

Application Bulletin 113/3 e

Determination of cadmium, lead and copper in foodstuffs, waste water and sewage sludge by anodic stripping voltammetry after digestion

Summary

Cadmium, lead and copper can be determined simultaneously in oxalate buffer by anodic stripping voltammetry (ASV) after digestion with sulfuric acid and hydrogen peroxide. Tin present in the sample does not interfere with the determination of lead.

For the voltammetric determination of tin, refer to Application Bulletin 176.

Instruments

VA instrument

capable of operating a Multi-Mode Electrode and supporting differential pulse (DP) measuring mode

«Hach Digesdahl» digestion apparatus with special quartz vessels for open wet digestion

Electrodes

WE	Multi-Mode Electrode pro Mercury drop capillary	6.1246.120 6.1226.030 or 6.1226.050
RE	Ag/AgCl reference electrode Ag/AgCl/KCl (3 mol/L) Electrolyte vessel	6.0728.x20 6.1245.010
۸.	Filled with c(KCI) = 3 mol/L	0.004000
AE	Pt rod electrode	6.0343.x00

Sample preparation

Digestion

Approx. 250 mg sample is weighed exactly into the flask of the digestion apparatus. 4 mL w(H_2SO_4) = 96% is added and the mixture heated to 200 °C. When any water present has evaporated and the mixture taken on a brown color, 1 mL w(H_2O_2) = 30% is added through the dropping funnel. After the reaction has finished and the digestion solution turned brown again, another 1 mL w(H_2O_2) = 30% is added. The

mixture is now heated up to 350 ... 400 °C. The addition of H_2O_2 has to be repeated at this temperature until the digestion solution stays clear and colorless at the boiling point of sulfuric acid. Typically, a total of up to 5 mL hydrogen peroxide solution is needed to achieve complete digestion.

After cooling down, the digestion flask is made up to 100 mL with ultrapure water. An aliquot of this digestion solution is then used for the voltammetric determination.

Removing the excess of sulfuric acid

The described digestion procedure can also be used for other applications. If the voltammetric determination is to be carried out in a supporting electrolyte with higher pH value, then the high content of sulfuric acid in the digestion solution has to be neutralized first. A large quantity of sodium hydroxide solution would be necessary for this, resulting in high blanks.

To avoid this, the sulfuric acid is evaporated almost to dryness in the digestion flask over the flame of a Bunsen burner. After cooling down, make up to 100 mL with ultrapure water.

Determination of Cd, Pb and Cu

Reagents

All of the used reagents must be of purest quality possible (for analysis or for trace analysis*).

- Sulfuric acid, w(H₂SO₄) = 96%, for trace analysis*, CAS 7664-93-9
- Hydrogen peroxide solution, w(H₂O₂) = 30%, for trace analysis*, CAS 7722-84-1
- Hydrochloric acid, w(HCI) = 30%, for trace analysis*, CAS 7647-01-0
- Ammonium oxalate $(NH_4)_2C_2O_4 \cdot 1$ H_2O , for analysis, CAS 6009-70-7
- Ammonium chloride NH₄Cl, for trace analysis*, CAS 12125-02-9



- Cadmium standard stock solution, β(Cd²+) = 1 g/L (commercially available)
- Lead standard stock solution, β(Pb²⁺) = 1 g/L (commercially available)
- Copper standard stock solution, β(Cu²⁺) = 1 g/L (commercially available)
- Ultrapure water, resistivity >18 MΩ·cm (25 °C), type I grade (ASTM D1193)
- * e.g., Merck suprapur®, Honeywell Fluka TraceSelect® or equivalent

Solutions

Oxalate buffer (pH = 1)	$c((NH_4)_2C_2O_4) = 0.25 \text{ mol/L}$ $c(NH_4CI) = 0.35 \text{ mol/L}$ $c(HCI) = 0.3 \text{ mol/L}$ $35.5 \text{ g } (NH_4)_2C_2O_4 \cdot 1 \text{ H}_2O, 18.7 \text{ g}$ $NH_4CI \text{ and } 31.6 \text{ mL w(HCI)} = 30\%$ are dissolved in warm water. After cooling down to room temperature, the solution is made
	up to 1 L with ultrapure water.

Standard solutions

Standard solutions with lower concentrations (e.g. 1 mg/L) are prepared from the corresponding stock solutions (1 g/L) by dilution with c(HCI) = 0.01 mol/L or $c(HNO_3) = 0.015$ mol/L.

Analysis

Measuring solution:

5 mL oxalate buffer

+ 10 mL (diluted) digestion solution

If the metals to be determined are present in concentrations above the linear working range, the digestion solution has to be diluted accordingly with ultrapure water.

The concentrations are determined by standard addition.

Parameters

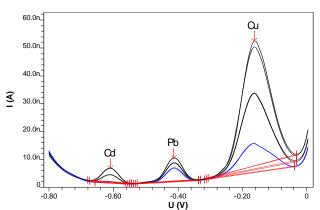
Voltammetric				
HMDE				
DP - Differential pulse				
2000 min ⁻¹				
-0.8 V				

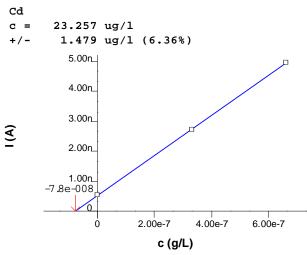
Application Bulletin 113/3 e

Determination of cadmium, lead and copper in foodstuffs, waste water and sewage sludge by anodic stripping voltammetry after digestion

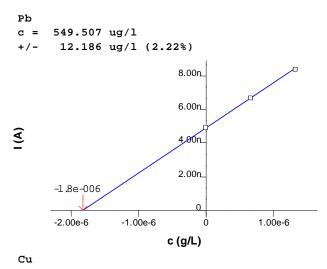
Waiting time 1	60 s		
Equilibration time	10 s		
Sweep			
Start potential	-0.8 V		
End potential	0 V		
Potential step	0.006 V		
Potential step time	0.15 s		
Sweep rate	0.04 V/s		
Pulse amplitude	0.05 V		
Substance			
Name	Cadmium		
Characteristic potential	-0.6 V		
Name	Lead		
Characteristic potential	-0.41 V		
Name	Copper		
Characteristic potential	-0.17 V		

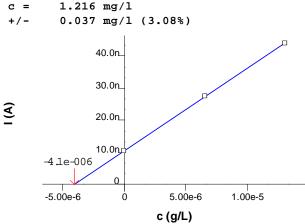
Example











Result

Sample	Artichoke juice		
Sample size	1 mL		
β(Cd)	0.023 mg/L		
β(Pb)	0.549 mg/L		
β(Cu)	1.216 mg/L		

Comments

- The digestion procedure can also be applied for the determination of other non-volatile metals.
- For volatile analytes (e.g. As or Hg) a closed digestion apparatus is necessary, e.g. a microwave digestion system or a «High Pressure Asher HPA».
- In order to remain within the linear range of the method, the concentrations of Cd, Pb and Cu in the measuring vessel should not exceed 50 ... 80 μg/L each. The total concentration of the three metals should not exceed 200 μg/L. Higher concentrations overload the working

Application Bulletin 113/3 e

Determination of cadmium, lead and copper in foodstuffs, waste water and sewage sludge by anodic stripping voltammetry after digestion

electrode during deposition. To avoid this, the time for deposition (Waiting time 1) can be reduced.

• Limits of quantitation in the digestion solution (deposition time 60 s):

Cd 0.1 μg/LPb 0.1 μg/LCu 0.5 μg/L

Lower limits of quantitation can be obtained by increasing the time for deposition (Waiting time 1).



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Appendix

Full report for the determination of Cd, Pb and Cu in artichoke juice using the 757 VA Computrace

Determinati Sample ID Creator met Creator det Modified by	on : 0428 : dige .hod : .erm.:	0943_digested plan	ested art nt e Dat Dat Dat	e:	28	Time: Time:	09:43:13
Method Title Remark1 Remark2	: AB11 : Cd , : Dige : 5mL	3CdPbCu.i Pb and (sted Art: 0xalate	mth Cu determ ichoke Ju ouffer +	ination acco ice: 1 mL di 10mL sample	rding to AB1 gested, fill (5mL digeste	113 Led up t ed sampl	co 100 mL) Le + 5mL water)
Sample amou Cell volume	: 15.	000 mL					
Substance Conc. Conc.dev. Amount Add.amount							
VR V	nA	I.mean	Std.Dev	. I.delta	Comments		
1 - 1 -0.6	0 0 524	0 531	0 01	0 000			
1 - 2 - 0.6 $2 - 1 - 0.6$	0.539 0.539 2.722	2.715	0.01	0 2.184			
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	09 2.708 09 4.910 09 5.000	4.955	0.06	0 2.184 4 2.240			
Substance Conc. Conc.dev. Amount Add.amount	: Pb : 1. : 0. : 27. : 10.	832 ug/L 041 ug/L 475 ng 000 ng	(2.	22%)			
VR V	nA	I.mean	Std.Dev	. I.delta	Comments		
1 - 1 -0.4	13 4.911	4.888		2 0.000			
2 - 1 -0.4	13 4.865 13 6.653 6.644 0.012 1.756 13 6.636 13 8.285 8.346 0.088 1.702						
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	6.636 13 8.285 13 8.408	8.346	0.08	8 1.702			
Substance Conc. Conc.dev. Amount Add.amount	: Cu : 4. : 0. : 60. : 100.	052 ug/L 125 ug/L 782 ng 000 ng	(3.	08%)			
				. I.delta	Comments		
				8 0.00			
2 - 1 -0.163	.63 27.38	27.46	0.11	5 17.04			
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	63 42.94 63 44.96	43.95	1.42	4 16.49			
Substance	Calibr.	Y.re	g/offset	Slope	Mean deviat	. Corr.	Coeff.
Cd Pb Cu	std.add. std.add. std.add.	5 4 1	.192e-010 .897e-009 .043e-008	6.697e-00 2.674e-00 2.574e-00	3 5.552e-0 3 7.119e-0 3 5.283e-0)11)11)10	0.99985 0.99962 0.99909
Final resul	ts			+/- Res. dev	. % Co	omments	
Cd: default				1.479	6.361		
Pb: default	=	549.507	ug/l	12.186	2.218		
Cu: default	=	1.216	mg/l	0.037	3.082		





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Method print for the determination of Cd, Pb and Cu with the 757 VA Computrace

```
Method parameters
         : AB113CdPbCu.mth
            : Cd , Pb and Cu determination according to AB113
Remark1 : Digested Artichoke Juice: 1 mL digested, filled up to 100 mL)
Remark2 : 5mL Oxalate buffer + 10mL sample (5mL digested sample + 5mL water)
                            : Standard addition
Calibration
Technique
                         : 15.000
: 5.000
: digested plant e
Cell volume (mL)
Sample amount (mL)
Sample ID
Voltammetric parameters
                                   : DP - Differential Pulse
                               : 1 mA
: 100 nA
Highest current range
Lowest current range
Electrode
                                   : HMDE
Drop size (1..9)
Stirrer speed (rpm)
                                   : 2000
No. of additions
No. of replications
Measure blank
Addition purge time (s)
Initial purge time (s)
                                   :
Conditioning cycles
Start potential (V)
                                           0.000
End potential (V)
No. of cycles
                                           0.000
                                     :
Cleaning potential (V)
Cleaning time (s)
Deposition potential (V)
Deposition time (s)
                                          -0.800
Equilibration time (s)
Start potential (V)
End potential (V)
                                          -0.800
                                          0.000
Voltage step (V)
Voltage step time (s)
Sweep rate (V/s)
Pulse amplitude (V)
                                           0.006
                                          0.150
                                           0.050
Pulse time (s)
                                           0.040
Cell off after measurement :
                                            Yes
Peak evaluation
Peak evaluation
                                    : Height
Peak evaluation
Minimum peak width (V.steps): 5
Minimum peak height (A): 1.000e-010
Smooth factor
Reverse peaks
Reverse sweep
                                    : No
Substances
Cd : -0.600 \text{ V} +/- 0.050 \text{ V} Standard solution : 1 0.100 \text{ mg/L} Addition volume (mL) : 0.050 \text{ default} : Final result (Cd) = (le+006 / 0.01) * Mass.conc + 0 - 0
                                : -0.400 V +/- 0.050 V
                              : 2 1.000 mg/L
: 0.010
Standard solution
Addition volume (mL)
                              : Final result (Pb) = (1e+0.06 / 0.01) * Mass.conc + 0 - 0
                               : -0.150 V
                                               +/- 0.050 V
Standard solution : 3 1.000 mg/L
Addition volume (mL) : 0.100
default : Final result (Cu) = (1000 / 0.01) * Mass.conc + 0 - 0
```