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Trace Level Analysis of Herbicides in Drinking and Surface Water by Online-SPE LC/Triple Quadrupole MS to the Lower ppt Range

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Introduction

According to the requirements of the European Union drinking water directive 98/83/EC, pollutants like herbicides have to be monitored in drinking water. The currently demanded limit of detection (LOD) is 25 ng/L (25 ppt). To achieve this limit of detection with an entry level or mid-range triple quadrupole mass spectrometer, a larger volume of the water sample has to be enriched on a trapping column. Then, the compounds must be eluted to the analytical HPLC column for separation. This work describes a method to enrich and separate different trace level herbicides. The performance of the system is demonstrated by the mass spectrometric detection of a suite of 28 common neutral herbicides down to an LOD of less than 10 ppt.

Experimental

Instrumentation

Agilent 1200 Infinity Series Online-SPE solution system:

- Quaternary Pump with internal degasser
- Standard Autosampler with sample thermostat
- Flexible Cube with two 2-position/10-port valves (G4232B)
- Thermostatted Column Compartment
- Agilent 6460 Triple Quadrupole LC/MS with Agilent Jet Stream Technology

LC/MS Method:

Quaternary Pump:

- Solvent A: Water, 5 mM ammonium formate + 0.1% formic acid
- Solvent B: AcN + 5% water, 5 mM ammonium formate + 0.1% formic acid
- Flow rate: 0.4 mL/min
- Gradient: 0 min 5% B, 5 min 5% B, 20 min 98% B
- Stop time: 25 minutes
- Post time: 10 min

Thermostatted Column Compartment:

Column temperature 40 °C

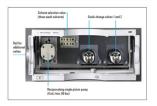
Flexible Cube (Figure 1):

- Valves: 2-position/10-port QuickChange valve heads
- Pump: 1.5 mL/min, Solvent: A1—Water, B1—AcN
- 0 minutes Pump 300 s, Solvent A1
- 5 minutes right valve change position
- 7 minutes Pump 180 s, Solvent B1
- 11 minutes Pump 300 s, Solvent A1

Standard Autosampler

- Injection volume: 1,800 μL (automated multi draw of 2 times 900 μL)
- Needle wash in vial (MeOH)
- Draw and eject speed: 1,000 µL/min
- Sample temperature: 10°C

Experimental



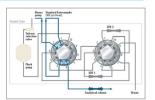
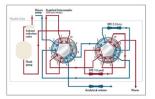


Figure 1: Flexible Cube LC module housing up to two quick-change valves, a single piston pump and 3 solvent selection valves

Figure 2A: Plumbing configuration for direct on-column injection



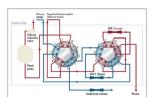


Figure 2B: **Plumbing** configuration for injection onto SPE trapping columns. The Quaternary Pump is connected with the left valve with the trapping columns (SPE2) toward the analytical column (left valve position, blue flow path). At beginning of the analysis, the piston pump is delivering the sample from the autosampler to the trapping column SPE1 (red flow path).

Figure 2C: **Plumbing** configuration for injection onto SPE trapping columns. The Quaternary Pump is connected with the left valve and the trapping columns toward analytical column for SPE (left valve position, blue flow path). After loading the SPE1, the left valve is switched and brings SPE1 in line with the analytical column (red flow path). The SPE2 is cleaned and equilibrated for the subsequent sample (red flow path).

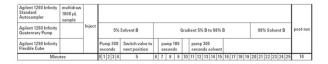


Table 1: Summary of the LC method for the Autosampler, the Quaternary Pump and the Flexible Cube.

Experimental

200.00		12-21-02-	Precursor ion		Fragment ion		Fragment ion	
Name	RT	Precursor	[M+H]+	Fragmentor	(quantifier)	CE	(qualifier)	CE
Atrazine desisopropyl	10.52	173.05	174.1	105	96.1	16	104.0	24
Carbendazim	11.15	191.07	192.1	110	160.0	16	132.0	32
Metamitron	11.67	202.10	203.1	105	175.1	12	104.1	20
Fenuron	11.81	164.09	165.1	85	72.1	16	46.1	12
Atrazine desethyl	11.93	187.06	188.0	105	146.0	16	104.0	28
Chloridazon	11.96	221.04	222.0	125	104.0	20	92.1	24
Carbetamide	13.34	236.12	237.1	75	118.1	8	192.1	4
Metoxuron	13.55	228.07/230.07	229.1/231.1	110	72.1	20	72.1	20
Monuron	13.79	198.06/200.06	199.1/201.1	95	72.1	16	72.1	16
Simazine	13.80	201.08	202.1	120	132.0	16	124.0	16
Cyanazine	14.03	240.09	241.1	120	214.1	12	104.0	32
Methabenz thiazuron	14.83	221.06	222.1	95	165.0	12	150.0	36
Chlorotoluron	14.85	212.07/214.07	213.1/215.1	100	72.1	16	72.1	16
Desmetryn	14.92	213.10	214.1	115	172.1	12	82.1	32
Atrazine	15.31	215.09	216.1	125	174.0	12	104.0	28
Isoproturon	15.48	206.14	207.1	100	72.1	16	46.1	16
Diuron	15.64	232.02/234.02	233.02/235.02	100	72.1	20	72.1	20
Monolinuron	15.71	214.05	215.1	85	126.0	12	148.0	8
Propazine	16.62	229.11	230.1	120	146.0	20	188.0	12
Linuron	16.85	248.01	249.0	90	159.9	16	182.0	12
Terbuthylazine	16.92	229.11	230.1	110	174.0	15	104.0	32
Chloroxuron	17.21	290.08/292.08	291.1/293.1	120	72.1	20	72.1	20
Irgarol 1051	17.52	253.14	254.1	120	198.1	16	83.1	28
Prometryn	17.61	241.14	242.1	125	158.0	20	200.1	16
Diflubenzuron	17.76	310.03	311.0	90	158.0	8	141.0	32
Terbutryn	17.85	241.14	242.1	110	186.0	16	68.1	48
Trietazine	18.11	229.11	230.1	125	99.0	24	132.0	20
Neburon	18.71	274.06	275.1	120	88.1	12	57.1	24

Triple Quadrupole MS method with Agilent Jet Stream thermal gradient focusing technology:

- Gas temperature: 325°C
- Gas flow: 9 L/min
- Nebulizer: 35 psi
- Sheath gas temperature: 350 °C
- Sheath gas flow: 12 L
- · Capillary: 4,000 V
- Nozzle 0 V

Table 2:

MRM and dynamic MRM MS method, showing the identified optimum fragmentor [v] and collision energy [eV] values for the individual pesticides as well as for the quantifier and qualifier ions. The retention time was used to develop the dynamic MRM method with a window of ± 3 times the peak width around the compound retention time. For some chlorinated compounds, the transition from both chlorine isotopes to the same fragment were used when other transitions were of lower abundance.

Results and Discussion

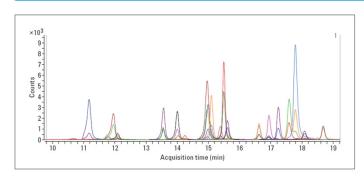
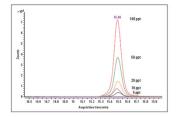
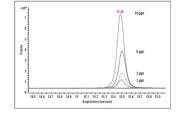


Figure 3: Chromatograms of a calibration standard with a concentration of 100 ppt (ng/L) each for the 28 pesticides measured by the final SPE-LC dynamic MRM method with quantifier and qualifier ion.

Calibration curves for each individual compound were obtained by diluting the stock solution containing all 28 pesticides at a concentration of 100 ng/L (100 ppt) in a dilution series down to 0.5 ng/L (0.5 ppt). The pesticides were measured with the developed online-SPE LC method using dynamic MRM. Each calibration standard was injected four times with a volume of 1,800 μ L and enriched on the SPE trapping column. The value at a signal-to-noise (S/N) ratio of 3 was used for the LOD and the value at a S/N ratio of 10 was used as LOQ. The calibration curve was calculated from LOQ up to 100 ng/L (Figure 4).





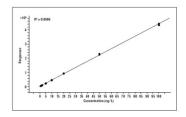


Figure 4: A) Dynamic MRM chromatograms for the quantifier transition m/z 207.1 & 72.1 of isoproturon, at a concentration of 5 to 100 ppt. B) 1 to 10 ppt. C) Calibration curve of isoproturon at a concentration of 1 ppt–100 ppt (seven levels, all used, 28 points, all used), linear coefficient 0.9986, LOQ 1ppt.

Results and Discussion

Figures 4A and 4B show the quantifier transition (m/z 207.1 & m/z 72.1) of isoproturon at a concentration range of 100 ppt to 5 ppt (Figure 4A) and for 10 ppt to the LOQ of 1 ppt (Figure 4B) at a retention time of 15.48 minutes. The calibration curve of the seven levels from the LOQ of 1 ppt to 100 ppt was calculated including all 28 injections and resulted in a linear coefficient of 0.9986 (Figure 5C).

Finally, water samples from the Rhine river, tap water, and spring water were spiked with all 28 pesticides to a final concentration of 25 ppt. Analysis of all samples yielded comparable intensities for a large number of the spiked herbicides independent of the source of the water sample (Figure 5). These results indicate that residual salt contaminations from the water samples or other contaminants with high ion strength which might cause ion suppression were effectively flushed out of the SPE column. The spiked tap water and river water samples were rich in calcium hydrogen carbonate. The measured concentrations of all pesticides shown in Figure 5 were averaged dependent on the source of water. The calculated concentration precision was between 2.3% and 2.8%. The concentration accuracy was always above 90%.

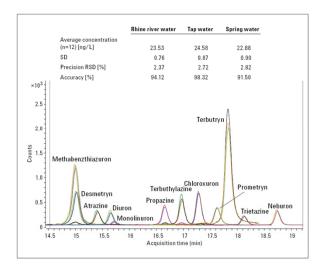


Figure 5: Water samples from a tap, the Rhine river, and a spring spiked with 25 ppt of all 28 pesticides. The retention time window from 14.5 minutes to 19.1 minutes is shown. The table shows the average measured concentrations of all pesticides (n=12) within the retention time window dependent on the sources of water together with precision RSD and accuracy.

Name	r.t.	L0Q (ng/L) (S/N=10)	R ²	LOD (ng/L) (S/N=3)	Area RSD (%)	r.t. RSD (%)	Recovery (%)
Atrazine desisopropyl	10.52	5	0.9969	2.0	5.0	0.20	84.3
Carbendazim	11.15	1	0.9971	0.5	7.8	0.10	88.8
Metamitron	11.67	5	0.9988	2.0	5.2	0.30	87.8
Fenuron	11.81	2	0.9985	1.0	7.0	1.00	96.1
Atrazine desethyl	11.93	5	0.9971	2.0	7.0	0.10	92.2
Chloridazon	11.96	2	0.9977	1.0	6.9	0.10	96.8
Carbetamide	13.34	2	0.9981	1.0	6.9	0.70	98.5
Metoxuron	13.55	2	0.9982	1.0	7.6	0.05	96.8
Monuron	13.79	2	0.9981	1.0	6.8	0.03	97.0
Simazine	13.80	5	0.9986	2.0	7.5	0.04	97.9
Cyanazine	14.03	5	0.9965	2.0	7.6	0.06	92.0
Methabenz thiazuron	14.83	1	0.9982	0.5	3.7	0.03	95.5
Chlorotoluron	14.85	1	0.9982	0.5	5.1	0.03	94.9
Desmetryn	14.92	1	0.9986	0.5	4.7	0.10	95.6
Atrazine	15.31	2	0.9982	1.0	6.6	0.04	96.9
Isoproturon	15.48	1	0.9986	0.5	6.8	0.03	98.0
Diuron	15.64	2	0.9986	1.0	6.4	0.80	82.1
Monolinuron	15.71	5	0.9976	2.0	4.8	0.05	92.3
Propazine	16.62	2	0.9980	1.0	4.8	0.03	94.6
Linuron	16.85	5	0.9981	2.0	7.6	80.0	87.1
Terbuthylazine	16.92	1	0.9920	0.5	4.2	0.05	100.9
Chloroxuron	17.21	1	0.9983	0.5	5.2	0.02	105.5
Irgarol 1051	17.52	1	0.9965	0.5	8.7	0.07	89.8
Pormetryn	17.61	1	0.9988	2.0	4.2	0.10	94.3
Diflubenzuron	17.76	5	0.9954	2.0	6.0	0.06	78.0
Terbutryn	17.85	1	0.9988	0.5	5.6	0.80	97.4
Trietazine	18.11	5	0.9984	2.0	5.9	0.02	97.3

Table 3: Performance data for all pesticide compounds present in the study. (R2 = linear coefficient, RSD = Relative standard deviation, r.t. = retention time, RSD (%) and recovery (%)).

The recovery of the SPE trapping process was determined by comparing the peak areas of an injection onto the SPE column to a direct injection of the same concentration level and volume (900 μL of 50 ppt standard) onto the analytical column.

Conclusions

This work demonstrates the use of the Online-SPE solution for enrichment, separation, and detection of trace level pesticide residues in water samples by HPLC with triple quadrupole MS detection.

It was demonstrated that an LOD of 0.5 ppt and an LOQ as low as 1 ppt could be achieved. The methodology shows a high sample-to-sample reproducibility with area deviation of less than 7%.

The efficient online-SPE trapping process allows pesticide detection in real drinking water samples well below regulatory limits with high precision and accuracy.