface Water by HPLC incl. Validation

Extension of the enforcement method EM F 01/98 - 0 for Iodosulfuron-methyl-sodium in Drinking Water to its metabolite Metsulfuron-methyl incl. Validation

Iodosulfuron-methyl-sodium (AE F115008) Metsulfuron-methyl (AE F075736)

Relevant residue

Iodosulfuron-methyl-sodium (AE F115008)
Metsulfuron-methyl (AE F075736)

Test commodity

Drinking water
Surface water

Principle of the method

Metsulfuron-methyl in drinking water:

The water sample is adjusted to pH 2.5 with phosphoric acid (2 N) and sucked through an C18-cartridge (conditioned with 5 mL methanol and 5 mL water). AE F075736 is eluted with 5 mL methanol. Metsulfuron-methyl in the final solution in acetonitrile/water (1/1, v/v) is determined by HPLC/UV.

Metsulfuron-methyl and iodosulfuron-methyl-sodium in surface water:

The water sample is adjusted to pH 2.5 with phosphoric acid (2 N) and filtered through a glass microfibre filter and a cellulose nitrate filter (0.45 µm). The sample is sucked through a NH₂ / C18-cartridge (conditioned with 5 mL methanol and 5 mL water). AE F115008 and AE F075736 are eluted with 15 mL methanol/water (60:40, v/v). After reducing to dryness, the residue is dissolved in 20 mL toluene and sucked through a Silicagel-cartridge (conditioned with 5 mL toluene). AE F115008 and AE F075736 are eluted with 30 mL toluene/methanol (95:5, v/v). Iodosulfuron-methyl-sodium and metsulfuron-methyl in the final solution in acetonitrile/water (1/1, v/v) are determined by HPLC/UV.

Calibration

A curve of the form $y = a + bx + cx^2$ is applicable over the tested range of 0.1 to 2.0 µg metsulfuron-methyl/mL and 0.1 to 2.0 µg iodosulfuron-methyl-sodium/mL.

Recovery efficiency, relative standard deviation (RSD)

Recovery experiments were conducted at 0.1 µg/L and 1.0 µg/L for AE F075736 in drinking water and at 0.1 µg/L and 1.0 µg/L for AE F075736 and AE F115008 in surface water.

The mean recoveries for AE F075736 in drinking water were found at 79 and 92 %, with an RSD of up to 9 %.

In surface water the mean recoveries for AE F075736 were found at 85 and 105 %, with an RSD of up to 10 %, the confirmation method gave mean recovery values of 92 and 106 %, with an RSD of up to 18 %.

In surface water the mean recoveries for AE F115008 were found at 92 and 103 %, with an RSD of up to 9 %, the confirmation method gave mean recovery values of 85 and 96 %, with an RSD of up to 20 %.

The results are summarized in the table below.

Limit of quantification

The limit of quantification (LOQ) for metsulfuron-methyl in drinking water was established and validated at 0.1 µg/L.

The limit of quantification (LOQ) for iodosulfuron-methyl-sodium and metsulfuron-methyl in surface water was established and validated at 0.1 µg/L.

Specificity

Control samples of drinking water were analysed for AE F075736 and control samples of surface water were analysed for AE F115008 and AE F075736. In none of the samples apparent residues were determined.

The specificity of the method was demonstrated by a confirmatory technique using a HPLC column with a different stationary phase.

Summary table of recoveries:

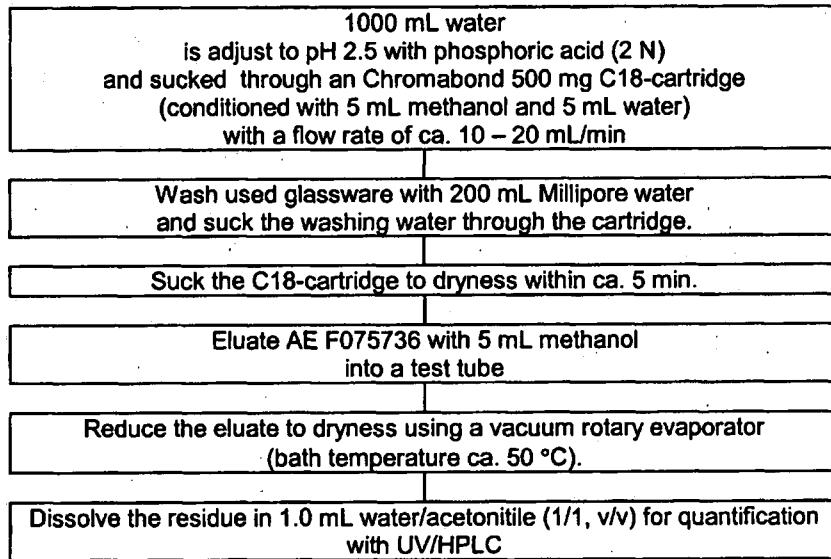
Analyte	Matrix	Fortification level [µg/L]	Recovery [%]	Mean recovery [%]	RSD (a) [%]	n
AE F075736	Drinking water (Vittel)	0.10	82, 75, 87, 83, 70	79	9	5
		1.0	96, 91, 87, 93, 93	92	4	5
	Surface water	0.10	90, 78, 91, 92, 74	85	10	5
		1.0	98, 102, 105, 109, 109	105	5	5
	Surface water Confirmation method	0.10	74, 110, 92, 106, 108, 90, 67	92	18	7
		1.0	102, 106, 106, 108, 109	106	3	5
AE F115008	Surface water	0.10	108, 92, 94, 106, 113	103	9	5
		1.0	89, 90, 90, 94, 95	92	3	5
	Surface water Confirmation method	0.10	75, 78, 74, 114, 98, 73	85	20	6
		1.0	92, 97, 97, 97, 98	96	2	5

a) RSD: relative standard deviation

RSD = SD / Mean recovery • 100 %

Analytical method flow sheet**Metsulfuron-methyl in drinking water**

*Extraction AE F075736 and
C18-cartridge clean-up*



Iodosulfuron-methyl-sodium and metsulfuron-methyl in surface water**Extraction**

AE F115008 and
AE F075736 and
NH₂ / C18-cartridge
clean-up

1000 mL water
is adjust to pH 2.5 with phosphoric acid (2 N),
filtered through a glass microfibre filter and a cellulose nitrate filter (0.45 µm)
the sample is sucked through a Chromabond 500 mg NH₂ / 500 mg C18-cartridge
(conditioned with 5 mL methanol and 5 mL water)
with a flow rate of ca. 10 – 20 mL/min

Wash used glassware with 200 mL Millipore water
and suck the washing water through the cartridge.

Suck the NH₂ / C18-cartridge to dryness within ca. 5 min.

Wash the NH₂ / C18-cartridge with 10 mL methanol/water (30:70, v/v).

Eluate AE F115008 and AE F075736 with 15 mL methanol/water (60:40, v/v)

Reduce the eluate to dryness using a vacuum rotary evaporator
(bath temperature ca. 50 °C).

**Silicagel-cartridge
clean-up**

Dissolve the residue in 20 mL toluene (if necessary use an ultrasonic bath)
suck through a Silicagel-cartridge (conditioned with 5 mL toluene)

Discard the eluate, suck the Silicagel-cartridge to dryness

Wash the round-bottom flask with 30 mL toluene/methanol (95:5, v/v)
and elute AE F115008 and AE F075736 with this solution
Reduce the eluate to dryness using a vacuum rotary evaporator
(bath temperature ca. 40 °C).

HPLC

Dissolve the residue in 1.0 mL water/acetonitrile (1/1, v/v) for quantification with
UV/HPLC

1 Organization and Personnel

Sponsor: AgrEvo GmbH
D - 65926 Frankfurt am Main

Test facility: AgrEvo GmbH
Rückstände und Verbrauchersicherheit
D - 65926 Frankfurt am Main
Head: Dr. M. Uihlein

Study No.: CR 99/029
Study director: Dr. A. Wrede
Address: see Test facility

Method No.: EM F 12/99-0

Archiving:

The study plan, the raw data and the original report will be archived at

Hoechst Schering AgrEvo GmbH
Rückstände und Verbrauchersicherheit
D-65926 Frankfurt am Main

Retained samples of the test and reference substances for the analytical work will be archived at

Hoechst Schering AgrEvo GmbH
Produktanalytik
D-65926 Frankfurt am Main

for at least the duration of the period prescribed in the GLP guidelines.

Quality assurance:

Hoechst Schering AgrEvo GmbH
GLP-Qualitätssicherungseinheit
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Time schedule:

Study plan signed by study director: 29 Sep 1999
Start of analysis: 22 Nov 1999
End of analysis: 25 Jan 2000

2 Objectives

The objective of this study was the development and validation of an analytical method for the determination of residues of metsulfuron-methyl (AE F 075736) in drinking water and metsulfuron-methyl (AE F075736) and iodosulfuron-methyl-sodium (AE F115008) in surface water by HPLC using UV-detection.

The validation for AE F115008 in drinking water is given in the analytical method EM F 01/98 – 0 (ref. 1).

3 Test commodities

The drinking water used for the validation was Vittel.

The samples of surface water used for the validation were taken from the small lake at building F821 (Industriepark Höchst).

The characteristics¹ of the surface water are:

pH	7.90 ± 0.02
DOC ¹ [mg/L] ¹	7.3 ± 0.3
total hardness [° dH] ²	10.1

¹ dissolved organic content

² degree german hardness [mg CaO+MgO / 100 mL water]

4 Relevant residue and reference substances

4.1 Relevant residue

The relevant residue consists of the metabolite metsulfuron-methyl (AE F075736) and the parent compound iodosulfuron-methyl-sodium (AE F115008).

¹ Determination of the characteristics of the surface water was not done under GLP.

4.2 Test and reference substances

Iodosulfuron-methyl-sodium (AE F115008)

Chemical name (IUPAC): methyl 4-iodo-2-[3-(4-methoxy-6-methyl-1,3,5-triazin-2-yl)ureido-sulfonyl]benzoate, sodium salt

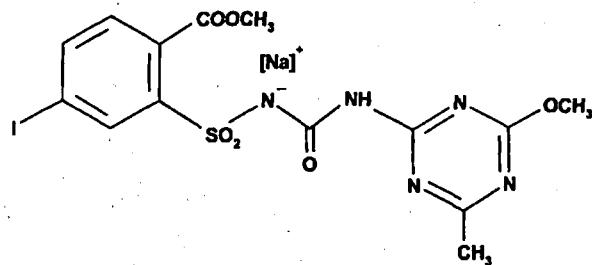
Empirical formula: C₁₄H₁₃IN₅NaO₈S

Molecular weight: 529.3

Solubility (20 °C):

Solvent	Solubility	Source
acetone	> 380 g/L	ref. 2
dichloromethane	> 500 g/L	ref. 2
ethyl acetate	23 g/L	ref. 2
n-hexane	1.2 • 10 ⁻³ g/L	ref. 2
methanol	12 g/L	ref. 2
n-heptane	1.1 • 10 ⁻³ g/L	ref. 2
2-propanol	4.4 g/L	ref. 2
toluene	2.1 g/L	ref. 2
acetonitrile	52 g/L	ref. 2
DMSO	> 500 g/L	ref. 2
PEG	87 g/L	ref. 2

Structural formula:



Certificate of analysis:

AZ 07931

Drawn up by:

Hoechst Schering AgrEvo GmbH

Produktanalytik

D-65926 Frankfurt am Main, Germany

97.3 % (w/w)

Purity:
Expiry date (d/m/y):

30 May 2000

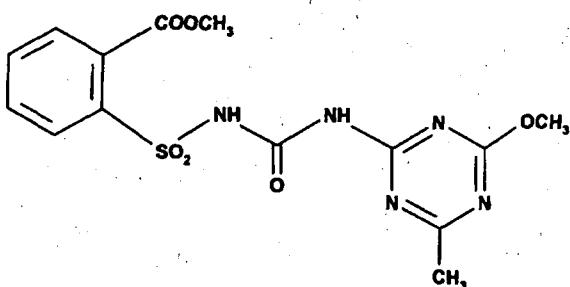
Metsulfuron-methyl (AE F075736)

Chemical name (IUPAC): methyl 2-[3-(4-methoxy-6-methyl-1,3,5-triazin-2-yl) ureidosulfonyl]-benzoate

Empirical formula: C₁₄H₁₅N₅O₆S

Molecular weight: 381.4

Structural formula:



Certificate of analysis: AZ 06892

Drawn up by: Hoechst Schering AgrEvo GmbH

Produktanalytik

D-65926 Frankfurt am Main, Germany

98.4 % (w/w)

07 May 2000

Purity:

Expiry date (d/m/y):

5 Procedures

5.1 Principle of analytical method

The method flow sheets are presented in Appendix I.

Metsulfuron-methyl in drinking water:

The water sample is adjust to pH 2.5 with phosphoric acid (2 N) and sucked through an C18-cartridge (conditioned with 5 mL methanol and 5 mL water). AE F075736 is eluted with 5 mL methanol. AE F075736 in the final solution in acetonitrile/water (1/1, v/v) is determined by HPLC/UV.

Metsulfuron-methyl and iodosulfuron-methyl-sodium in surface water:

The water sample is adjust to pH 2.5 with phosphoric acid (2 N) and filtered through a glass microfibre filter and a cellulose nitrate filter (0.45 µm). The sample is sucked through a NH₂ / C18-cartridge (conditioned with 5 mL methanol and 5 mL water). AE F115008 and AEF075736 are eluted with 15 mL methanol/water (60:40, v/v). After reducing to dryness, the residue is dissolved in 20 mL toluene and sucked through a Silicagel-cartridge (conditioned with 5 mL toluene). AE F115008 and AE F075736 are eluted with 30 mL toluene/methanol (95:5, v/v). AE F115008 and AE F075736 in the final solution in acetonitrile/water (1/1, v/v) are determined by HPLC/UV.

5.2 Reagents

- methanol Chromasolv, cat. No. 34860 (Riedel-de Haën, Germany)
- acetonitrile Chromasolv p.A., cat. No. 34851 (Riedel-de Haën, Germany)
- deionized water
- water (e.g. prepared with Milli-Q-Plus, Millipore)
- phosphoric acid 2 N, cat. No. 30417 (Riedel-de Haën, Germany)
- toluene Pestanal, cat. No. 34494 (Riedel-de Haën, Germany)
- AE F075736, analytical standard (AgrEvo GmbH, Germany)
- AE F115008, analytical standard (AgrEvo GmbH, Germany)
- C18 – cartridge, 500 mg, cat. No. 730013 (Chromabond)
- Glass microfibre filter, cat. No. 1827070 (Whatman)
- Cellulose nitrate filter (0.45 µm), cat. No. 11306-50-N (Satorius)
- NH₂/C18 – cartridge, 500mg NH₂, 500mg C18, cat. No. 730618 (Chromabond)
- Silicagel-cartridge, 500 mg, ISOLUTE, cat. No. 460-0050-H (ICT)

Stock solutions of the analytical standards were prepared by dissolving about 50 mg of analytical standard of AE F075736 and 50 mg of the analytical standard of AE F115008 in ca. 50 mL acetonitrile / triethylamine (0.02 mol/L), 4:1, v/v. Concentration of the stock solutions was 1.0 mg/mL. Working solutions were prepared from the stock solution by further dilution with acetonitrile / water, 1:1, v/v.

5.3 Apparatus

The following list contains the apparatus used in the laboratory of the author for validation.
Suitable alternatives can be taken.

- standard laboratory glassware
- rotary vacuum evaporator with water bath
- HPLC system with UV-detector
- chromatography column, Prodigy ODS, 150 mm x 4.6 mm, 5 µm
- chromatography column, Nucleosil C18, 5 µm, 250 mm x 4 mm (confirmation method)

5.4 Preparation of samples and storage

The samples of drinking water (Vittel) were bought November 1999.

The samples of surface water were taken from the small lake at building F821 (Industriepark Höchst) on 14 Jan 2000.

Samples were stored at room temperature.

5.5 Laboratory steps

5.5.1 Metsulfuron-methyl in drinking water

5.5.1.1 Extraction and C18-cartridge clean up

1000 mL of the water sample is adjust to pH 2.5 with phosphoric acid (2 N) and sucked through a Chromabond 500 mg C18-cartridge (conditioned with 5 mL methanol and 5 mL water) with a flow rate of ca. 10 – 20 mL/min. Wash used glassware with 200 mL Millipore water and suck the washing water through the cartridge. Suck the C18-cartridge to dryness within ca. 5 min. Eluate AE F075736 with 5 mL methanol into a test tube. Reduce the eluate to dryness using a vacuum rotary evaporator (bath temperature ca. 50 °C).

5.5.1.2 Preparation of the final solution

Dissolve the residue in 1.0 mL acetonitrile/water (1/1, v/v).

5.5.1.3 Determination of residues

The following conditions have been used successfully during validation of this analytical method. If different equipment and columns are used, modifications of the given conditions may be necessary.

HPLC-conditions

Instrument:	Beckmann
System	IBM-PC System 2 8570 Model 70386
Pump:	226 Beckmann
Detector:	Diode Array Detector 168 Beckmann
Injector:	Autosampler 507 Beckmann
Injection volume:	100 µL
Column temperature:	30 °C
Column	Prodigy ODS, 5 µm, 150 mm x 4,6 mm
Wavelength:	233 nm
Flow rate:	1.0 mL/min
Mobile phase:	
Eluent A	Acetonitrile Chromasolv
Eluent B	Phosphoric acid $C_{H_3PO_4} = 0.01\text{mol/L}$

Gradient program for the determination of AE F075736

Time [min]	Total flow pump A + B [mL/min]	Pump A (eluent A)		Pump B (eluent B) phosphoric acid $C_{H_3PO_4} = 0.01\text{mol/L}$
		[%]	[%]	
0	1.0	20		80
10	1.0	50		50
20	1.0	50		50
30	1.0	80		20
35	1.0	80		20
45	1.0	20		80
47	1.0	20		80
55	1.0			

Under these conditions the retention time for AE F075736 is about 21.0 min.

The chromatography data were recorded and evaluated with TURBOCHROM® Client/Server system, PERKIN ELMER.

5.5.2 Iodosulfuron-methyl-sodium and metsulfuron-methyl in surface water

5.5.2.1 Extraction and NH₂/C18-cartridge clean up

1000 mL of the water sample is adjust to pH 2.5 with phosphoric acid (2 N) and filtered through a glass microfibre filter and a cellulose nitrate filter (0.45 µm). The sample is sucked through a Chromabond 500 mg NH₂ / 500 mg C18-cartridge (conditioned with 5 mL methanol and 5 mL water) with a flow rate of ca. 10 – 20 mL/min. Wash used glassware with 200 mL Millipore water and suck the washing water through the cartridge. Suck the NH₂ / C18-cartridge to dryness within ca. 5 min. and wash the NH₂ / C18-cartridge with 10 mL methanol/water (30:70, v/v). Eluate AE F115008 and AE F075736 with 15 mL methanol/water (60:40, v/v) into a round-bottom flask. Reduce the eluate to dryness using a vacuum rotary evaporator (bath temperature ca. 50 °C).

5.5.2.2 Silicagel-cartridge clean up

Dissolve the residue in 20 mL toluene (if necessary use an ultrasonic bath) and suck through a Silicagel-cartridge (conditioned with 5 mL toluene). Discard the eluate, suck the Silicagel-cartridge to dryness. Wash the round-bottom flask with 30 mL toluene/methanol (95:5, v/v), if necessary use an ultrasonic bath, and elute AE F115008 and AE F075736 with this solution. Reduce the eluate to dryness using a vacuum rotary evaporator (bath temperature ca. 40 °C).

5.5.2.3 Preparation of the final solution

Dissolve the residue in 1.0 mL acetonitrile/water (1/1, v/v).

5.5.2.4 Determination of residues

The following conditions have been used successfully during validation of this analytical method. If different equipment and columns are used, modifications of the given conditions may be necessary.

HPLC-conditions

Instrument:	Beckmann
System	IBM-PC System 2 8570 Model 70386
Pump:	226 Beckmann
Detector:	Diode Array Detector 168 Beckmann
Injector:	Autosampler 507 Beckmann
Injection volume:	100 µL
Column oven:	Beckmann
Column temperature:	30 °C
Column	Prodigy ODS, 5 µm, 150 mm x 4,6 mm (validation) Nucleosil C18, 5 µm, 250 mm x 4 mm (confirmation method)
Wavelength:	233 nm
Flow rate:	1.0 mL/min
Mobile phase:	
Eluent A	Acetonitrile Chromasolv
Eluent B	Phosphoric acid c _{H3PO4} = 0.01mol/L

Gradient program for the determination of AE F075736

Time [min]	Total flow pump A + B [mL/min]	Pump A (eluent A)	Pump B (eluent B)
		Acetonitrile Chromasolv	phosphoric acid $\text{CH}_3\text{PO}_4 = 0.01\text{ mol/L}$
0	1.0	20	80
10	1.0	50	50
20	1.0	50	50
30	1.0	80	20
35	1.0	80	20
45	1.0	20	80
47	1.0	20	80
55	1.0		

Under these conditions the retention time for AE F075736 is about 21.0 min and for AE F115008 about 25.4 min.

The chromatography data were recorded and evaluated with TURBOCHROM® Client/Server system, PERKIN ELMER.

Confirmatory method

For confirmatory purposes a different stationary phase was used:

HPLC-Column: Nucleosil C18, 5 μm , 250 mm x 4 mm

Under these conditions the retention time for AE F075736 is about 21.5 min and for AE F115008 about 25.5 min.

5.6 Calibration

The concentration of AE F075736 and AE F115008 were calculated using external standards at 4 different concentrations over a range from 0.1 ng/ μL up to 1 or 2 ng/ μL . The lowest concentration was 0.1 ng/ μL . The highest concentration was 2 ng/ μL .

The recommended order of samples / test solutions for setting up a sequence for HPLC-determination is 'test solution – sample - test solution - sample'. If different equipment is used and /or more or less samples are worked up, modifications of this order may be necessary.

5.7 Calculations

Determination of concentration of the analytical target in the final solution

The concentrations of the analytes in control samples, fortified samples and treated samples were calculated using external standard procedures with multi level or single level calibration.

Single level calibration (one point calibration):

$$C_s = \frac{P_s}{P_R} \cdot C_R \cdot \frac{I_R}{T_4} \quad \left[\text{pg}/\mu\text{L} = \frac{\text{counts}}{\text{counts}} \cdot \text{pg}/\mu\text{L} \cdot \frac{\mu\text{L}}{\mu\text{L}} \right] \quad (1)$$

C_s Concentration in final sample solution V_{end} (identical with conc. in T_4)

(treated, untreated and recovery)

[$\text{pg}/\mu\text{L}$] = [ng/mL]

C_R Concentration in reference solution

[$\text{pg}/\mu\text{L}$] = [ng/mL]

P_s Peak area or peak height of the sample solution

[counts]

P_R Peak area or peak height of the reference solution

[counts]

T_4 Injection volume of the sample solution

[μL]

I_R Injection volume of the reference solutions

[μL]

Multi level calibration (calibration curve):

For the calibration peak areas (heights) of the standards were plotted versus the corresponding concentrations. An optimized calibration curve of the following form

$$f(C_s) = P = a + bC_s + cC_s^2 \quad (2)$$

is calculated, where $f(C_s)$ is the peak area (height), C_s the concentration of the analyte in the final sample extract and a , b , c are constants.

Determination of residues

Calculation of residues was carried out by a data handling software according to the following procedure

$$\text{Res} = \frac{C_s \cdot V_{\text{end}} \cdot f}{W} \quad \left[\mu\text{g/L} = \frac{(\text{ng/mL}) \cdot \text{mL} \cdot 1}{\text{mL}} \right] \quad (3)$$

$$f = \frac{V_1 \cdot V_2 \cdot V_n}{T_1 \cdot T_2 \cdot T_n} \quad \left[1 = \frac{\text{mL} \cdot \text{mL} \cdot \text{mL}}{\text{mL} \cdot \text{mL} \cdot \text{mL}} \right] \quad (4)$$

Res	Residue	[μg/L]
C_s	Concentration in final sample solution <i>V_{end}</i> (treated, untreated and recovery)	[ng/mL]
W	Sample weight	[mL]
f	Dilution factor	without dimension
V₁	Volume for primary extraction	[mL]
V₂	Volume after making up of aliquot <i>T₁</i>	[mL]
V_n	Volume after making up of aliquot <i>T_{n-1}</i> (<i>n</i> = 3, 4 and so on)	[mL]
V_{end}	Final sample solution (identical with <i>V₂</i> or <i>V₃</i> or <i>V_n</i> depending on the method)	[mL]
T₁	Aliquot of <i>V₁</i>	[mL]
T₂	Aliquot of <i>V₂</i>	[mL]
T_n	Aliquot of <i>V_n</i> (<i>n</i> = 3, 4 and so on)	[mL]

Determination of recovery rates

Calculation of recovery rates were carried out by a data handling software according to the following procedure

$$\text{Res}_{\text{d}} = \text{Res}_{(\text{Rec})} - \text{Res}_{(\text{Unt})} \quad \left[\frac{\mu\text{g}}{\text{L}} = \frac{\mu\text{g}}{\text{L}} - \frac{\mu\text{g}}{\text{L}} \right] \quad (5)$$

$$\text{Rec} = \frac{\text{Res}_{\text{d}}}{\text{Res}_{\text{f}}} \cdot 100 \quad \left[\% = \frac{\mu\text{g/L}}{\mu\text{g/L}} \cdot \% \right] \quad (6)$$

Res_(Rec)	Residue in the sample solution of the recovery test calculated with equation (3) and (4)	[μg/L]
Res_(Unt)	Residue in the sample solution of the corresponding untreated control sample calculated with equation (3) and (4)	[μg/L]
Rec	Recovery rate	[%]
Res_f	Concentration spiked for fortification	[μg/L]
Res_d	Concentration detected by analytical method	[μg/L]

6 Results

Calculation sheets and calibration curves for all laboratory analyses are given in Annex II and III with examples of chromatograms given in Annex VI.

6.1 Recoveries

Recovery experiments for method validation were conducted at 0.1 and 1.0 µg/L. At each level for drinking water or surface water a min. of 5 recovery values were determined. Recoveries are calculated with calibration curve according to the procedure described in section 5.8 with correction for any apparent residues.

The mean recoveries for AE F075736 in **drinking water** were found at 79 and 92 %, with an RSD of up to 9 %.

In **surface water** the mean recoveries for AE F075736 were found at 85 and 105 %, with an RSD of up to 10 %, the confirmation method gave mean recovery values of 92 and 106 %, with an RSD of up to 18 %.

In **surface water** the mean recoveries for AE F115008 were found at 92 and 103 %, with an RSD of up to 9 %, the confirmation method gave mean recovery values of 85 and 96 %, with an RSD of up to 20 %.

A summary of all recoveries for AE F075736 and AE F115008 are given in the following tables.

Summary table of recoveries:
Analyte: AE F075736

Matrix	Fortification level [µg/L]	Recovery [%]	Mean recovery [%]	RSD (a) [%]	n
Drinking water (Vittel)	0.10	82			
		75			
		87			
		83			
		70	79	9	5
	1.0	96			
		91			
		87			
		93			
		93	92	4	5
Surface water	0.10	90			
		78			
		91			
		92			
		74	85	10	5
	1.0	98			
		102			
		105			
		109			
		109	105	5	5
Surface water Confirmation method	0.10	74			
		110			
		92			
		106			
		108			
		90			
		67			
	1.0	35 (b)	92	18	7
		102			
		106			

The calculation was done with calibration function of peak areas.

a)
$$\text{RSD} = \frac{\text{S.D.}}{\text{Mean Recovery}} \times 100\%$$

$$\text{S.D.} = \sqrt{\frac{\sum (R_i - R_m)^2}{n - 1}}$$

R_i: recovery
 R_m: mean recovery
 n: number of recoveries

- b) Outliner Injection problems, HPLC system used to much solution for washing, so the injection volume was too small

Summary table of recoveries:
Analyte: AE F115008

Matrix	Fortification level [µg/L]	Recovery [%]	Mean recovery [%]	RSD (a) [%]	n
Surface water	0.10	108			
		92			
		94			
		106			
		113	103	9	5
	1.0	89			
		90			
		90			
		94			
		95	92	3	5
Surface water Confirmation method	0.10	75			
		78			
		74			
		114			
		138 (b)			
		98			
		73			
	1.0	36 (c)	85	20	6
		92			
		97			
		97			
		98	96	2	5

The calculation was done with calibration function of peak areas.

a) RSD = S.D. / Mean Recovery • 100 %

$$S.D. = \left[\frac{\sum (R_i - R_m)^2}{n - 1} \right]^{1/2}$$

R_i: recovery
 R_m: mean recovery
 n: number of recoveries

- b) Outliner Unexpected high value (results including this value: mean recovery = 93 %, RSD = 27 %, n = 7)
 c) Outliner Injection problems, HPLC system used to much solution for washing, so the injection volume was too small

6.2 Limit of quantification (LOQ)

Drinking water:

The lowest level at which metsulfuron-methyl has been quantified in this study is 0.1 µg/L. This level is therefore considered to be an appropriate limit of quantification for AE F075736 in drinking water.

The LOQ (0.1 µg/L) for AE F115008 in drinking water is given in the analytical method EM F 01/98 – 0 (ref. 1).

Surface water

The lowest level at which iodosulfuron-methyl-sodium and metsulfuron-methyl have been quantified in this study is 0.1 µg/L. This level is therefore considered to be an appropriate limit of quantification for AE F115008 and AE F075736 in surface water.

6.3 Blank values

Drinking water:

Analysis of control samples has shown that apparent residues of AE F075736 observed were n.d. (not detectable, < 0.3 x LOQ). This demonstrates that 0.1 µg/L is a feasible level for recognition of residues with reasonable certainty.

Surface water:

Analysis of control samples has shown that apparent residues of AE F115008 and AE F075736 observed were n.d. (not detectable, < 0.3 x LOQ). This demonstrates that 0.1 µg/L is a feasible level for recognition of residues with reasonable certainty.

6.4 Critical steps of the method

There are no critical steps of the method.

6.5 Time for analysis

From extraction of the samples to preparation of the final solutions for HPLC/UV determination, it is normally possible to analyse a batch of 14 samples in three day.

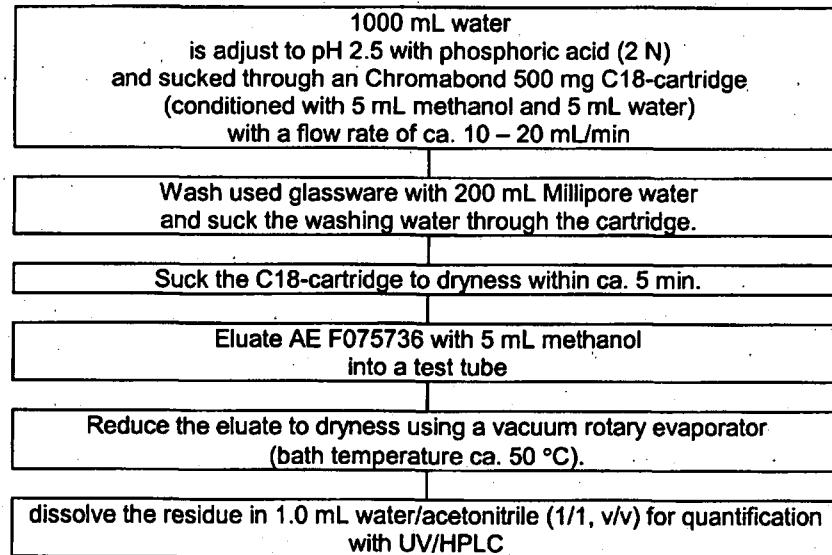
REFERENCES

No	Doc No	Report No	Author(s) Title Source and Date
1	C000710	EM F 01/98 - 0	Wrede A., 1998 Code AE F115008 Analytical method and validation for the determination of residues of AE F115008 and its metabolite AE F075736 in water using HPLC
2	A54684	CP 94/070	Sadowsky-Dunkmann I., Schmidt W., 1995 Substance, pure Code: HOE 115008 00 ZB97 0001 Solubility in organic solvents/vehicles according to Commission Directive 94/37/EEC (1994)

Annex I: Analytical method flow sheet**Metsulfuron-methyl in drinking water**

*Extraction AE F075736 and
C18-cartridge clean-up*

HPLC



Iodosulfuron-methyl-sodium and metsulfuron-methyl in surface water

Extraction
AE F115008 and
AE F075736 and
NH₂ / C18-cartridge
clean-up

1000 mL water
is adjust to pH 2.5 with phosphoric acid (2 N),
filtered through a glass microfibre filter and a cellulose nitrate filter (0.45 µm)
the sample is sucked through a Chromabond 500 mg NH₂ / 500 mg C18-cartridge
(conditioned with 5 mL methanol and 5 mL water)
with a flow rate of ca. 10 – 20 mL/min

Wash used glassware with 200 mL Millipore water
and suck the washing water through the cartridge.

Suck the NH₂ / C18-cartridge to dryness within ca. 5 min.

Wash the NH₂ / C18-cartridge with 10 mL methanol/water (30:70, v/v).

Eluate AE F115008 and AE F075736 with 15 mL methanol/water (60:40, v/v)

Reduce the eluate to dryness using a vacuum rotary evaporator
(bath temperature ca. 50 °C).

**Silicagel-cartridge
clean-up**

Dissolve the residue in 20 mL toluene (if necessary use an ultrasonic bath)
suck through a Silicagel-cartridge (conditioned with 5 mL toluene)

Discard the eluate, suck the Silicagel-cartridge to dryness

Wash the round-bottom flask with 30 mL toluene/methanol (95:5, v/v)
and elute AE F115008 and AE F075736 with this solution
Reduce the eluate to dryness using a vacuum rotary evaporator
(bath temperature ca. 40 °C).

HPLC

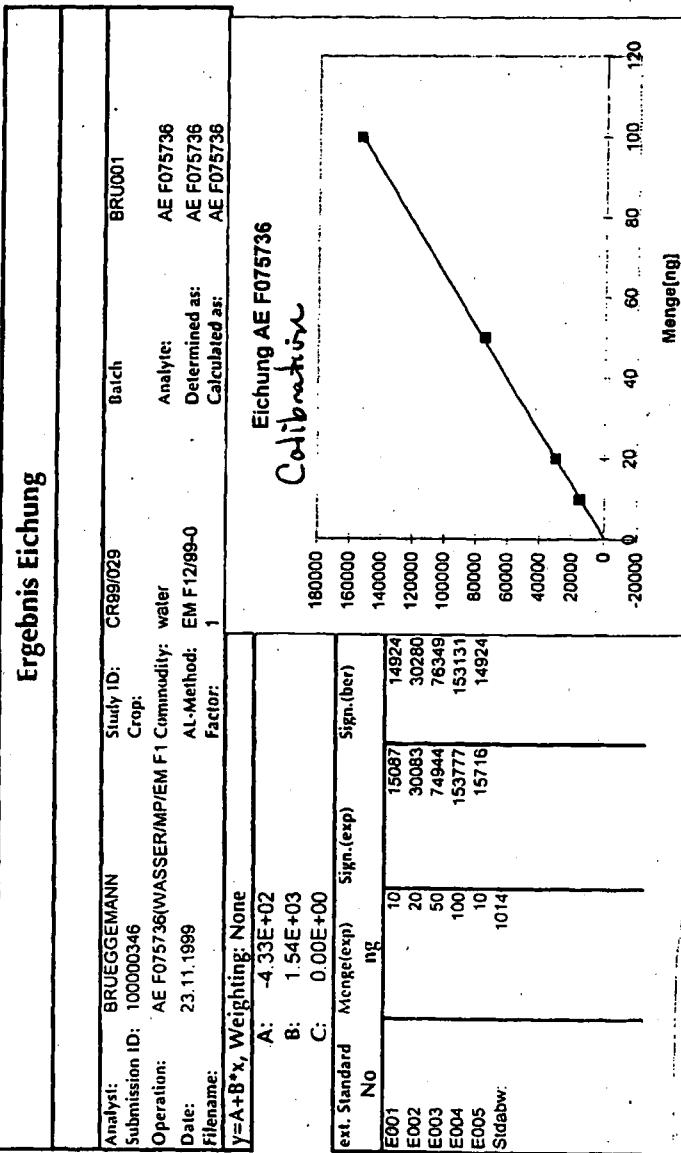
dissolve the residue in 1.0 mL water/acetonitrile (1/1, v/v) for quantification with
UV/HPLC

Annex II: Calibration curves

AE F075736

(drinking water)

Calibration Results



Calibration curves

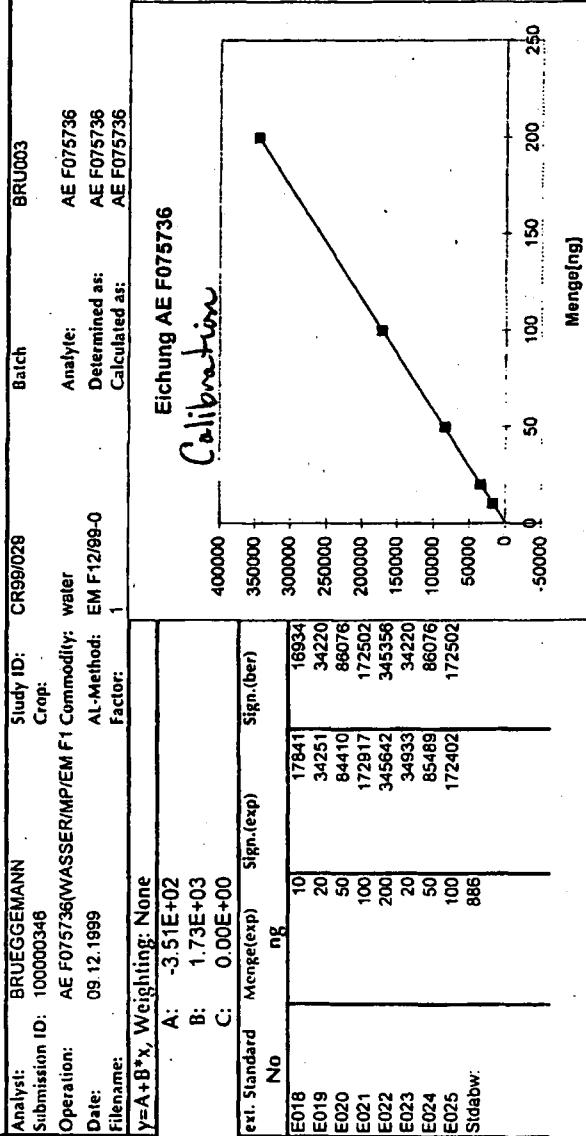
AE F075736

||

(drinking water)

Calibration Results

Ergebnis Eichung



Calibration curves

AE F075736

III

(surface water)

Calibration Results

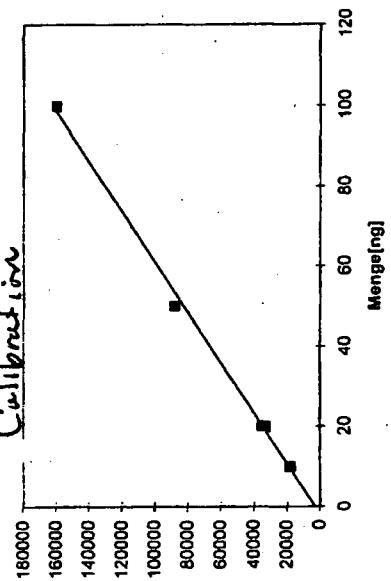
Ergebnis Eichung

Analyst:	BRÜGGEIMANN	Study ID:	CR99/029	Batch:	BRUQ07
Submission ID:	100000346	Crop:		Analyte:	AE F075736
Operation:	AE F075736(WASSERIMP/EM F1	Commodity:	water	Determined as:	AE F075736
Date:	24.01.2000	AI-Method:	EM F12/99-0	Calculated as:	AE F075736
Filename:		Factor:	1		

est. Standard No	Menge(exp)	Sign.(exp)	Sign.(ber)
E054	10	16536	18977
E055	20	32959	34881
E056	50	88301	82596
E057	100	159773	162120
E058	10	17758	18977
E059	20	35105	34881
Sidabw.	3297		60000

Eichung AE F075736

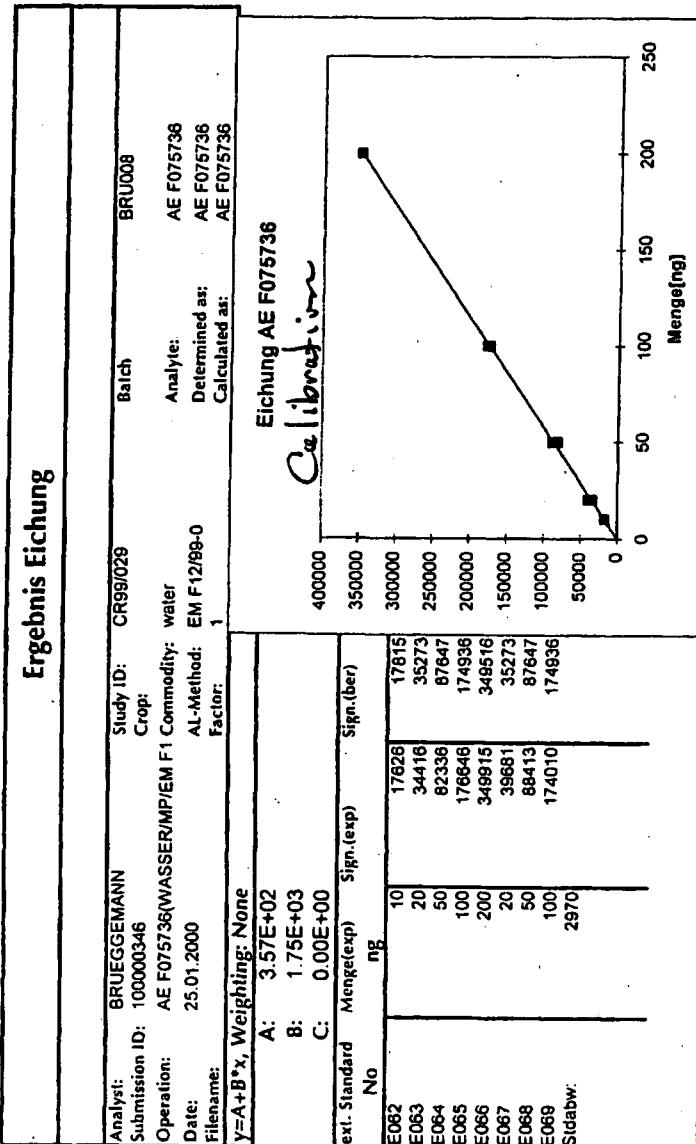
Calibration



Calibration curves

 AE F075736
 IV

(surface water)

Calibration Results


Calibration curves

AE F075736

V

(surface water)
(confirmation method)**Calibration Results****Ergebnis Eichung**

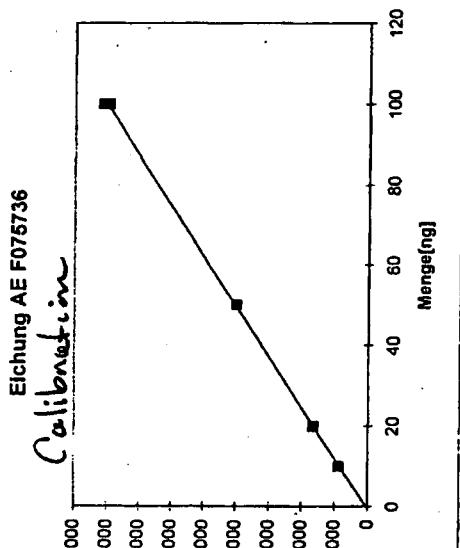
Analyst:	BRUEGGMANN	Study ID:	CR99/029	Batch:	BRU005
Submission ID:	100000346	Crop:		Analyte:	AE F075736
Operation:	AE F075736(WASSER/MP/EM F1	Commodity:	water	Determined as:	AE F075736
Date:	18.01.2000	AL-Method:	EM F12/99-0	Calculated as:	AE F075736
Filename:		Factor:	1		

y=A+B*x, Weighting: None

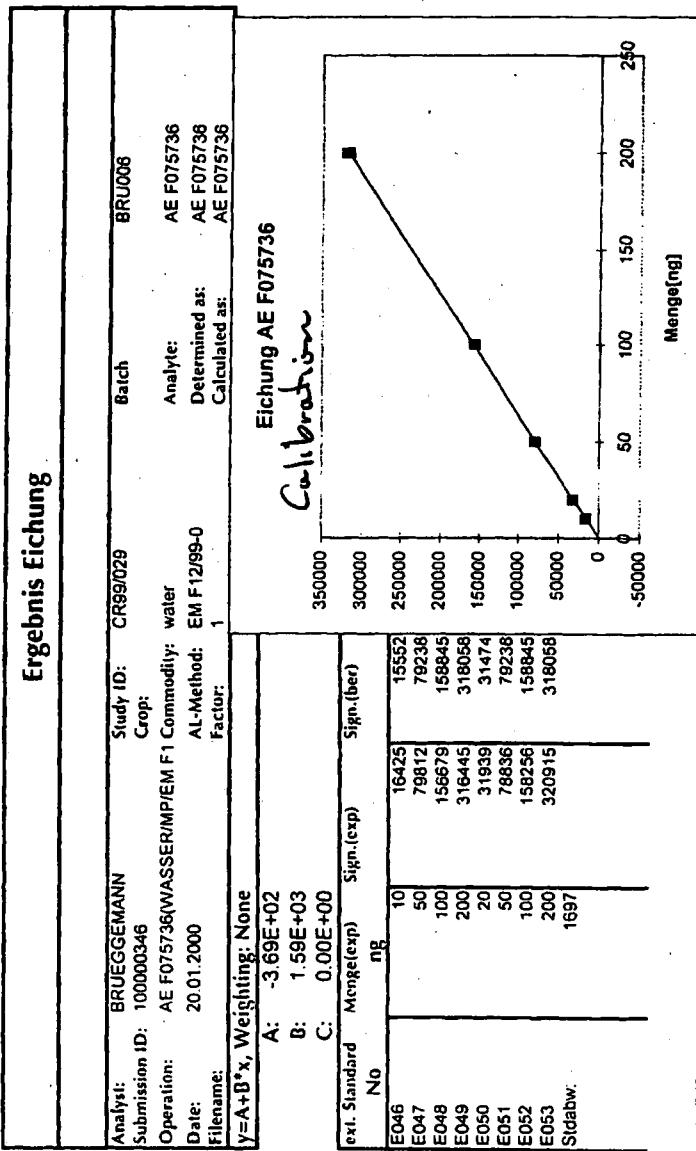
A: 1.11E+03

B: 1.60E+03

C: 0.00E+00



No.	Standard	Menge(exp)	Sign.(exp)	Sign.(ber)
E038		10	17428	17066
E039		20	32714	33028
E040		50	80124	80912
E041		100	159251	160718
E042		10	17698	17066
E043		20	33157	33028
E044		50	79847	80912
E045		100	163029	160718
Sidabw.		1303		

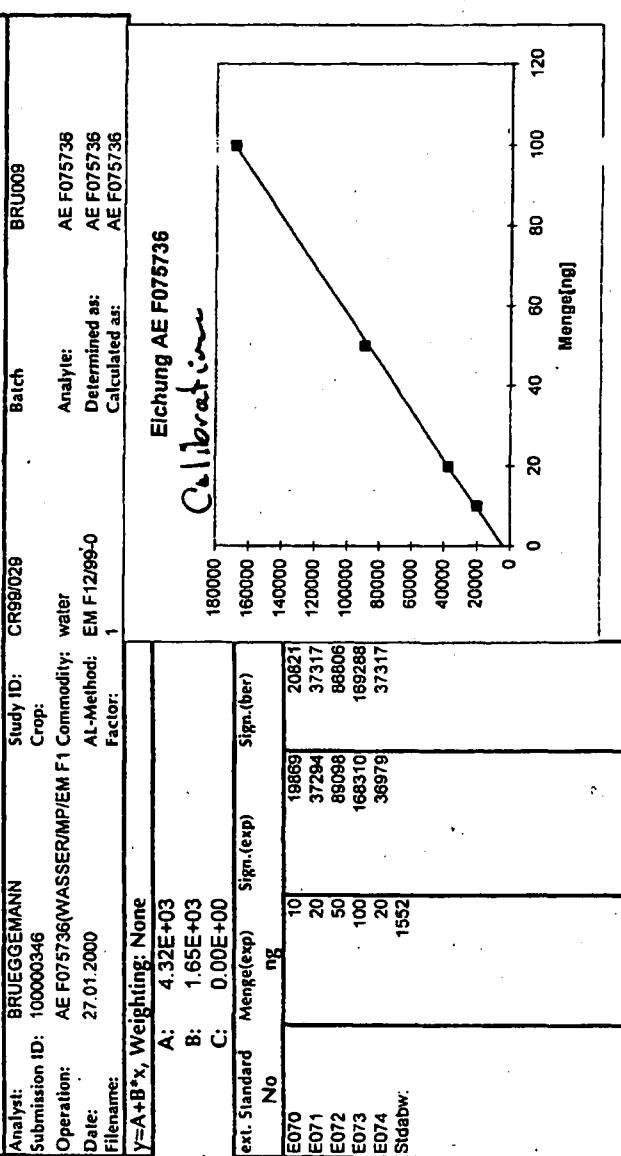
Calibration curves
AE F075736
VI
(surface water)
(confirmation method)
Calibration Results
Ergebnis Eichung


Calibration curves

AE F075736
VII(surface water)
(confirmation method)

Calibration Results

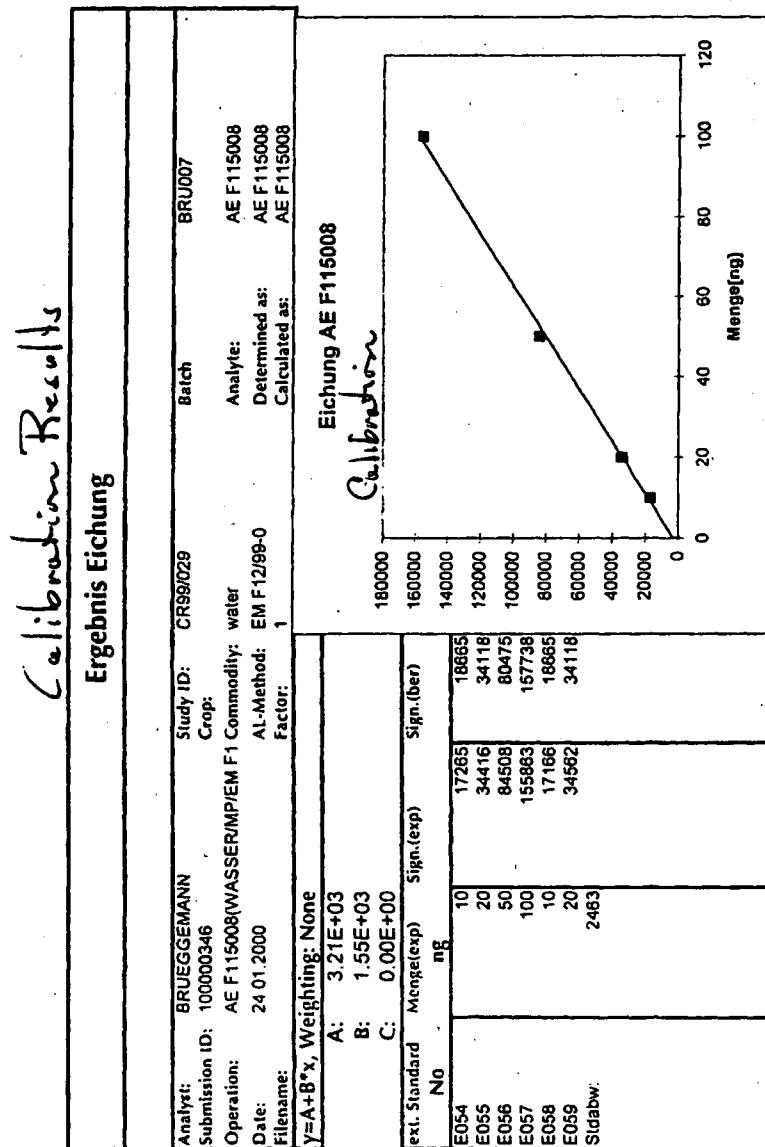
Ergebnis Eichung

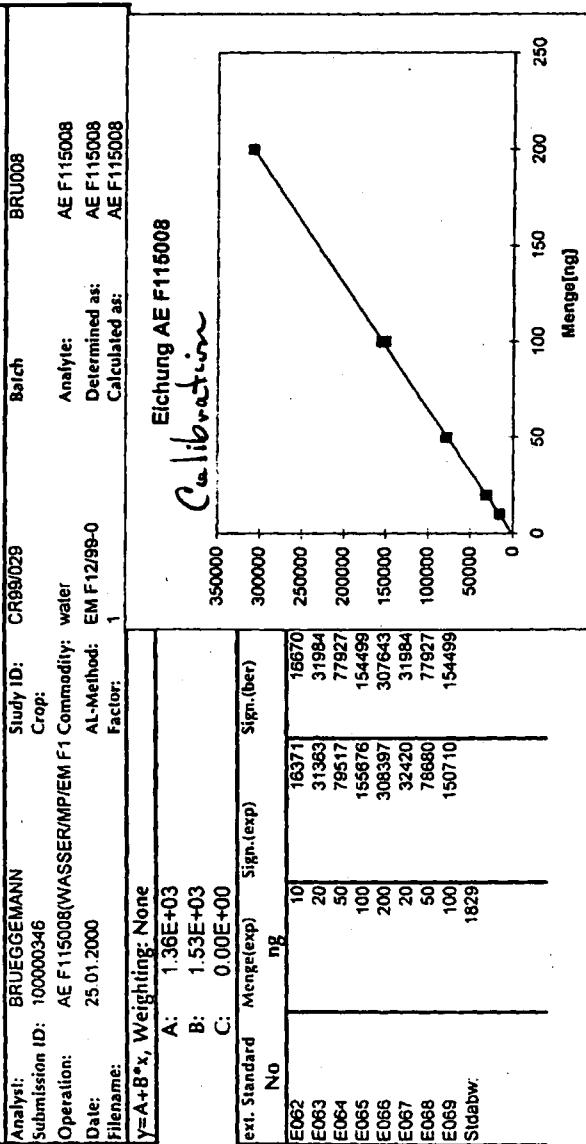


Calibration curves

AE F115008

(surface water)



Calibration curves
AE F115008
||
(surface water)
Calibration Results
Ergebnis Eichung


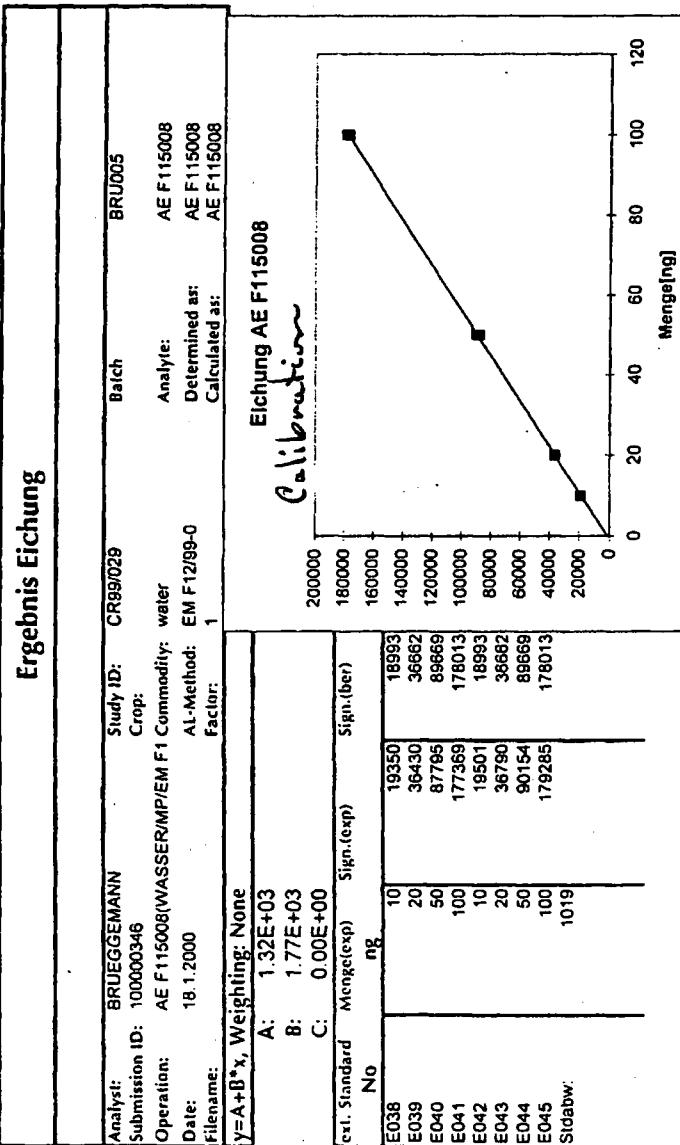
Calibration curves

AE F115008

III

(surface water)
(confirmation method)

Calibration Results



Calibration curves

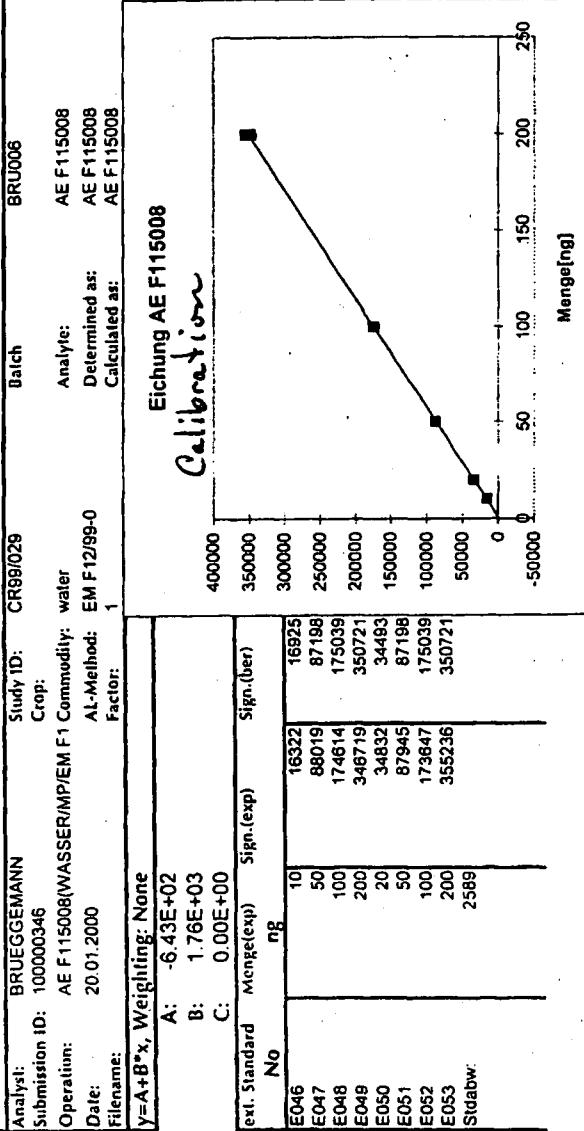
AE F115008

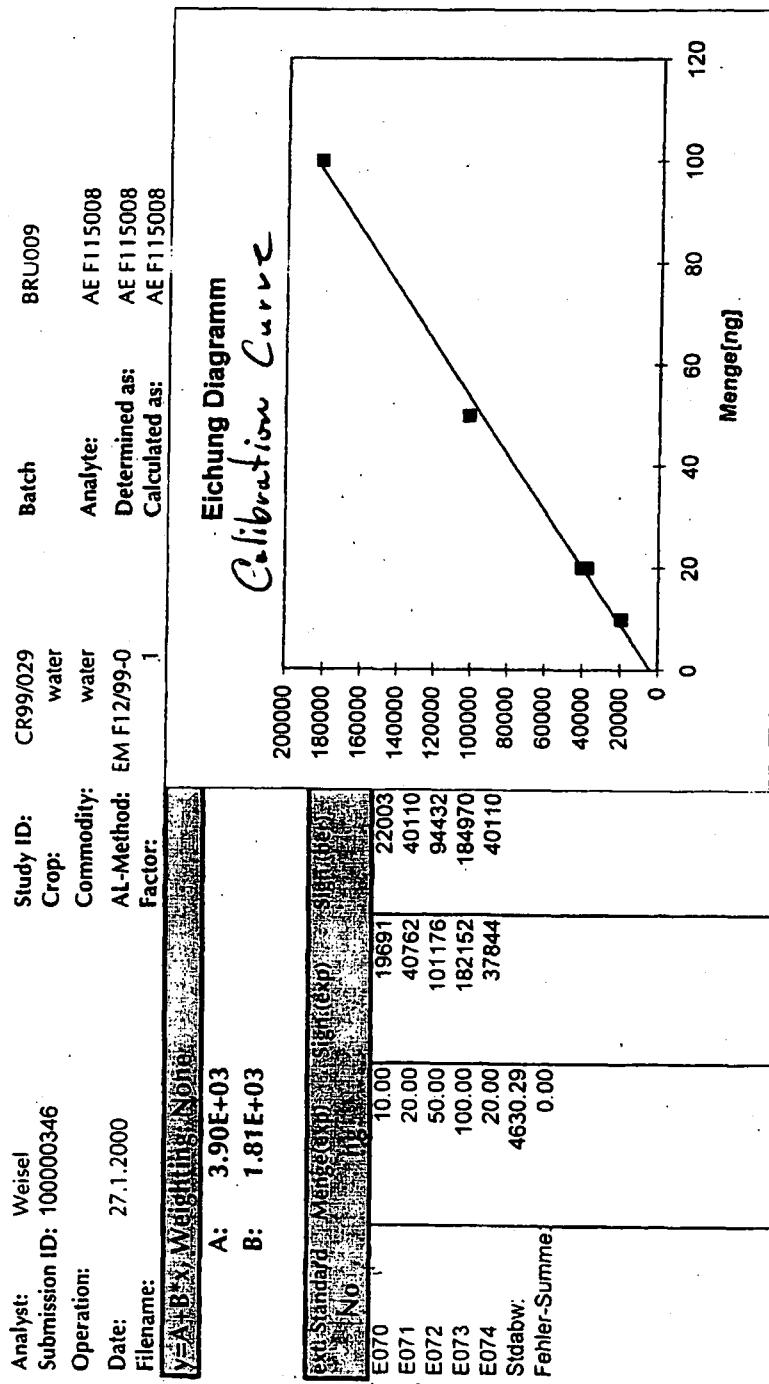
IV

(surface water)
(confirmation method)

Calibration Results

Ergebnis Eichung



Calibration curves
AE F115008
V
(surface water)
(confirmation method)


Annex III: Calculation sheets

AE F075736
Drinking water

0.1 µg/L

Calculation of Calibration Curve

Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc²Conc)

Sample_ID	Task-ID	Lab	W	V1	T1	V2	T2	V3	T3	V4	T4	Sample	File	A	B	C	counts	counts	counts	<	spiked	Reid	LQ	Sample	ug/l	%				
				Code	ml	Counts	Counts	Counts	Counts	ug/ml																				
200016106 BLI	300030073 U001	1	1	1	1	1	1	1	1	1	1	100	0	U001	-4.33E+02	1.54E+03	0.00E+00	n.d.	0.00E+00	0.021	n.d.	0.00E+00	0.021	n.d.	0.00E+00	0.021	n.d.	0.00E+00	0.021	n.d.
200016107 BLI	300030074 U002	1	1	1	1	1	1	1	1	1	1	100	2850	U002	-4.33E+02	1.54E+03	0.00E+00	0.021	0.00E+00	0.021	n.d.	0.00E+00	0.021	n.d.	0.00E+00	0.021	n.d.	0.00E+00	0.021	n.d.
200016130 AUF	300030068 R001	1	1	1	1	1	1	1	1	1	1	100	13841	R001	-4.33E+02	1.54E+03	0.00E+00	0.093	0.00E+00	0.093	MBV	0.1	0.1	62	0.1	0.1	62	0.1	0.1	62
200016131 AUF	300030071 R002	1	1	1	1	1	1	1	1	1	1	100	12673	R002	-4.33E+02	1.54E+03	0.00E+00	0.085	0.00E+00	0.085	MBV	0.1	0.1	75	0.1	0.1	75	0.1	0.1	75
200016132 AUF	300030069 R003	1	1	1	1	1	1	1	1	1	1	100	14513	R003	-4.33E+02	1.54E+03	0.00E+00	0.097	0.00E+00	0.097	MBV	0.1	0.1	87	0.1	0.1	87	0.1	0.1	87
200016133 AUF	300030072 R004	1	1	1	1	1	1	1	1	1	1	100	14017	R004	-4.33E+02	1.54E+03	0.00E+00	0.094	0.00E+00	0.094	MBV	0.1	0.1	83	0.1	0.1	83	0.1	0.1	83
200016134 AUF	300030070 R005	1	1	1	1	1	1	1	1	1	1	100	11920	R005	-4.33E+02	1.54E+03	0.00E+00	0.080	0.00E+00	0.080	MBV	0.1	0.1	70	0.1	0.1	70	0.1	0.1	70

Calculation sheets
AE F075736
 Drinking water

1.0 µg/L

 Calculation of Calibration Curve
 Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc*Conc)

Sample_ID	Task-ID	Lab	W	V1	T1	V2	T2	V3	T3	V4	T4	Sample	File	A	B	C	Resid.	< counts/ng	counts/ng	counts/ng	counts/ng	LQQ:	mean Blank:	0.01846	0.0100	0.03		
200016110 BLI	300030668 R005		1	1	1	1	1	1	1	1	1	100	2869	0.005	-3.51E+02	1.73E+03	0.00E+00	0.019	n.d.									
200016111 BLI	300030667 U006		1	1	1	1	1	1	1	1	1	100	2810	0.006	-3.51E+02	1.73E+03	0.00E+00	0.018	n.d.									
200016280 AUF	300030661 U009		1	1	1	1	1	1	1	1	1	100	168915	R009	-3.51E+02	1.73E+03	0.00E+00	0.98		200016110	1	96						
200016281 AUF	300030664 R010		1	1	1	1	1	1	1	1	1	100	159882	R010	-3.51E+02	1.73E+03	0.00E+00	0.93		200016111	1	91						
200016282 AUF	300030662 R011		1	1	1	1	1	1	1	1	1	100	153625	R011	-3.51E+02	1.73E+03	0.00E+00	0.89		200016110	1	87						
200016283 AUF	300030665 R012		1	1	1	1	1	1	1	1	1	100	163234	R012	-3.51E+02	1.73E+03	0.00E+00	0.95		200016111	1	93						
200016284 AUF	300030663 R013		1	1	1	1	1	1	1	1	1	100	162907	R013	-3.51E+02	1.73E+03	0.00E+00	0.94		200016110	1	93						

Calculation sheets

AE F075736
Surface water

0.1 µg/L

Calculation of Calibration Curve

Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc²)

Auswertung[Eichkurve: Counts=A+B*Conc+C*Conc*Conc]																
Analyst:	BRUEGGERMANN	Study ID:	CR960029	Batch:	BRU007	SEND	LOQ: 0.1								LOQ: 0.01	
							Analyte:				Determined as:				mean Blank:	STD:
Submission ID:	100000346	Crop:	water	Analyte:	AE F075736		mean Recov:	0	STD:	0	Determined as:	AE F075736		RSD:	0	
Operation:	AE F075736(WASSER)	Sample Mat.:	water	Determined as:	AE F075736		Calculated as:	AE F075736	STD:	0.46	Calculated as:	AE F075736		RSR:	10	
Date:	24.01.2000	AL-Method:	EMF 12/98-0	Factor:	1.00		mean LabID:	0	STD:	0	Factor:	1.00		Sample ID:	0	
Filename:	BR081331	Lab:	W	V1	T1	V2	T2	V3	T3	V4	T4	Sample File	A	B	Spiked	
Sample_ID	Type	Task-ID	Code	l	ml	ml	ml	ml	ml	ml	ml	Counts	counts	counts	Sample	
2000016122 BLI		300031929	U013	1	1	1	1	1	1	1	1	0	U013	3.07E+03	<	upfr %
2000016123 BLI		300031931	U014	1	1	1	1	1	1	1	1	0	U014	1.59E+03	<	upfr %
20000161951 AUF		300031919	R027	1	1	1	1	1	1	1	1	100	17441	1027	3.07E+03	0.1
3000031925 AUF		300031928	R028	1	1	1	1	1	1	1	1	100	15474	R028	3.07E+03	0.1
2000016911 AUF		300031921	R029	1	1	1	1	1	1	1	1	100	11604	R029	3.07E+03	0.1
2000016918 AUF		300031927	R030	1	1	1	1	1	1	1	1	100	17685	R030	3.07E+03	0.1
2000016919 AUF		300031923	R031	1	1	1	1	1	1	1	1	100	14837	R031	3.07E+03	0.1

Calculation sheets

AE F075736
Surface water

1.0 µg/L

Calculation of Calibration Curve

Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc²)

Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc*Conc)															LOG: 0.1										
Analyte:	BRÜGEGEMANN	Study ID:	CR980028	Batch:	BRD008	SEND	LOG: 0.1							LOG: 0.34 LOQ							0.03				
							Crop:	water	Analyte:	AE F075736	Determined as:	AE F075736	Calculated as:	AE F075736	mean Blank:	0	STD:	0	mean Recov:	105	STD:	4.83	R&D:	6	STD:
Submission ID:	100000348						AE F075736(WASSER)	Sample Mat:	water																
Operation:	AE F075736																								
Date:	25.01.2000						AI-Methode:	EM F1299-0																	
Filename:	BR008443						Factor:	1.00																	
Sample_ID	Type	Task-ID	Lab	W	V1	T1	V2	T2	V3	T3	V4	T4	Sample_File	A	B	C	Resid	<	spiked	Spiked	Sample	ug/ml	ug/ml		
				Code	l	ml	ml	ml	ml	ml	ml	ml	Counts	counts	counting	counts	ug/ml	LOQ	Sample	ug/ml	ug/ml	%			
200016124	BLI	300032117	U015	1	1	1	1	1	1	1	1	1	100	0	U015	3.57E+02	1.75E-03	0.00E+00	0.00E+00	0.00E+00	n.d.	n.d.			
200016125	BLI	300032119	U016	1	1	1	1	1	1	1	1	1	100	0	U016	3.57E+02	1.75E-03	0.00E+00	0.00E+00	0.00E+00	MBV	1			
2000161667	AUF	300032107	R032	1	1	1	1	1	1	1	1	1	100	171221	R032	3.57E+02	1.75E-03	0.00E+00	0.98	MBV	1	102			
200016568	AUF	300032113	R033	1	1	1	1	1	1	1	1	1	100	178003	R033	3.57E+02	1.75E-03	0.00E+00	1.0	MBV	1	105			
200016569	AUF	300032109	R034	1	1	1	1	1	1	1	1	1	100	185506	R034	3.57E+02	1.75E-03	0.00E+00	1.0	MBV	1	109			
200016870	AUF	300032115	R035	1	1	1	1	1	1	1	1	1	100	180084	R035	3.57E+02	1.75E-03	0.00E+00	1.1	MBV	1	109			
200016871	AUF	300032111	R036	1	1	1	1	1	1	1	1	1	100	191374	R036	3.57E+02	1.75E-03	0.00E+00	1.1	MBV	1	109			
200016871	AUF	300032111	R036	1	1	1	1	1	1	1	1	1	100	191374	R036	3.57E+02	1.75E-03	0.00E+00	1.1	MBV	1	109			

Calculation sheets

AE F075736

Surface water (confirmation method)

0.1 µg/L

Calibration of Calibration Curve

Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc^2)

Sample_ID	Type	Task-ID	Lab	W	V1	T1	V2	T2	V3	T3	V4	T4	Sample	File	A	B	C	Reald	<	spiked	Spiked	Rec
					Code	ml	Counts	Counts	Counts	LOQ	Sample	ug/L	%									
200016118 BLI	300031470 U009	1	1	1	1	1	1	1	1	1	1	1	100	0	0.008	1.11E+03	1.60E+03	0.00E+00	0.000 n.d.	0.009 n.d.	0.00E+00	0.009
200016119 BLI	300031472 U010	1	1	1	1	1	1	1	1	1	1	1	100	2479	U010	1.11E+03	1.60E+03	0.00E+00	0.009	0.009 n.d.	0.00E+00	0.009
200016484 AUF	300031460 R017	1	1	1	1	1	1	1	1	1	1	1	100	15586	R017	1.11E+03	1.60E+03	0.00E+00	0.078	MBV	0.1	74
200016485 AUF	300031466 R018	1	1	1	1	1	1	1	1	1	1	1	100	19301	R018	1.11E+03	1.60E+03	0.00E+00	0.11	MBV	0.1	110
200016486 AUF	300031462 R019	1	1	1	1	1	1	1	1	1	1	1	100	18461	R019	1.11E+03	1.60E+03	0.00E+00	0.096	MBV	0.1	92
200016487 AUF	300031464 R020	1	1	1	1	1	1	1	1	1	1	1	100	18728	R020	1.11E+03	1.60E+03	0.00E+00	0.11	MBV	0.1	106
200016488 AUF	300031468 R021	1	1	1	1	1	1	1	1	1	1	1	100	18972	R021	1.11E+03	1.60E+03	0.00E+00	0.11	MBV	0.1	108

Calculation sheets

AE F075736

Surface water (confirmation method)

1.0 µg/L

 Calibration of Calibration Curve
 Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc*Conc)

Sample_ID	Task_ID	Lab	W	V1	T1	V2	T2	V3	T3	V4	T4	Counts	File	A	B	C	Resid	<	spiked	Spiked	Rec	ug/l	ug/l	%
200016120 BLI	300031620 U011	1	1	1	1	1	1	1	1	1	1	100	2371	U011	-3.69E+02	1.59E+03	0.00E+00	0.0017	nd	0.00E+00	0.000	0.00E+00	0.00E+00	0.00E+00
200016121 BLI	300031622 U012	1	1	1	1	1	1	1	1	1	1	100	0	U012	-3.69E+02	1.59E+03	0.00E+00	0.000	nd	0.00E+00	0.000	0.00E+00	0.00E+00	0.00E+00
200016128 AUF	300031610 R022	1	1	1	1	1	1	1	1	1	1	100	163250	R022	-3.69E+02	1.59E+03	0.00E+00	1.0	MBV	1	102	MBV	1	106
200016129 AUF	300031616 R023	1	1	1	1	1	1	1	1	1	1	100	170449	R023	-3.69E+02	1.59E+03	0.00E+00	1.1	MBV	1	106	MBV	1	106
200016130 AUF	300031612 R024	1	1	1	1	1	1	1	1	1	1	100	169455	R024	-3.69E+02	1.59E+03	0.00E+00	1.1	MBV	1	106	MBV	1	106
200016131 AUF	300031618 R025	1	1	1	1	1	1	1	1	1	1	100	172371	R025	-3.69E+02	1.59E+03	0.00E+00	1.1	MBV	1	108	MBV	1	108
200016132 AUF	300031614 R026	1	1	1	1	1	1	1	1	1	1	100	174106	R026	-3.69E+02	1.59E+03	0.00E+00	1.1	MBV	1	109	MBV	1	109

Calculation sheets

AE F075736

Surface water (confirmation method)

0.1 μg/L

Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc²)

Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc)

Calculation sheets

AE F115008
Surface water

0.1 µg/L

Calculation of Calibration Curve

Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc²)

Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc^2Conc)																
Analyst:	BRUEGEMANN	Study ID:	CR860239	Batch:	BR0007			SEND	LOG: 0.1			0.5%LOQ			0.03	
					Crop:	water	Analyte:		mean Blank:	0	STD:	0	mean Recov:	103	STD:	9.5
Substation ID:	100000346	Sample ID/N:	AE F115008	Determined as:	AE F115008	Calculated as:	AE F115008	mean Leab:	0	STD:	0	mean Leab:	0	STD:	0	MBV
Operation:	AE F115008	AL-Method:	EM F1298-0	Factor:	1.00	Code:		C	Resid	<	spiked	spiked	Rec	Leab/		
Date:	24.01.2000	Filename:	BR0007	Task-ID	Lab	W	V1	T1	V2	T2	V3	T3	V4	T4		
Sample ID	Type	Code	1	m	m	m	m	m	m	m	m	m	m	m		
300031928	BLT	U013	1	1	1	1	1	1	1	1	1	1	1	1		
300031930	BLI	U014	1	1	1	1	1	1	1	1	1	1	1	1		
300031919	BLB	U027	1	1	1	1	1	1	1	1	1	1	1	1		
300031924	RO28	1	1	1	1	1	1	1	1	1	1	1	1	1		
300031920	RO29	1	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U020	1	1	1	1	1	1	1	1	1	1	1	1		
300031919	AUF	U021	1	1	1	1	1	1	1	1	1	1	1	1		
300031926	AUF	U030	1	1	1	1	1	1	1	1	1	1	1	1		
300031927	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031919	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031920	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031920	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031921	AUF	U031	1	1	1	1	1	1	1	1	1	1	1	1		
300031922	AUF	U														

Calculation sheets

AE F115008
Surface water

1.0 µg/L

Calibration of Calibration Curve

Calculation sheets

AE F115008

Surface water (confirmation method)

0.1 µg/L

Calculation of Calibration Curve

Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc*Conc)

Sample_ID	Type	Task-ID	Lab	W	V1	T1	V2	T2	V3	T3	V4	T4	Sample	File	A	B	C	Resid.	<	Spiked	Spiked	Rec.
				ml	Counts	Counts	counts	counts	counts	ug/ml	ug/ml	ug/ml	ug/ml	%								
200016118	BLI	300031469	U009	1	1	1	1	1	1	1	1	1	100	0	0.009	1.32E+03	1.77E+03	0.00E+00	0.000	0.000	n.d.	
200016119	BLI	300031471	U010	1	1	1	1	1	1	1	1	1	100	0	0.010	1.32E+03	1.77E+03	0.00E+00	0.000	0.000	n.d.	
200016484	AUF	300031459	R017	1	1	1	1	1	1	1	1	1	100	14508	R017	1.32E+03	1.77E+03	0.00E+00	0.075	0.075	MBV	0.1
200016485	AUF	300031465	R018	1	1	1	1	1	1	1	1	1	100	15150	R018	1.32E+03	1.77E+03	0.00E+00	0.078	0.078	MBV	0.1
200016486	AUF	300031461	R019	1	1	1	1	1	1	1	1	1	100	14341	R019	1.32E+03	1.77E+03	0.00E+00	0.074	0.074	MBV	0.1
200016487	AUF	300031463	R020	1	1	1	1	1	1	1	1	1	100	21408	R020	1.32E+03	1.77E+03	0.00E+00	0.11	0.11	MBV	0.1
200016488	AUF	300031467	R021	1	1	1	1	1	1	1	1	1	100	25649	R021	1.32E+03	1.77E+03	0.00E+00	0.14	0.14	MBV	0.1

Calculation sheets
AE F115008

Surface water (confirmation method)

1.0 µg/L
Calculation of Calibration Curve.
Auswertung(Eichkurve: Counts=A+B*Conc+C*Conc)

Analyst:	BRUEGEMANN	Study ID:	CR9a029	Batch:	BR0008 SEND	LOQ:	0.3700																			
Submission ID:	10000346	Crop:	water	Analyte:	AE F115008	mean Blank:	0																			
Operation:	AE F115008/WASSER/N Sample Mat.:	water	Determined as:	AE F115008	mean Recov.:	96																				
Date:	2001.2000	AL-Method:	EM F 298-0	Calculated as:	AE F115008	STD:	0																			
Filename:	BR08430	Factor:	1.00			STD:	2.46																			
Sample_ID	Type	Task-ID	Lab	W	V1	T1	V2	T2	V3	T3	V4	T4	Counts	File	A	B	C	Resid	<	spiked	Spiked	Rec	Sample	Sign	%	
200016120	BLI	300031619	U011	1	1	1	1	1	1	1	1	1	100	U011	-6.43E-02	1.76E-03	0.00E+00	0.00E+00	n.d.	0.00E+00	0.00E+00	0.00E+00	0.00E+00	nd		
200016121	BLI	300031621	U012	1	1	1	1	1	1	1	1	1	100	U012	-6.43E-02	1.76E-03	0.00E+00	0.00E+00	n.d.	0.00E+00	0.00E+00	0.00E+00	0.00E+00	nd		
200016828	AUF	300031609	R022	1	1	1	1	1	1	1	1	1	100	160784	R022	-6.43E-02	1.76E-03	0.00E+00	0.92	MBV	1	92				
200016829	AUF	300031615	R023	1	1	1	1	1	1	1	1	1	100	169113	R023	-6.43E-02	1.76E-03	0.00E+00	0.97	MBV	1	97				
200016830	AUF	300031611	R024	1	1	1	1	1	1	1	1	1	100	168966	R024	-6.43E-02	1.76E-03	0.00E+00	0.97	MBV	1	97				
200016831	AUF	300031617	R025	1	1	1	1	1	1	1	1	1	100	168945	R025	-6.43E-02	1.76E-03	0.00E+00	0.97	MBV	1	97				
200016832	AUF	300031613	R026	1	1	1	1	1	1	1	1	1	100	172108	R026	-6.43E-02	1.76E-03	0.00E+00	0.98	MBV	1	98				

Calculation sheets**AE F115008**

Surface water (confirmation method)

0.1 µg/L

Analyst:	Weisel		Study ID:		CR990029		Batch:		BRU009		LOD:		0.1		0.3*LOQ		0.03							
	Submission ID:	100000346	Crop:	water	Commodity:	water	Analyte:	AE F115008	Determined as:	AE F115008	Calculated as:	AE F115008 <th>mean Blank:</th> <td>0</td> <th>STD:</th> <td>0</td> <th>mean Recov.:</th> <td>69</td> <th>STD:</th> <td>31.23</td> <th>mean Lstab:</th> <td>0</td> <th>STD:</th> <td>46</td>	mean Blank:	0	STD:	0	mean Recov.:	69	STD:	31.23	mean Lstab:	0	STD:	46
Date:	27.1.2000	AL-Method:	EM F12/99-0	Factor:	1.00	W	V1	T1	V2	T2	V3	T3	V4	T4	Sample	Chainell/ Sequenz.	A	B	C	Residue mg/kg	Spiked mg/kg	Rec. %		
Lab	Spiked	Code	W	V1	T1	V2	T2	V3	T3	V4	T4	ml	ml	ml	ml	Counts	Chainell/ Sequenz.	A	B	C	Residue mg/kg	Spiked mg/kg	Rec. %	
	Code	Code	g	ml	ml	ml	ml	ml	ml	ml	ml	ml	ml	ml	ml	ml	ml	ml	ml	LOD	0.100	0.03		
U017				1	1	1	1	1	1	1	1	1	1	1	1	0	U017	3895.481	1810.746	0	0.000	n.d.		
R037	U017			1	1	1	1	1	1	1	1	1	1	1	1	1	21588	R037	3895.481	1810.746	0	0.098	-	
R038	U017			1	1	1	1	1	1	1	1	1	1	1	1	1	17720	R038	3895.481	1810.746	0	0.073	-	
R039	U017			1	1	1	1	1	1	1	1	1	1	1	1	1	10357	R039	3895.481	1810.746	0	0.036	-	

Annex IV: Sample history and recovery efficiencies

Analyte: AE F075736

Matrix	Lab code	Start of sample work-up	Date of data acquisition	Fortification level [$\mu\text{g/L}$]	Recovery [%]	Mean recovery [%]	RSD (a) [%]	n
Drinking water (Vittel)	R001	22.11.1999	23.11.1999	0.10	82			
	R002	22.11.1999	23.11.1999	0.10	75			
	R003	22.11.1999	23.11.1999	0.10	87			
	R004	22.11.1999	23.11.1999	0.10	83			
	R005	22.11.1999	23.11.1999	0.10	70	79	9	5
	R009	06.12.1999	08.12.1999	1.0	96			
	R010	06.12.1999	08.12.1999	1.0	91			
	R011	06.12.1999	08.12.1999	1.0	87			
	R012	06.12.1999	08.12.1999	1.0	93			
	R013	06.12.1999	08.12.1999	1.0	93	92	4	5

The calculation was done with calibration function of peak areas.
 a) RSD = S.D. / Mean Recovery • 100 %

$$\text{S.D.} = \sqrt{\frac{\sum (R_i - R_m)^2}{n-1}}$$

R_i: recovery
 R_m: mean recovery
 n: number of recoveries

Analyte: AE F115008

Matrix	Lab code	Start of sample work-up	Date of data acquisition	Fortification level [µg/L]	Recovery [%]	Mean recovery [%]	RSD (a) [%]	n
Surface water	R027	20.01.2000	21.01.2000	0.1	108			
	R028	20.01.2000	21.01.2000	0.1	92			
	R029	20.01.2000	21.01.2000	0.1	94			
	R030	20.01.2000	22.01.2000	0.1	106			
	R031	20.01.2000	22.01.2000	0.1	113	103	9	5
	R032	24.01.2000	24.01.2000	1.0	89			
	R033	24.01.2000	25.01.2000	1.0	90			
	R034	24.01.2000	25.01.2000	1.0	90			
	R035	24.01.2000	25.01.2000	1.0	94			
	R036	24.01.2000	25.01.2000	1.0	95	92	3	5

The calculation was done with calibration function of peak areas.
 a) RSD = S.D. / Mean Recovery • 100 %

$$S.D. = \sqrt{\frac{\sum (R_i - R_m)^2}{n - 1}}$$

R_i: recovery
 R_m: mean recovery
 n: number of recoveries

Analyte: AE F075736

Matrix	Lab code	Start of sample work-up	Date of data acquisition	Fortification level [µg/L]	Recovery [%]	Mean recovery [%]	RSD (a) [%]	n
Surface water	R027	20.01.2000	21.01.2000	0.1	90			
	R028	20.01.2000	21.01.2000	0.1	78			
	R029	20.01.2000	21.01.2000	0.1	91			
	R030	20.01.2000	22.01.2000	0.1	92			
	R031	20.01.2000	22.01.2000	0.1	74			
	R032	24.01.2000	24.01.2000	1.0	98			
	R033	24.01.2000	25.01.2000	1.0	102			
	R034	24.01.2000	25.01.2000	1.0	105			
	R035	24.01.2000	25.01.2000	1.0	109			
	R036	24.01.2000	25.01.2000	1.0	109			
					105		5	5

The calculation was done with calibration function of peak areas.
 a) RSD = S.D. / Mean Recovery • 100 %

$$S.D. = \sqrt{\frac{\sum (R_i - R_m)^2}{n-1}}$$

R_i: recovery
 R_m: mean recovery
 n: number of recoveries

Analyte: AE F115008

Matrix	Lab code	Start of sample work-up	Date of data acquisition	Fortification level [µg/L]	Recovery [%]	Mean recovery [%]	RSD (a) [%]	n
Surface water	R017	17.01.2000	17.01.2000	0.1	75			
Confirmation method	R018	17.01.2000	17.01.2000	0.1	78			
	R019	17.01.2000	18.01.2000	0.1	74			
	R020	17.01.2000	18.01.2000	0.1	114			
	R021	17.01.2000	18.01.2000	0.1	138 (b)			
	R037	25.01.2000	25.01.2000	0.1	98			
	R038	25.01.2000	25.01.2000	0.1	73			
	R039	25.01.2000	25.01.2000	0.1	36 (c)			
					85	20	6	
	R022	19.01.2000	19.01.2000	1.0	92			
	R023	19.01.2000	19.01.2000	1.0	97			
	R024	19.01.2000	20.01.2000	1.0	97			
	R025	19.01.2000	20.01.2000	1.0	97			
	R026	19.01.2000	20.01.2000	1.0	98	96	2	5

The calculation was done with calibration function of peak areas.
 a) RSD = S.D. / Mean Recovery • 100 %

$$S.D. = \left[\frac{\sum (R_i - R_m)^2}{n - 1} \right]^{1/2}$$

R_i: recovery
 R_m: mean recovery
 n: number of recoveries

- b) Outliner
 c) Outliner
- Unexpected high value (results including this value: mean recovery = 93 %, RSD = 27 %, n = 7)
 Injection problems, HPLC system used to much solution for washing, so the injection volume was too small

Analyte: AE F075736

Matrix	Lab code	Start of sample work-up	Date of data acquisition	Fortification level [µg/L]	Recovery [%]	Mean recovery [%]	RSD (a) [%]	n
Surface water	R017	17.01.2000	17.01.2000	0.1	74			
Confirmation method	R018	17.01.2000	17.01.2000	0.1	110			
	R019	17.01.2000	18.01.2000	0.1	92			
	R020	17.01.2000	18.01.2000	0.1	106			
	R021	17.01.2000	18.01.2000	0.1	108			
	R037	25.01.2000	25.01.2000	0.1	90			
	R038	25.01.2000	25.01.2000	0.1	67			
	R039	25.01.2000	25.01.2000	0.1	35 (b)			
					92	18	7	
	R022	19.01.2000	19.01.2000	1.0	102			
	R023	19.01.2000	19.01.2000	1.0	106			
	R024	19.01.2000	20.01.2000	1.0	106			
	R025	19.01.2000	20.01.2000	1.0	108			
	R026	19.01.2000	20.01.2000	1.0	109			
					106	3	5	

The calculation was done with calibration function of peak areas.
 a) RSD = S.D. / Mean Recovery • 100 %

$$S.D. = \sqrt{\frac{\sum (R_i - R_m)^2}{n-1}}$$

R_i: recovery
 R_m: mean recovery
 n: number of recoveries

b) Outliner Injection problems, HPLC system used to much solution for washing, so the injection volume was too small

Annex V: Sample history and apparent residue levels
Analyte: AE F075736

Matrix	Lab code	Start of sample work-up	Date of data acquisition	Apparent residue level AE F075736 [µg/L]	Mean apparent residue level AE F075736 [µg/L]
Drinking water (Vittel)	U001	22.11.1999	22.11.1999	0.000	n.d.
	U002	22.11.1999	22.11.1999	0.021	
	U005	06.12.1999	08.12.1999	0.019	
	U006	06.12.1999	08.12.1999	0.018	
Surface water	U013	20.01.2000	21.01.2000	0.000	n.d.
	U014	20.01.2000	21.01.2000	0.000	
	U015	24.01.2000	24.01.2000	0.000	
	U016	24.01.2000	24.01.2000	0.000	
Surface water Confirmation method	U009	17.01.2000	17.01.2000	0.000	n.d.
	U010	17.01.2000	17.01.2000	0.009	
	U011	19.01.2000	19.01.2000	0.017	
	U012	19.01.2000	19.01.2000	0.000	
	U017	25.01.2000	25.01.2000	0.000	

Analyte: AE F115008

Matrix	Lab code	Start of sample work-up	Date of data acquisition	Apparent residue level AE F115008 [µg/L]	Mean apparent residue level AE F115008 [µg/L]
Surface water	U013	20.01.2000	21.01.2000	0.000	n.d.
	U014	20.01.2000	21.01.2000	0.000	
	U015	24.01.2000	24.01.2000	0.000	
	U016	24.01.2000	24.01.2000	0.000	
Surface water Confirmation method	U009	17.01.2000	17.01.2000	0.000	n.d.
	U010	17.01.2000	17.01.2000	0.000	
	U011	19.01.2000	19.01.2000	0.000	
	U012	19.01.2000	19.01.2000	0.000	
	U017	25.01.2000	25.01.2000	0.000	

Annex VI: Typical chromatograms

The following chromatograms are given as examples of characteristic recorder plots.

The retention time of the target signal varied to a certain extent as the chromatographic system had to be fitted to the particular situation.

The computer plot offers the possibility of presenting only the important cut of the chromatogram with additional stretching of time or height. The electronic plots show the peak height in counts plotted vs. the retention time in minutes.

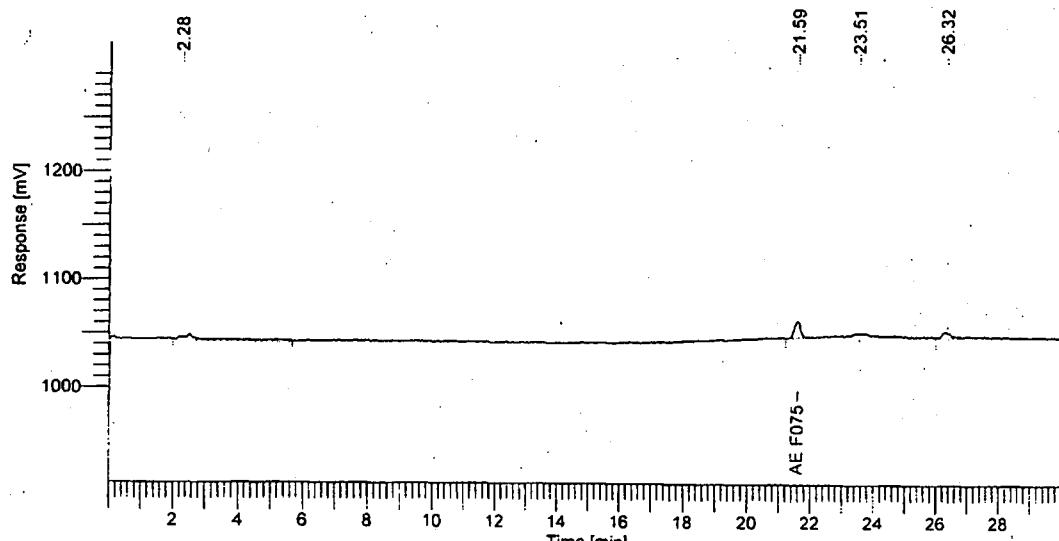
Sample material	Lab code	Description	Page
-	E001	Test 0.10 µg AE F075736/mL	61
	E002	Test 0.20 µg AE F075736/mL	62
	E003	Test 0.50 µg AE F075736/mL	63
	E004	Test 1.0 µg AE F075736/mL	64
Drinking water (Vittel)	U001	Control —	65
	R001	Recovery 0.10 µg AE F075736/L	66
	R002	Recovery 0.10 µg AE F075736/L	67
	U005	Control —	68
	R009	Recovery 1.0 µg AE F075736/L	69
	R010	Recovery 1.0 µg AE F075736/L	70
-	E054	Test 0.10 µg (AE F115008 + AE F075736)/mL	71
	E055	Test 0.20 µg (AE F115008 + AE F075736)/mL	72
	E056	Test 0.50 µg (AE F115008 + AE F075736)/mL	73
	E057	Test 1.0 µg (AE F115008 + AE F075736)/mL	74
Surface water	U013	Control —	75
	R027	Recovery 0.10 µg (AE F115008 + AE F075736)/L	76
	R028	Recovery 0.10 µg (AE F115008 + AE F075736)/L	77
	U015	Control —	78
	R032	Recovery 1.0 µg (AE F115008 + AE F075736)/L	79
	R033	Recovery 1.0 µg (AE F115008 + AE F075736)/L	80
(Confirmation method)	E038	Test 0.10 µg (AE F115008 + AE F075736)/mL	81
	E041	Test 1.0 µg (AE F115008 + AE F075736)/mL	82
Surface water (Confirmation method)	U009	Control —	83
	R017	Recovery 0.10 µg (AE F115008 + AE F075736)/L	84
	R018	Recovery 0.10 µg (AE F115008 + AE F075736)/L	85

Lab code Description
E001 Test 0.10 µg AE F075736/mL

Page 1 of 1

Software Version	: 6.1.0.2:G07	Date	: 22.11.1999 13:58:33
Operator	: Brueggemann	Sample Name	: E001
Sample Number	: 1	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s	Area Reject	: 0.000000
Volume Injected	: 100.000000 µL	Dilution Factor	: 1.00
Sample Amount	: 1.0000	Cycle	: 1
Data Acquisition Time	: 22.11.1999 13:08:46		

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 Sequence File : \\adeft003\Turbochrom Daten\Brueggemann\Sequenzen\CR99029\bru001.seq
 Sample Notes:
 10 ng AE F075736 in 100 µL



Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
2	AE F075736	21.589	211310	15087

Lab code Description

E002 Test 0.20 µg AE F075736/mL

Software Version	: 6.1.0.2:G07	Date	: 22.11.1999 15:39:02
Operator	: Brueggemann	Sample Name	: E002
Sample Number	: 3	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s	Area Reject	: 0.000000
Volume Injected	: 100.000000 µL	Dilution Factor	: 1.00
Sample Amount	: 1.0000	Cycle	: 3
Data Acquisition Time	: 22.11.1999 15:04:01		

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Result File : \\Adeft003\turbochrom daten\Brueggemann\Chromatogramme\CR99029\bru001\E002.rst

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Proc Method : \\Adeft003\Turbochrom Daten\brueggemann\methoden\75736emf1299 from \\Adeft003\turbochrom

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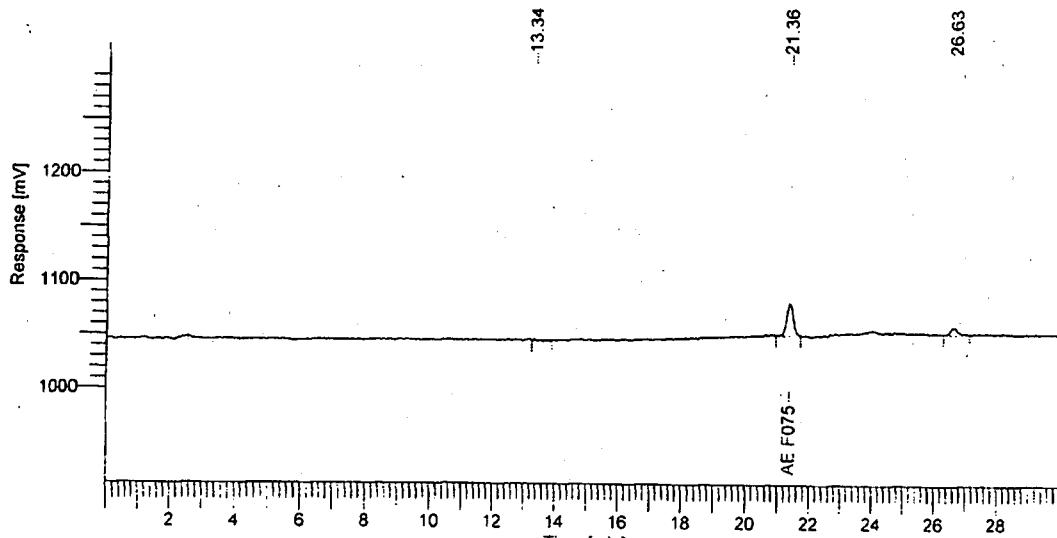
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Sample Notes:

20 ng AE F075736 in 100 µL



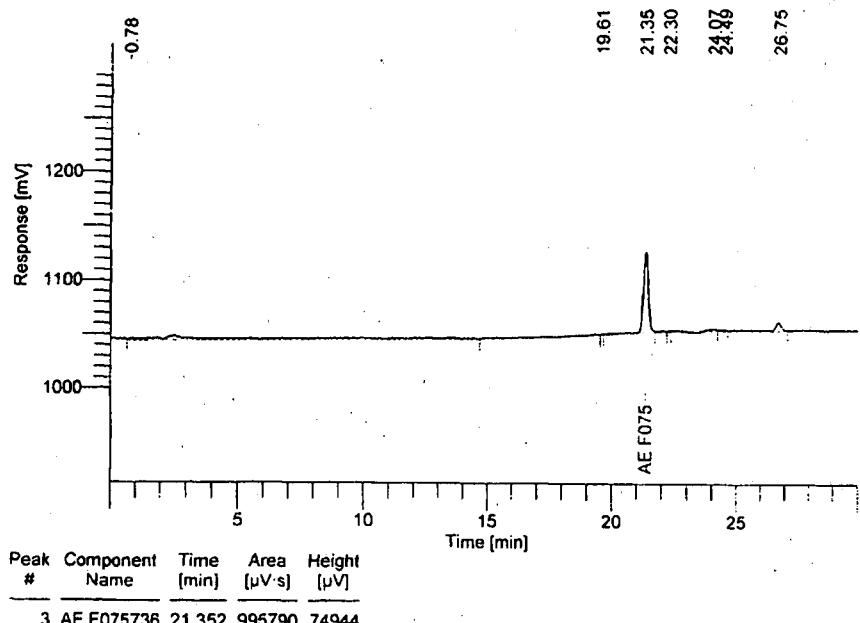
Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
2	AE F075736	21.359	420875	30083

Lab code Description

E003 Test 0.50 µg AE F075736/mL

Software Version	:	6.1.0.2:G07	Date	:	23.11.1999 08:00:10
Operator	:	Brueggemann	Sample Name	:	E003
Sample Number	:	5	Study	:	CR99/029
AutoSampler	:	NONE	Rack/Vial	:	0/0
Instrument Name	:	101	Channel	:	A
Interface Serial #	:	5237270013	A/D mV Range	:	10000
Delay Time	:	0.00 min	End Time	:	30.00 min
Sampling Rate	:	1.0000 pts/s	Area Reject	:	0.000000
Volume Injected	:	100.000000 µL	Dilution Factor	:	1.00
Sample Amount	:	1.0000	Cycle	:	5
Data Acquisition Time	:	22.11.1999 16:59:17			

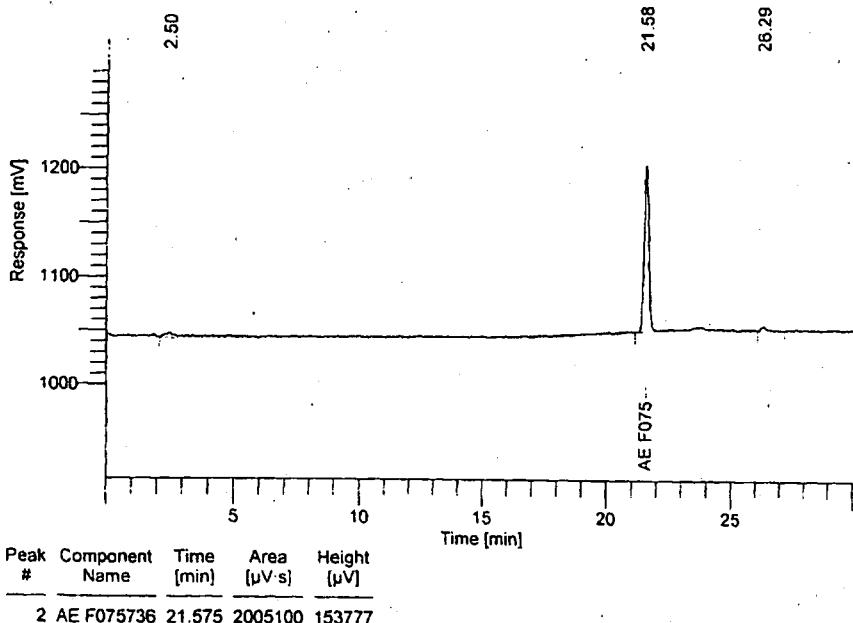
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 Sequence File : \Adef003\Turbochrom Daten\Brueggemann\Sequenzen\CR99029\bru001.seq
 Sample Notes:
 50 ng AE F075736 in 100 µL



Lab code Description
E004 Test 1.0 µg AE F075736/mL

Software Version	:	6.1.0.2:G07	Date	:	23.11.1999 08:01:43
Operator	:	Brueggemann	Sample Name	:	E004
Sample Number	:	7	Study	:	CR99/029
AutoSampler	:	NONE	Rack/Vial	:	0/0
Instrument Name	:	101	Channel	:	A
Interface Serial #	:	5237270013	A/D mV Range	:	10000
Delay Time	:	0.00 min	End Time	:	30.00 min
Sampling Rate	:	1.0000 pts/s	Area Reject	:	0.000000
Volume Injected	:	100.000000 µL	Dilution Factor	:	1.00
Sample Amount	:	1.0000	Cycle	:	7
Data Acquisition Time	:	22.11.1999 18:54:36			

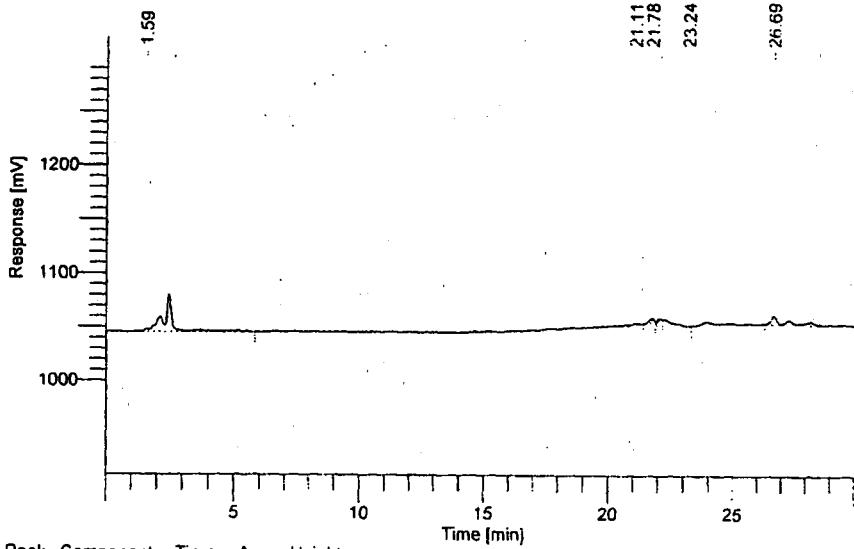
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 Sequence File : \Adef003\Turbochrom Daten\Brueggemann\Sequenzen\CR99029\bru001.seq
 Sample Notes:
 100 ng AE F075736 in 100 µL



Sample material	Lab code	Description
Drinking water (Vittel)	U001	Control

Software Version	: 6.1.0.2:G07	Date	: 23.11.1999 08:44:00
Operator	: Brueggemann	Sample Name	: U001
Sample Number	: 2	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s	Area Reject	: 0.000000
Volume Injected	: 100.000000 µL	Dilution Factor	: 1.00
Sample Amount	: 1.0000	Cycle	: 2
Data Acquisition Time	: 22.11.1999 14:06:23		

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 Sample Notes:
 1 mL / 100 µL

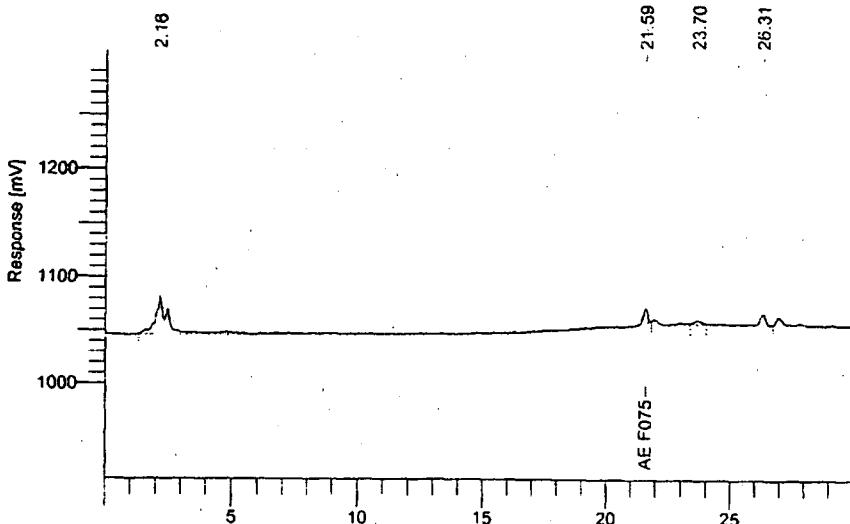


Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
-	AE F075736	21.500	0	0

Sample material	Lab code	Description
Drinking water (Vittel)	R001	Recovery 0.10 µg AE F075736/L

Software Version	: 6.1.0.2:G07	Date	: 23.11.1999 08:01:17
Operator	: Brueggemann	Sample Name	: R001
Sample Number	: 6	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s		
Volume Injected	: 100.000000 µL	Area Reject	: 0.000000
Sample Amount	: 1.0000	Dilution Factor	: 1.00
Data Acquisition Time	: 22.11.1999 17:56:56	Cycle	: 6

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 Sample Notes:
 1 mL / 100 µL

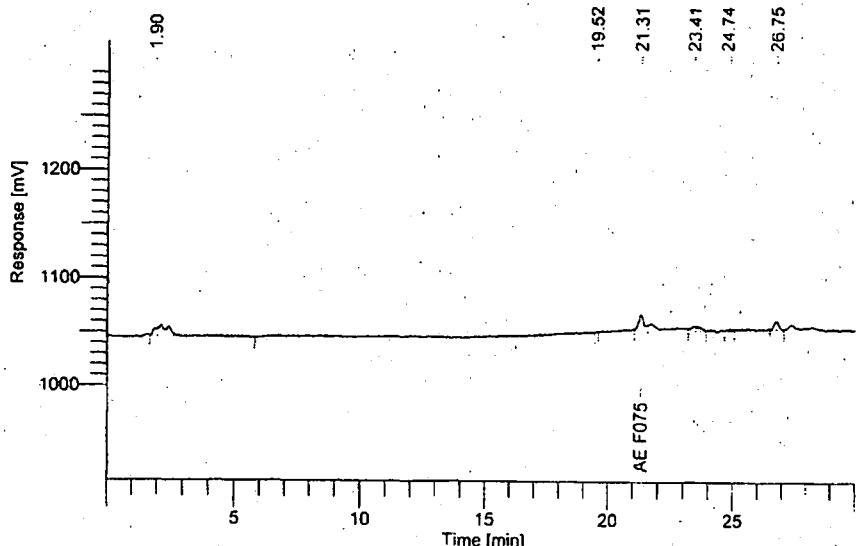


Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
2	AE F075736	21.587	170540	13841

Sample material	Lab code	Description
Drinking water (Vittel)	R002	Recovery 0.10 µg AE F075736/L

Software Version	: 6.1.0.2:G07	Date	: 23.11.1999 08:03:17
Operator	: Brueggemann	Sample Name	: R002
Sample Number	: 8	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s		
Volume Injected	: 100.000000 µL	Area Reject	: 0.000000
Sample Amount	: 1.0000	Dilution Factor	: 1.00
Data Acquisition Time	: 22.11.1999 19:52:16	Cycle	: 8

Raw Data File : \Adef003\Turbochrom
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 Sequence File : \Adef003\Turbochrom.Daten\Brueggemann\Sequenzen\CR99029\bru001.seq
 Sample Notes:
 1 mL / 100 µL

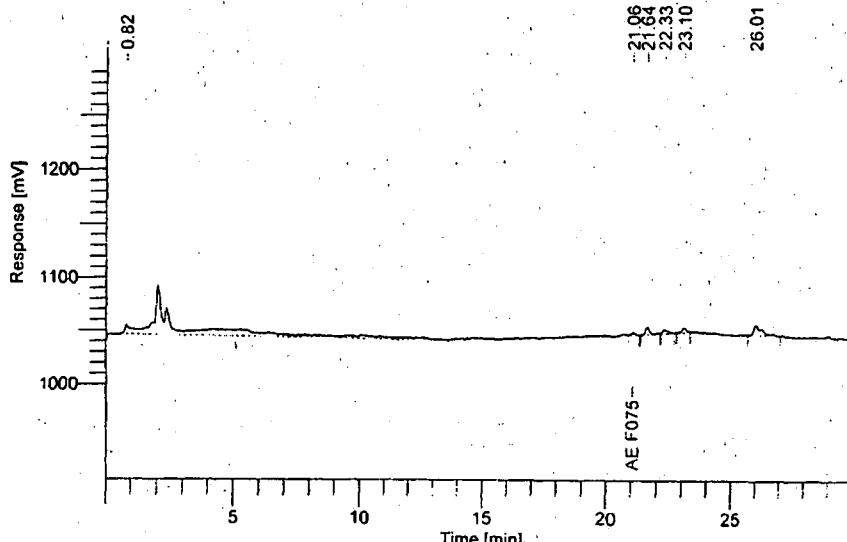


Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
3	AE F075736	21.313	138420	12673

Sample material	Lab code	Description
Drinking water (Vittel)	U005	Control

Software Version	: 6.1.0.2:G07	Date	: 08.12.1999 14:14:58
Operator	: Brueggemann	Sample Name	: U005
Sample Number	: 2	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s		
Volume Injected	: 100.000000 µL	Area Reject	: 0.000000
Sample Amount	: 1.0000	Dilution Factor	: 1.00
Data Acquisition Time	: 08.12.1999 09:42:55	Cycle	: 2

Raw Data File : \\Adef003\Turbochrom
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 Sequence File : \\Adef003\Turbochrom Daten\Brueggemann\Sequenzen\CR99029\bru003.seq
 Sample Notes:
 1 mL / 100 µL

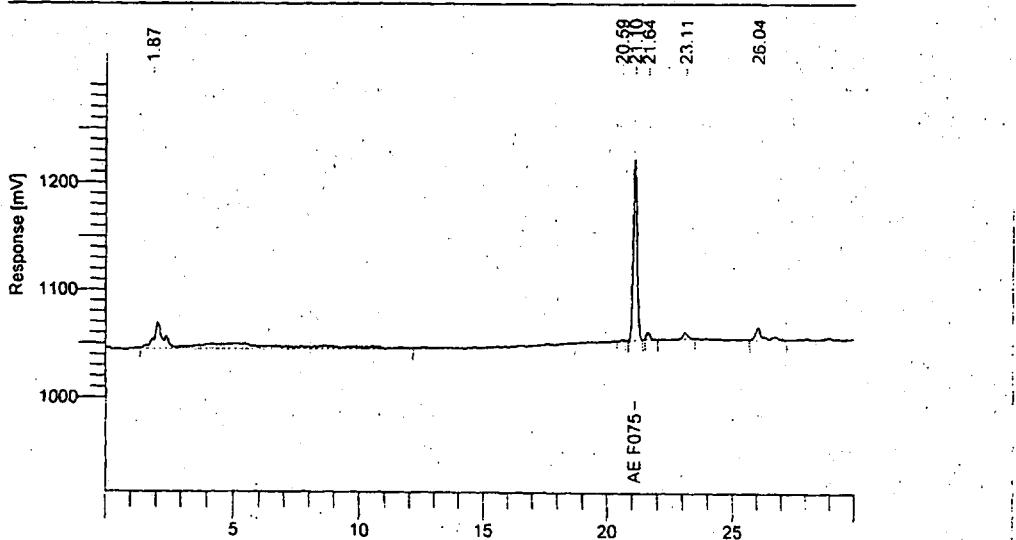


Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
2	AE F07536	21.065	38485	2869

Sample material	Lab code	Description
Drinking water (Vittel)	R009	Recovery 1.0 µg AE F075736/L

Software Version	: 6.1.0.2:G07	Date	: 08.12.1999 14:17:04
Operator	: Brueggemann	Sample Name	: R009
Sample Number	: 6	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s	Area Reject	: 0.000000
Volume Injected	: 100.000000 µL	Dilution Factor	: 1.00
Sample Amount	: 1.0000	Cycle	: 6
Data Acquisition Time	: 08.12.1999 13:33:00		

Raw Data File : \\adef003\Turbochrom
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 Sequence File : \\adef003\Turbochrom Daten\Brueggemann\Sequenzen\CR99029\bru003.seq
 Sample Notes:
 1 mL / 100 µL

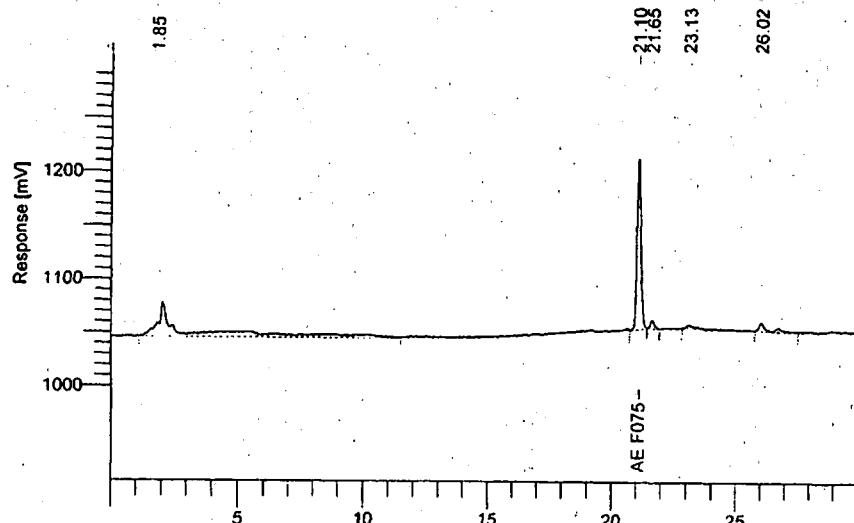


Peak #	Component Name	Time [min]	Area [µV·s]	Height (µV)
3	AE F075736	21.096	1829750	168915

Sample material	Lab code	Description
Drinking water (Vittel)	R010	Recovery 1.0 µg AE F075736/L

Software Version	: 6.1.0.2:G07	Date	: 09.12.1999 10:26:13
Operator	: Brueggemann	Sample Name	: R010
Sample Number	: 8	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s		
Volume Injected	: 100.000000 µL	Area Reject	: 0.000000
Sample Amount	: 1.0000	Dilution Factor	: 1.00
Data Acquisition Time	: 08.12.1999 15:28:06	Cycle	: 8

Raw Data File : \Adef003\Turbochrom
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 Sequence File : \Adef003\Turbochrom.Daten\Brueggemann\Sequenzen\CR99029\bru003.seq
 Sample Notes:
 1 mL / 100 µL



Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
2	AE F075736	21.096	1744620	159862

Lab code Description

E054 Test 0.10 µg (AE F115008 + AE F075736)/mL

Software Version	: 6.1.0.2:G07	Date	: 21.01.2000 15:05:50
Operator	: Brueggemann	Sample Name	: E054
Sample Number	: 1	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s		
Volume Injected	: 100.000000 µL	Area Reject	: 0.000000
Sample Amount	: 1.0000	Dilution Factor	: 1.00
Data Acquisition Time	: 21.01.2000 14:24:55	Cycle	: 1

Raw Data File : \\Adef003\Turbochrom Daten\Brueggemann\Chromatogramme\CR99029\bru007\E054.raw

Result File : \\Adef003\turbochrom daten\Brueggemann\Chromatogramme\CR99029\bru007\E054.rst

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Proc Method : \\Adef003\Turbochrom Daten\brueggemann\methoden\115008emf1299 from

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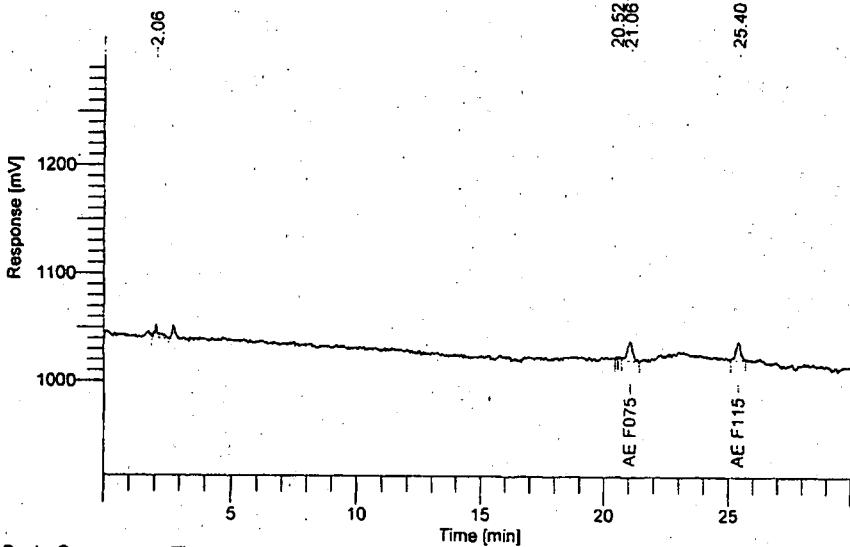
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Sequence File : \\Adef003\turbochrom Daten\Brueggemann\Sequenzen\CR99029\bru007.seq

Sample Notes:

10 ng AE F075736, 10 ng AE F115008 in 100 µL



Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
3	AE F075736	21.062	291500	18536
4	AE F115008	25.403	251150	17265

Lab code Description

E055 Test 0.20 µg (AE F115008 + AE F075736)/mL

Software Version	: 6.1.0.2:G07	Date	: 24.01.2000 07:47:54
Operator	: Brueggemann	Sample Name	: E055
Sample Number	: 3	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s		
Volume Injected	: 100.000000 µL	Area Reject	: 0.000000
Sample Amount	: 1.0000	Dilution Factor	: 1.00
Data Acquisition Time	: 21.01.2000 16:20:00	Cycle	: 3

Raw Data File : \Adept003\Turbochrom Daten\Brueggemann\Chromatogramme\CR99029\bru007\E055.raw
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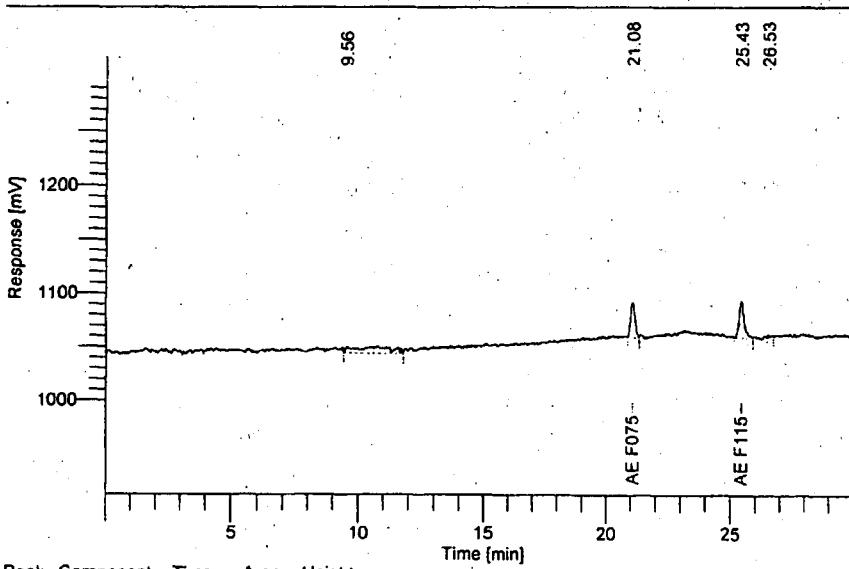
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Sequence File : \Adept003\Turbochrom Daten\Brueggemann\Sequenzen\CR99029\bru007.seq

Sample Notes:

20 ng AE F075736, 20 ng AE F115008 in 100 µL



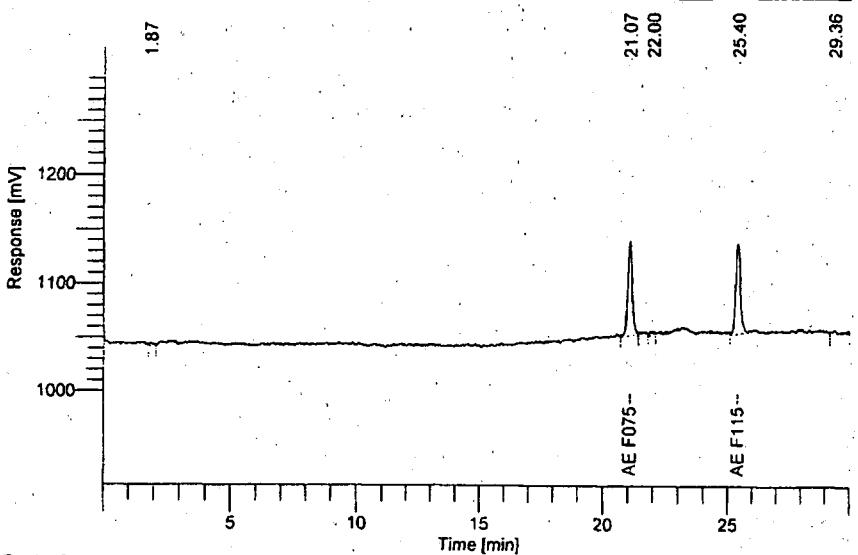
Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
2	AE F075736	21.080	412925	32959
3	AE F115008	25.431	482305	34416

Lab code Description

E056 Test 0.50 µg (AE F115008 + AE F075736)/mL

Software Version	: 6.1.0.2;G07	Date	: 24.01.2000 07:48:57
Operator	: Brueggemann	Sample Name	: E056
Sample Number	: 5	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s	Area Reject	: 0.000000
Volume Injected	: 100.000000 µL	Dilution Factor	: 1.00
Sample Amount	: 1.0000	Cycle	: 5
Data Acquisition Time	: 21.01.2000 18:15:05		

Raw Data File : \Adef003\Turbochrom Daten\Brueggemann\Chromatogramme\CR99029\bnu007\E056.raw
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 Sequence File : \Adef003\Turbochrom Daten\Brueggemann\Sequenzen\CR99029\bnu007.seq
 Sample Notes:
 50 ng AE F075736, 50 ng AE F115008 in 100 µL



Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
2	AE F075736	21.070	1050150	88301
4	AE F115008	25.405	1161025	84508

Lab code Description

E057 Test 1.0 µg (AE F115008 + AE F075736)/mL

Software Version	: 6.1.0.2:G07	Date	: 24.01.2000 07:51:02
Operator	: Brueggemann	Sample Name	: E057
Sample Number	: 7	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s	Area Reject	: 0.000000
Volume Injected	: 100.000000 µL	Dilution Factor	: 1.00
Sample Amount	: 1.0000	Cycle	: 7
Data Acquisition Time	: 21.01.2000 20:10:09		

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Result File : \\Adept003\Turbochrom daten\Brueggemann\Chromatogramme\CR99029\bru007\E057.rst

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Proc Method : \\Adept003\Turbochrom Daten\brueggemann\methoden\115008emf1299 from

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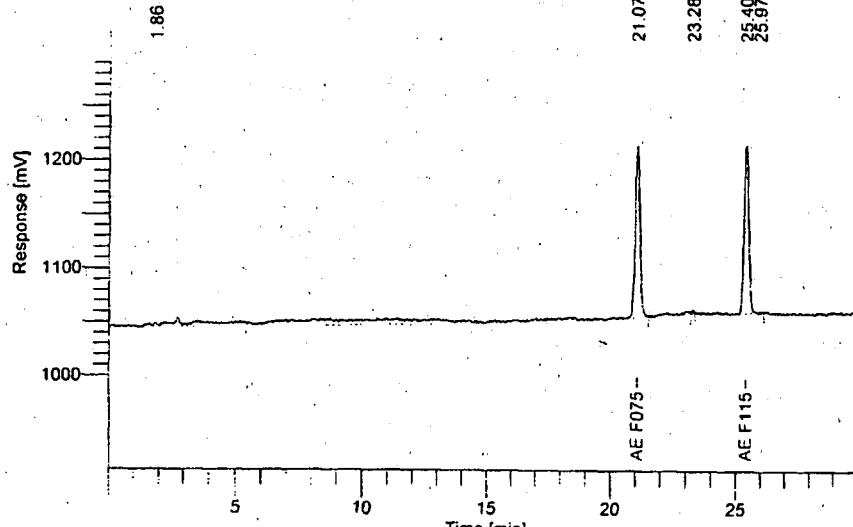
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Sequence File : \\Adept003\Turbochrom Daten\Brueggemann\Sequenzen\CR99029\bru007.seq

Sample Notes:

100 ng AE F075736, 100 ng AE F115008 in 100 µL

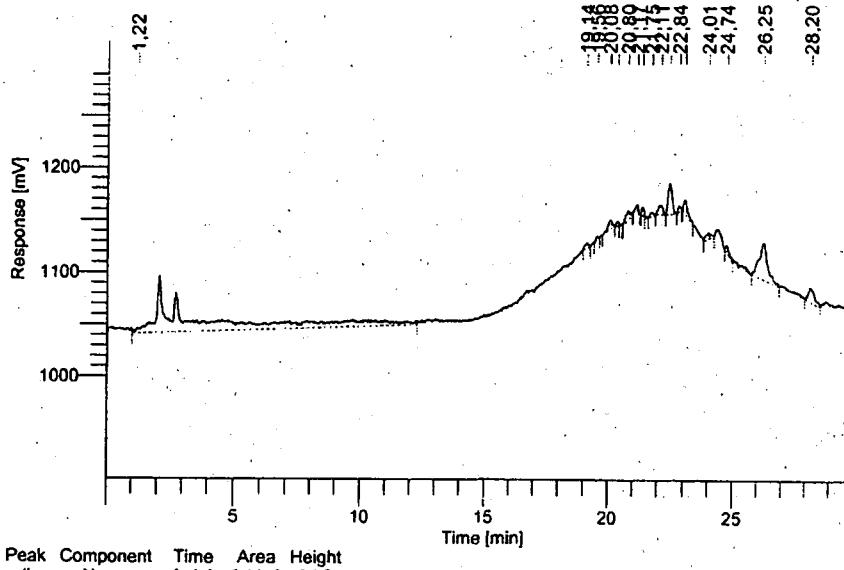


Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
2	AE F075736	21.067	2091435	159773
4	AE F115008	25.397	2136590	155863

Sample material	Lab code	Description
Surface water	U013	Control

Software Version	:	6.1.0.2:G07	Date	:	24.01.00 08:02:55
Operator	:	Brueggemann	Sample Name	:	U013
Sample Number	:	2	Study	:	CR99/029
AutoSampler	:	NONE	Rack/Vial	:	0/0
Instrument Name	:	101	Channel	:	A
Interface Serial #	:	5237270013	A/D mV Range	:	10000
Delay Time	:	0,00 min	End Time	:	30,00 min
Sampling Rate	:	1,0000 pts/s			
Volume Injected	:	100,000000 µL	Area Reject	:	0,000000
Sample Amount	:	1,0000	Dilution Factor	:	1,00
Data Acquisition Time	:	21.01.00 15:22:27	Cycle	:	2

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Result File : \adef003\Turbochrom Daten\Brueggemann\Chromatogramme\CR99029\bru007\U013.rst
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\adef003\Turbochrom Daten\Brueggemann\Chromatogramme\CR99029\bru007\U013.rst
Proc Method : \adef003\Turbochrom Daten\Brueggemann\Methoden\115008emf1299.mth
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Sequence File : \adef003\PenExe\TcCS\Ver6.1.0\Temp\U013-1328213012-20000124-075832.idx
Sample Notes:
1 mL / 100 µL

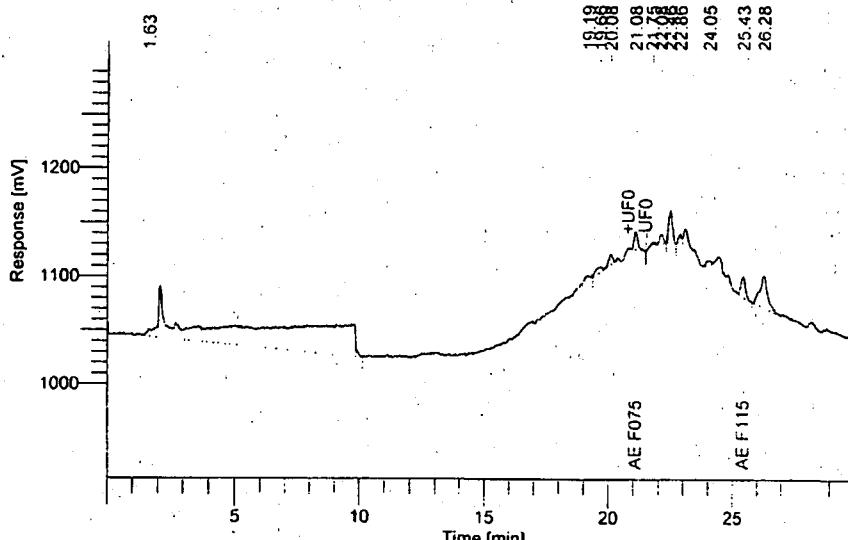


Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
-	AE F075736	21,070	0	0
-	AE F115008	25,400	0	0

Sample material	Lab code	Description
Surface water	R027	Recovery 0.10 µg (AE F115008 + AE F075736)/L

Software Version : 6.1.0.2:G07
 Operator : Brueggemann
 Sample Number : 6
 AutoSampler : NONE
 Instrument Name : 101
 Interface Serial # : 5237270013
 Delay Time : 0.00 min
 Sampling Rate : 1.0000 pts/s
 Volume Injected : 100.000000 µL
 Sample Amount : 1.0000
 Data Acquisition Time : 21.01.2000 19:12:38
 Date : 24.01.2000 07:50:39
 Sample Name : R027
 Study : CR99/029
 Rack/Vial : 0/0
 Channel : A
 A/D mV Range : 10000
 End Time : 30.00 min
 Area Reject : 0.000000
 Dilution Factor : 1.00
 Cycle : 6

Raw Data File : \\Adefit003\Turbochrom
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 Sample Notes:
 1 mL / 100 µL

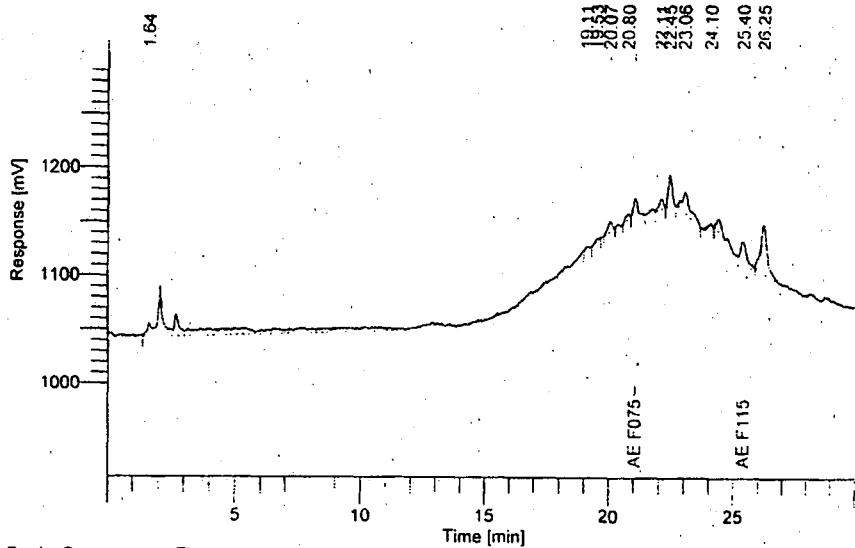


Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
5	AE F075736	21.083	187581	17441
12	AE F115008	25.429	256350	19968

Sample material	Lab code	Description
Surface water	R028	Recovery 0.10 µg (AE F115008 + AE F075736)/L

Software Version	: 6.1.0.2:G07	Date	: 24.01.2000 07:51:28
Operator	: Brueggemann	Sample Name	: R028
Sample Number	: 8	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s	Area Reject	: 0.000000
Volume Injected	: 100.000000 µL	Dilution Factor	: 1.00
Sample Amount	: 1.0000	Cycle	: 8
Data Acquisition Time	: 21.01.2000 21:07:41		

Raw Data File : \Adef003\Turbochrom
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 Sample Notes:
 1 mL / 100 µL

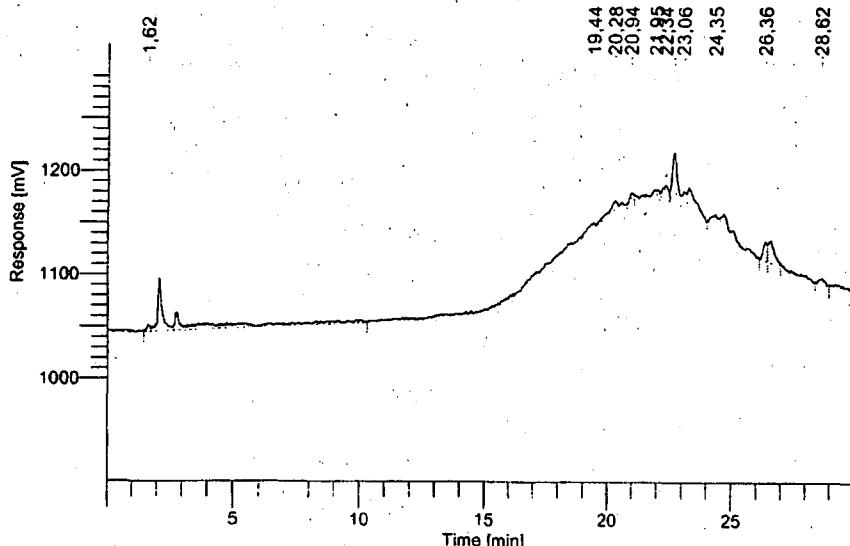


Peak #	Component Name	Time [min]	Area [µV*s]	Height [µV]
6	AE F075736	21.066	169860	15474
12	AE F115008	25.404	230545	17369

Sample material	Lab code	Description
Surface water	U015	Control

Software Version	:	6.1.0.2:G07	Date	:	25.01.00 07:56:33
Operator	:	Brueggemann	Sample Name	:	U015
Sample Number	:	2	Study	:	CR99/029
AutoSampler	:	NONE	Rack/Vial	:	0/0
Instrument Name	:	101	Channel	:	A
Interface Serial #	:	5237270013	A/D mV Range	:	10000
Delay Time	:	0,00 min	End Time	:	30,00 min
Sampling Rate	:	1,0000 pts/s	Area Reject	:	0,000000
Volume Injected	:	100,000000 µL	Dilution Factor	:	1,00
Sample Amount	:	1,0000	Cycle	:	2
Data Acquisition Time	:	24.01.00 18:52:43			

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 Sequence File : \\adef003\PenExe\TcCS\Ver6.1.0\Temp\U015-2010099093-20000125-075306.idx
 Sample Notes:
 1 mL / 100 µL

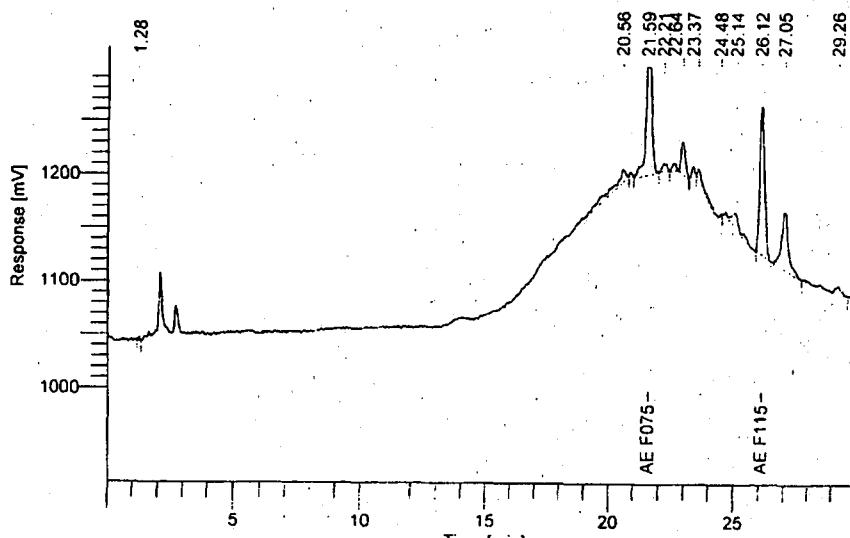


Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
-	AE F075736	21,500	0	0
-	AE F115008	25,400	0	0

Sample material	Lab code	Description
Surface water	R032	Recovery 1.0 µg (AE F115008 + AE F075736)/L

Software Version	: 6.1.0.2:G07	Date	: 25.01.2000 07:37:00
Operator	: Brueggemann	Sample Name	: R032
Sample Number	: 6	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s		
Volume Injected	: 100.000000 µL	Area Reject	: 0.000000
Sample Amount	: 1.0000	Dilution Factor	: 1.00
Data Acquisition Time	: 24.01.2000 22:42:53	Cycle	: 6

Raw Data File : \\adeft003\Turbochrom
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 Sequence File : \\adeft003\Turbochrom Daten\Brueggemann\Sequenzen\CR99029\bru008.seq
 Sample Notes:
 1 mL / 100 µL

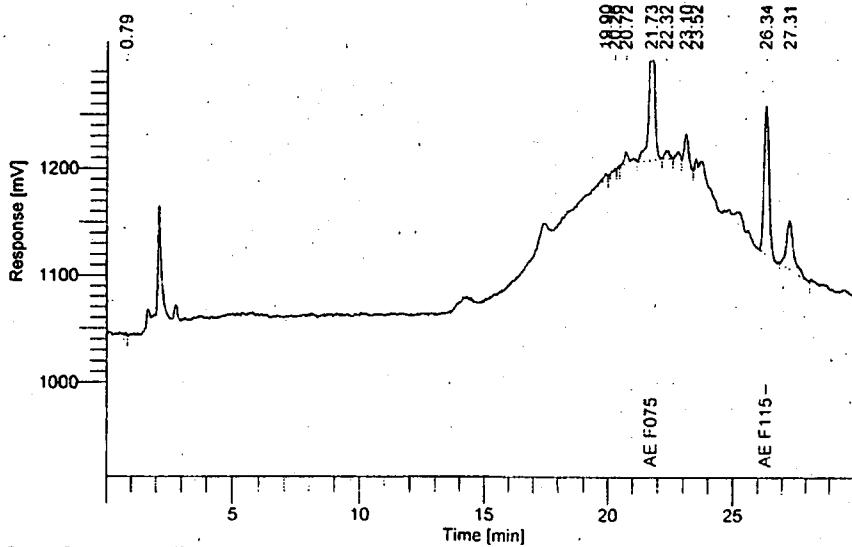


Peak #	Component Name	Time [min]	Area [µV*s]	Height [µV]
3	AE F075736	21.586	2205040	171221
11	AE F115008	26.118	1897340	137822

Sample material	Lab code	Description
Surface water	R033	Recovery 1.0 µg (AE F115008 + AE F075736)/L

Software Version	: 6.1.0.2:G07	Date	: 25.01.2000 07:38:17
Operator	: Brueggemann	Sample Name	: R033
Sample Number	: 8	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s		
Volume Injected	: 100.000000 µL	Area Reject	: 0.000000
Sample Amount	: 1.0000	Dilution Factor	: 1.00
Data Acquisition Time	: 25.01.2000 00:37:58	Cycle	: 8

Raw Data File : \adeft003\Turbochrom
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Sequence File : \adeft003\Turbochrom\daten\Brueggemann\Sequenzen\CR99029\bru008.seq
Sample Notes:
1 mL / 100 µL



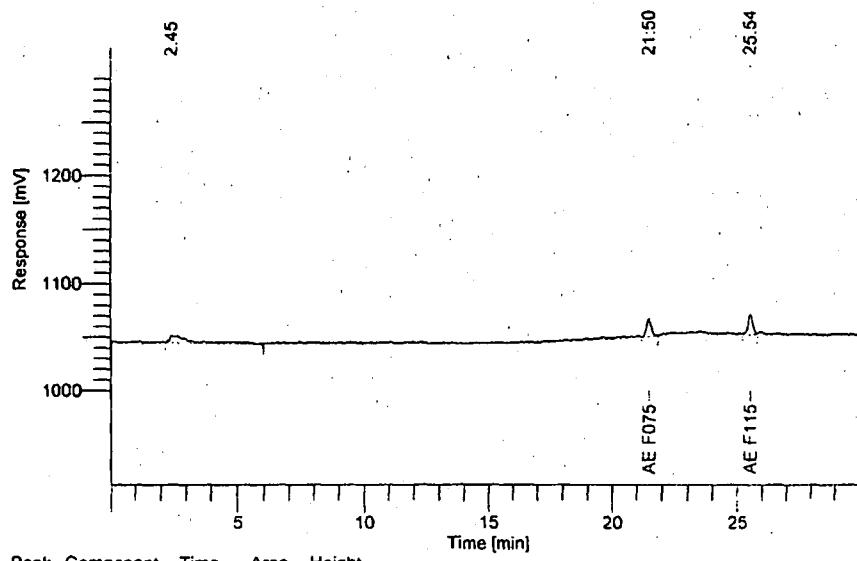
Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
5	AE F075736	21.731	2364410	178003
9	AE F115008	26.344	1973825	138633

Lab code Description

E038 Test 0.10 µg (AE F115008 + AE F075736)/mL
 Confirmation method

Software Version	: 6.1.0.2:G07	Date	: 18.01.2000 07:39:26
Operator	: Brueggemann	Sample Name	: E038
Sample Number	: 1	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s	Area Reject	: 0.000000
Volume Injected	: 100.000000 µL	Dilution Factor	: 1.00
Sample Amount	: 1.0000	Cycle	: 1
Data Acquisition Time	: 17.01.2000 16:54:51		

Raw Data File : \\adeft003\Turbochrom Daten\Brueggemann\Chromatogramme\CR99029\bru005\E038.raw
 Result File : \\Adef003\turbochrom daten\Brueggemann\Chromatogramme\CR99029\bru005\E038.rst
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 Sequence File : \\Adef003\Turbochrom Daten\Brueggemann\Sequenzen\CR99029\bru005.seq
 Sample Notes:
 10 ng AE F075736, 10 ng AE F115008 in 100 µL



Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
2	AE F075736	21.503	226270	17428
3	AE F115008	25.543	245910	19350

Lab code Description

E041 Test 1.0 µg (AE F115008 + AE F075736)/mL

Confirmation method

Software Version	: 6.1.0.2:G07	Date	: 18.01.2000 07:47:40
Operator	: Brueggemann	Sample Name	: E041
Sample Number	: 7	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s	Area Reject	: 0.000000
Volume Injected	: 100.000000 µL	Dilution Factor	: 1.00
Sample Amount	: 1.0000	Cycle	: 7
Data Acquisition Time	: 17.01.2000 22:40:09		

Raw Data File : \Adept003\Turbochrom Daten\Brueggemann\Chromatogramme\CR99029\bru005\E041.raw

Result File : \Adept003\turbochrom daten\Brueggemann\Chromatogramme\CR99029\bru005\E041.rst

Inst Method : \Adept003\Turbochrom Daten\brueggemann\methoden\115008emf1299 from \\Adept003\turbochrom daten\Brueggemann\Chromatogramme\CR99029\bru005\E041.rst

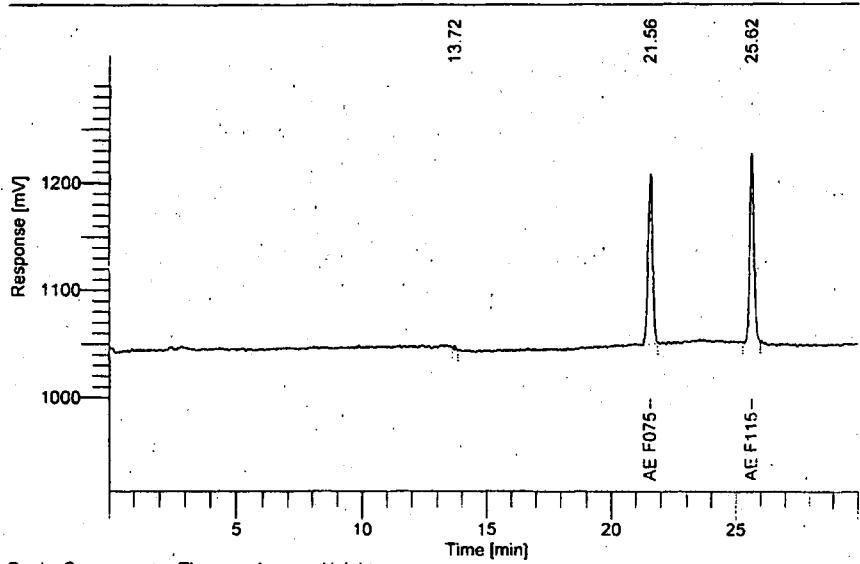
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Calib Method : \Adept003\Turbochrom Daten\brueggemann\methoden\115008emf1299 from \\Adept003\turbochrom daten\Brueggemann\Chromatogramme\CR99029\bru005\E041.rst

Sequence File : \Adept003\Turbochrom Daten\Brueggemann\Sequenzen\CR99029\bru005.seq

Sample Notes:

100 ng AE F075736, 100 ng AE F115008 in 100 µL

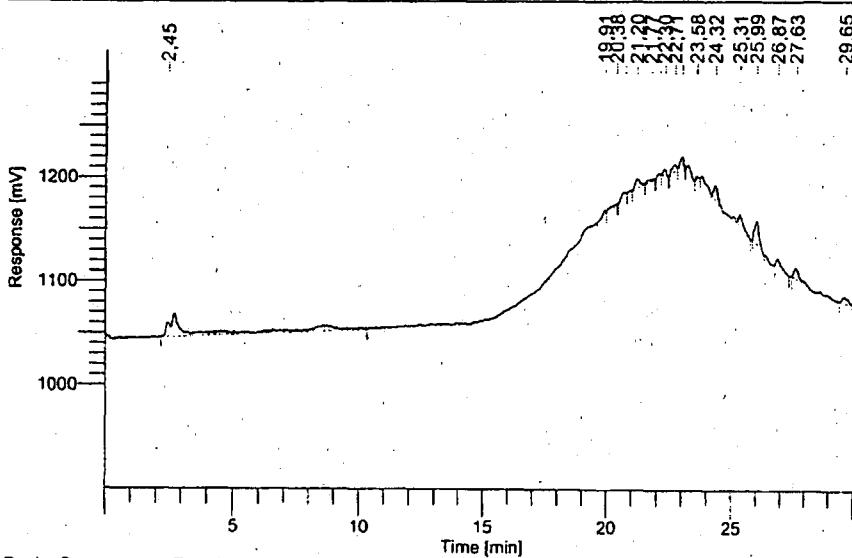


Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
2	AE F075736	21.562	2029860	159251
3	AE F115008	25.622	2181430	177369

Sample material	Lab code	Description
Surface water	U009	Control
Confirmation method		

Software Version	:	6.1.0.2:G07	Date	:	18.01.00 08:13:16
Operator	:	Brueggemann	Sample Name	:	U009
Sample Number	:	2	Study	:	CR99/029
AutoSampler	:	NONE	Rack/Vial	:	0/0
Instrument Name	:	101	Channel	:	A
Interface Serial #	:	5237270013	A/D mV Range	:	10000
Delay Time	:	0.00 min	End Time	:	30,00 min
Sampling Rate	:	1,0000 pts/s	Area Reject	:	0,000000
Volume Injected	:	100,000000 µL	Dilution Factor	:	1,00
Sample Amount	:	1,0000	Cycle	:	2
Data Acquisition Time	:	17.01.00 17:52:25			

Raw Data File : \adef003\Turbochrom
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 Sample Notes:
 1 mL / 100 µL

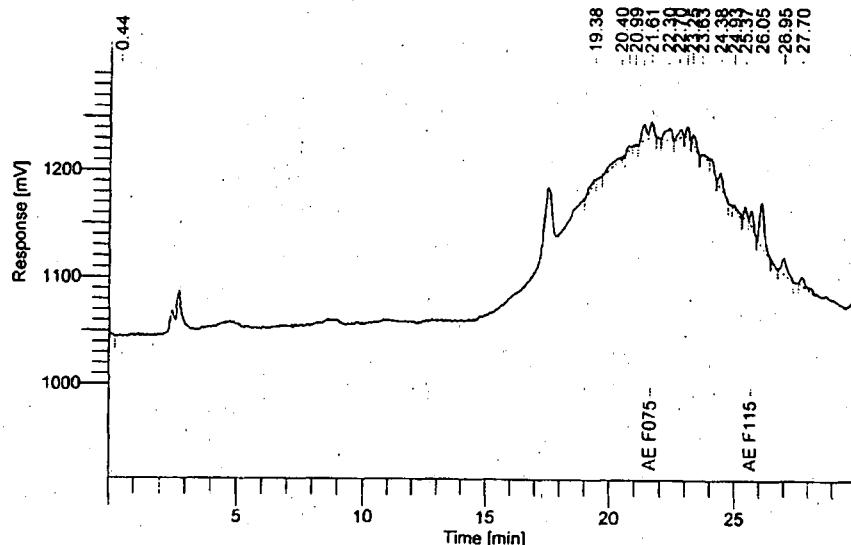


Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
-	AE F075736	21,600	0	0
-	AE F115008	25,500	0	0

Sample material	Lab code	Description
Surface water	R017	Recovery 0.10 µg (AE F115008 + AE F075736)/L
Confirmation method		

Software Version	: 6.1.0.2:G07	Date	: 18.01.2000 08:22:17
Operator	: Brueggemann	Sample Name	: R017
Sample Number	: 6	Study	: CR99/029
AutoSampler	: NONE	Rack/Vial	: 0/0
Instrument Name	: 101	Channel	: A
Interface Serial #	: 5237270013	A/D mV Range	: 10000
Delay Time	: 0.00 min	End Time	: 30.00 min
Sampling Rate	: 1.0000 pts/s	Area Reject	: 0.000000
Volume Injected	: 100.000000 µL	Dilution Factor	: 1.00
Sample Amount	: 1.0000	Cycle	: 6
Data Acquisition Time	: 17.01.2000 21:42:36		

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 \Adef003\turbochrom daten\Brueggemann\Chromatogramme\CR99029\bru005\R017.rst
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 Sample Notes:
 1 mL / 100 µL

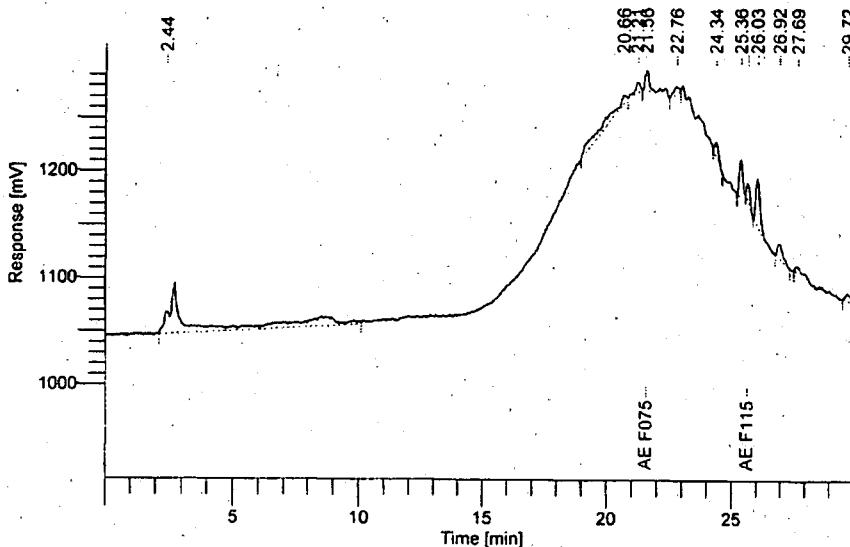


Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
7	AE F075736	21.610	127750	13586
16	AE F115008	25.641	142835	14508

Sample material	Lab code	Description
Surface water	R018	Recovery 0.10 µg (AE F115008 + AE F075736)/L
Confirmation method		

Software Version	:	6.1.0.2:G07	Date	:	18.01.2000 07:48:39
Operator	:	Brueggemann	Sample Name	:	R018
Sample Number	:	8	Study	:	CR99/029
AutoSampler	:	NONE	Rack/Vial	:	0/0
Instrument Name	:	101	Channel	:	A
Interface Serial #	:	5237270013	A/D mV Range	:	10000
Delay Time	:	0.00 min	End Time	:	30.00 min
Sampling Rate	:	1.0000 pts/s			
Volume Injected	:	100.000000 µL	Area Reject	:	0.000000
Sample Amount	:	1.0000	Dilution Factor	:	1.00
Data Acquisition Time	:	17.01.2000 23:37:41	Cycle	:	8

Raw Data File : \Adef003\Turbochrom
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 Sequence File : \Adef003\Turbochrom Daten\Brueggemann\Sequenzen\CR99029\bru005.seq
 Sample Notes:
 1 mL / 100 µL



Peak #	Component Name	Time [min]	Area [µV·s]	Height [µV]
4	AE F075736	21.564	209790	19301
8	AE F115008	25.627	154250	15150



HESSISCHES MINISTERIUM
FÜR UMWELT, ENERGIE,
JUGEND, FAMILIE UND
GESUNDHEIT

GLP-Bescheinigung

Bescheinigung

Hiermit wird bestätigt, daß die Prüfeinrichtung
Rückstände und Verbrauchersicherheit
in 65929 Frankfurt am Main

Brüningstraße 50

(Ort, Anschrift)

der Hoechst Schering AgrEvo GmbH, Werk Höchst

(Firma)

am 06.06.1997 und 10.09.1997

(Datum)

von der für die Überwachung zuständigen Behörden über
die Einhaltung der Grundsätze der Guten Laborpraxis
inspiziert worden ist.

Es wird hiermit bestätigt, daß folgende Prüfungen in
dieser Prüfeinrichtung nach den Grundsätzen der Guten
Laborpraxis durchgeführt werden:

Certificate

It is hereby certified that the test facility

Rückstände und Verbrauchersicherheit

in Frankfurt am Main

Brüningstraße 50

(location, address)

of Hoechst Schering AgrEvo GmbH, Werk Höchst

(company name)

on 06.06.1997 und 10.09.1997

(date)

was inspected by the competent authority
regarding compliance with the Principles of
Good Laboratory Practice.

It is hereby certified that studies in this
test facility are conducted in compliance with
the Principles of Good Laboratory Practice.

Prüfungen zur Bestimmung von Rückständen**Residues**

Im Auftrag

Jr. Hecker

(Dr. Hecker) Wiesbaden, den 11. Februar 1998

