

Pražské analytické centrum inovací

Projekt CZ.04.3.07/4.2.01.1/0002 spolufinancovaný ESF a Státním rozpočtem ČR

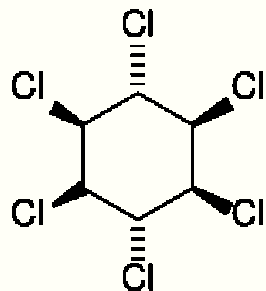


Trends in sample preparation: innovative strategies from trace elements to metalloproteins



Marco Aurélio Zezzi Arruda
Associate Professor
GEPAM - <http://gepam.iqm.unicamp.br>
e-mail: zezzi@iqm.unicamp.br





Sample preparation: why?

Lindane determination in vegetable samples

1,2,3,4,5,6-hexachlorocyclohexane



Homogenization

Can the sample be directly injected ???

Water: 90 – 96%

Fibers: 0.4 – 1.8%

Starch: 0 – 0.6%

Salts: 0.4 – 0.7%

Fat: 0 – 1.3%

Proteins: 0.7 – 1.2%

**NON VOLATILE MATERIALS
PRESENT IN SAMPLE MATRIX**

Incompatible with GC!!!

(columns/injectors occlusion)
(contaminant introduction)
(artifacts formation)



OUTLINE

CPE

→ Cd

→ Proteins

US/MAWE

→ Inorganic

→ Organic

MIP

→ Catechol

Miscellaneous

→ Metalloproteins

from trace elements to metalloproteins



Unicamp: Institute of Chemistry



87 teachers → 4 areas
ca. 400 under-graduation students
ca. 300 MSc/PhD students



Unicamp: Institute of Chemistry



Unicamp: GEPAM



- 4 main research areas
- Sample preparation
 - Bio-analytical
 - Spectrometry
 - Mechanization



4 under-graduation students
7 MSc/PhD students
2 Post-doc collaborators



Unicamp: GEPAM



**Adilson
Alessandra
Aline Lopes
Ana Cristi
Cristiana**

**Eduardo
Eraldo
Fabiana
Herbert
Marcelo
Márcia
Pedro
Renata**



Marco A. Z. Arruda

OUTLINE

CPE

→ **Cd**

→ **Proteins**

US/MIA/VE

→ Inorganic

→ Organic

IVFP

→ Catechol

Miscellaneous

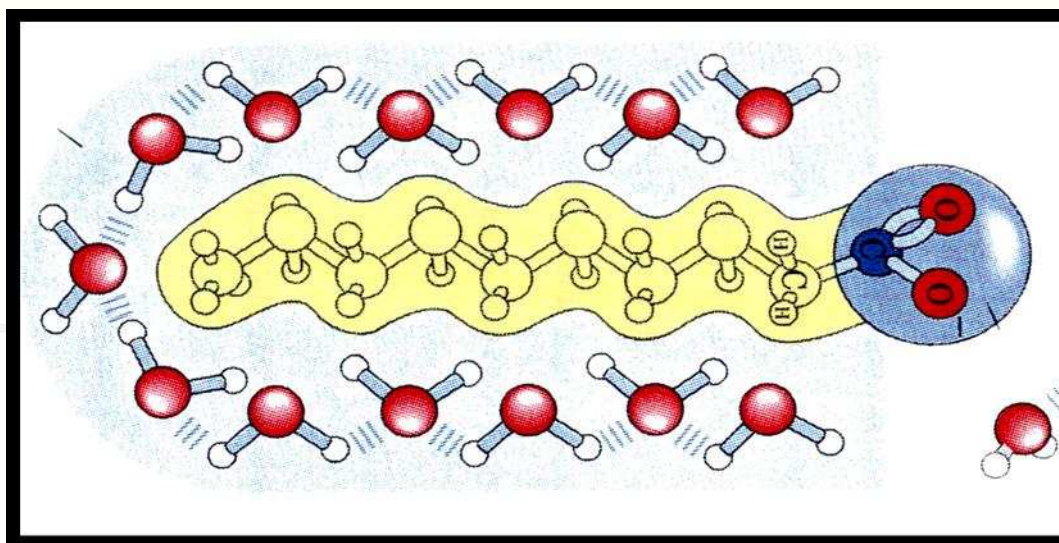
→ Metalloproteins

from trace elements to metalloproteins



SURFACTANT

(Surface active agent)



R-X structure

R: hydrocarbon chain (8-18 atoms)

X: polar or ionic head group

hydrophilic + hydrophobic groups → dissolution in water or other solvents

SURFACTANT

● Non-ionic

Brij 35, Polyoxyethylene(23); $\text{CH}_3(\text{CH}_2)_{11}(\text{OCH}_2\text{CH}_2)_{23}\text{OH}$

● Anionic

SDS, Sodium dodecil sulfate; $\text{CH}_3(\text{CH}_2)_{11}\text{OSO}_3^-\text{Na}$

● Cationic

CTAB, Cetyl trimetyl ammonium bromide; $\text{CH}_3(\text{CH}_2)_{15}\text{N}^+(\text{CH}_3)_3\text{Br}$

● Amphoteric (Zwitterionic)

DAB, 4-(Dodecyldimetyl ammonium) butirate

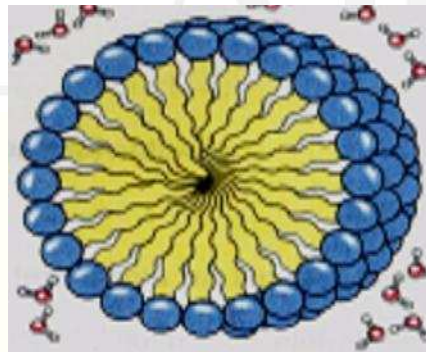
$\text{CH}_3(\text{CH}_2)_{11}\text{N}^+(\text{CH}_3)_2(\text{CH}_2)_3\text{COO}^-$



The aggregate micellar formation

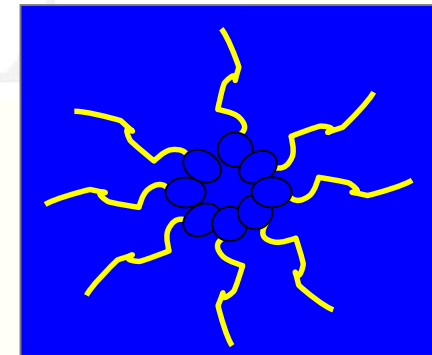
Amphiphilic molecules → self association in several solvents

organized medium
(microscopically ordered
molecular aggregates)



normal micelle

bi-layers
vesicles
micro emulsions

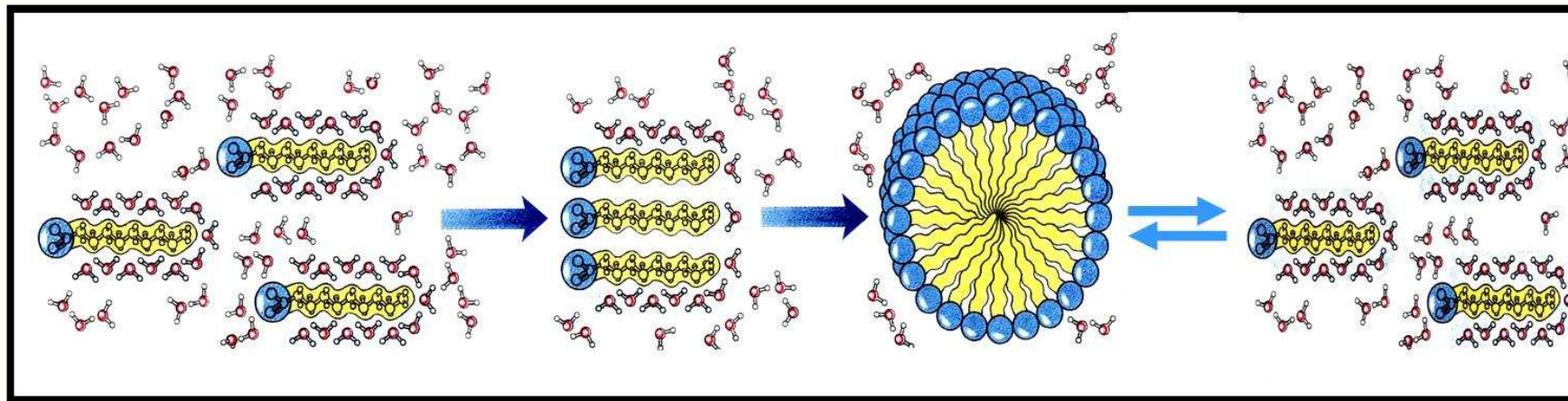


reversed micelle

final structure: monomer, solvent, surrounding ions



The aggregate micellar formation



**monomers
(below the CMC)**

**micelles and monomers in
dynamic equilibrium
(above the CMC)**



Micellar medium in analytical chemistry

- water solubilization of hydrophobic substances
 - enhancing detection (spectrophotometric methods)
 - improvement on transport and nebulization (AAS methods)
 - catalytic reactions
 - extraction
 - preconcentration
- } **CPE**

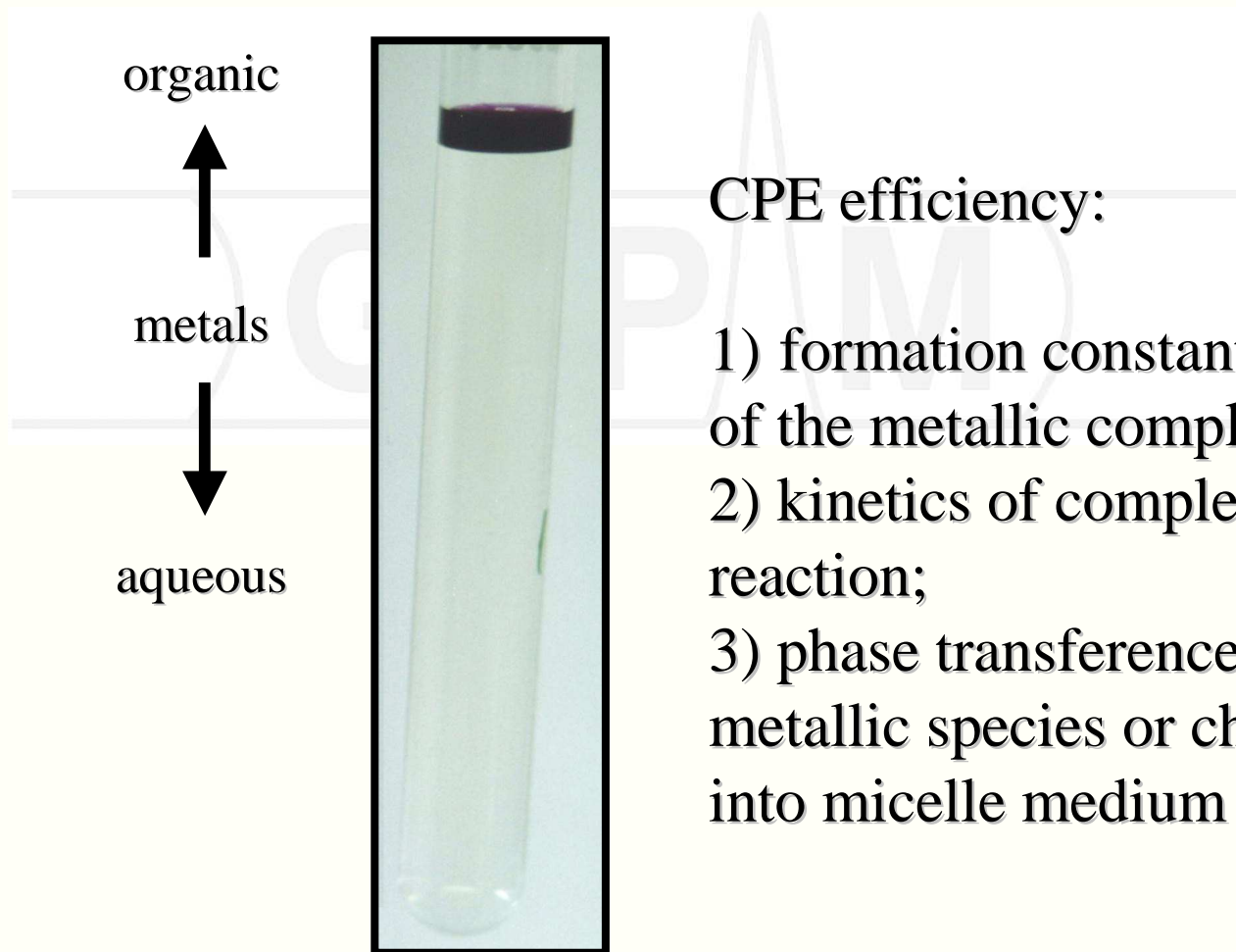


Cloud point extraction – CPE

Extraction mechanism

Bezerra et al., *App. Spectrosc. Review* 40(2005)269

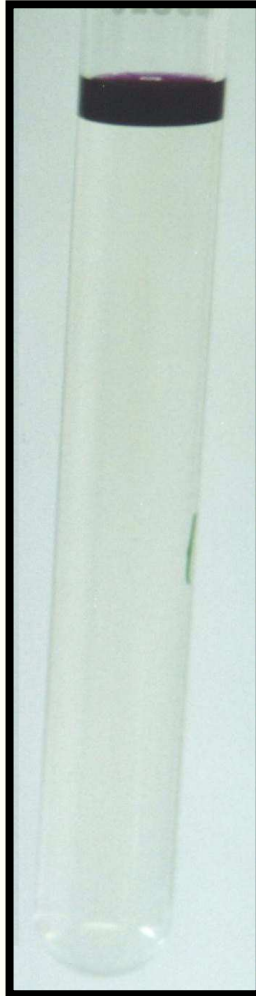
surfactant molecules act as organic solvent



Cloud point extraction – CPE

Extraction mechanism

Bezerra *et al.*, *App. Spectrosc. Review* 40(2005)269



- - hydrated nature of the surfactant phase → distribution coefficient (D) lower than conventional LLE
- LLE: > ionic strength does not seriously modify extraction efficiency; CPE: salt addition → increase on phase separation



Cloud point extraction – CPE

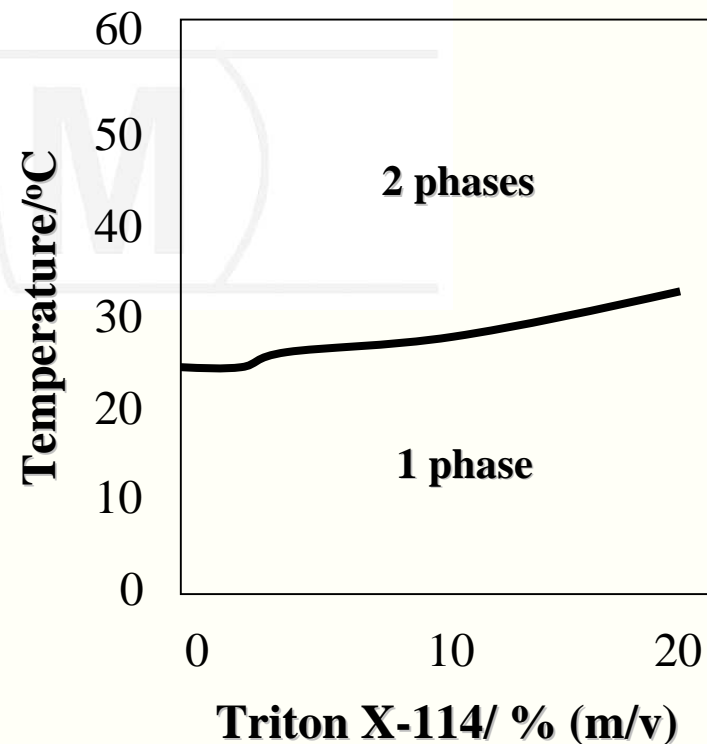
Extraction mechanism



Marco A. Z. Arruda

Gu & Sjöblom, *Colloids Surf.* 64(1992)39

Surfactant	T/°C	cmc/mmol L ⁻¹
C ₆ E ₃	40.5	
C ₁₀ E ₄	19.7	
C ₁₂ E ₅	28.9	
C ₁₄ E ₈	70.5	
Triton X-114	22.0	Triton X-100: 0.17 – 0.30 Triton X-114: 0.20 – 0.35
Triton X-100	ca. 64	



$$C(^{\circ}C) = 220 \log N_E - 5.5 N_C - 55$$

N_E = number of ethylene oxide

N_C = number of alkyl carbons

Cloud point extraction – CPE

Temperature modification

✓ **Ionic surfactants**

✓ **Inorganic salts**

✓ **Mineral acids**

✓ **Pressure**



Cloud point extraction – CPE

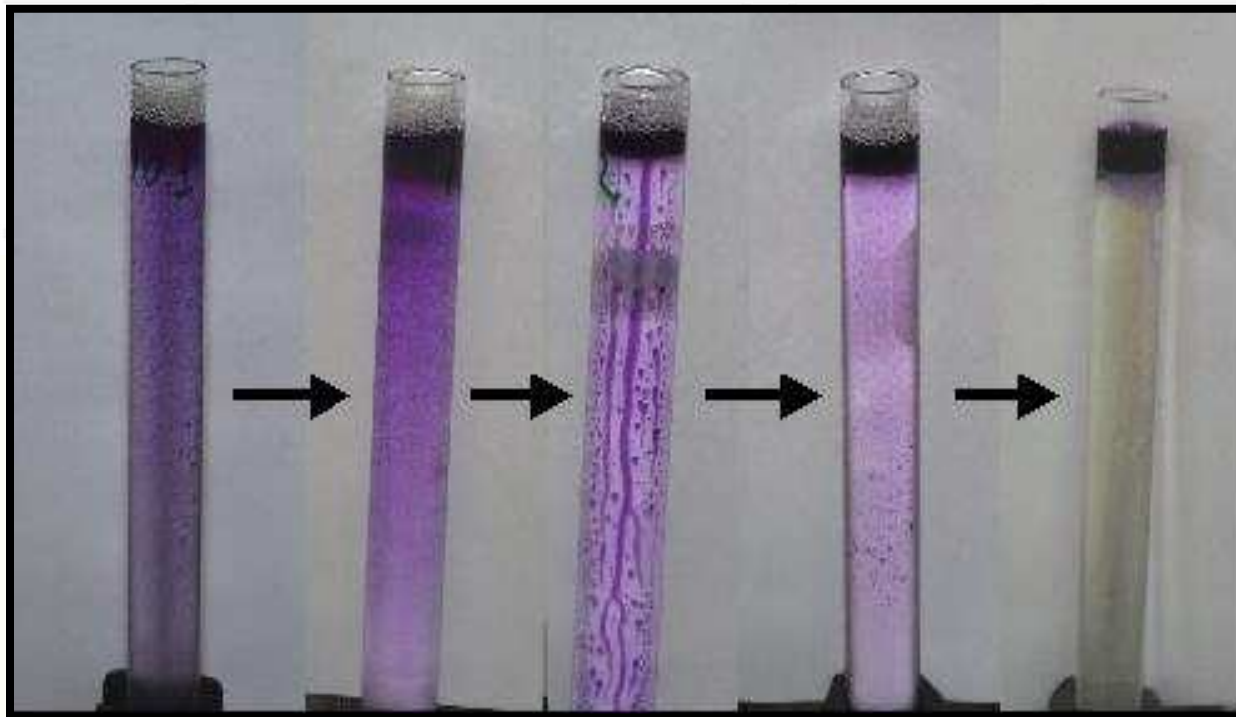
Co

+ 5-Br-PADAP

+ SDS + Triton X-100

+ NaCl

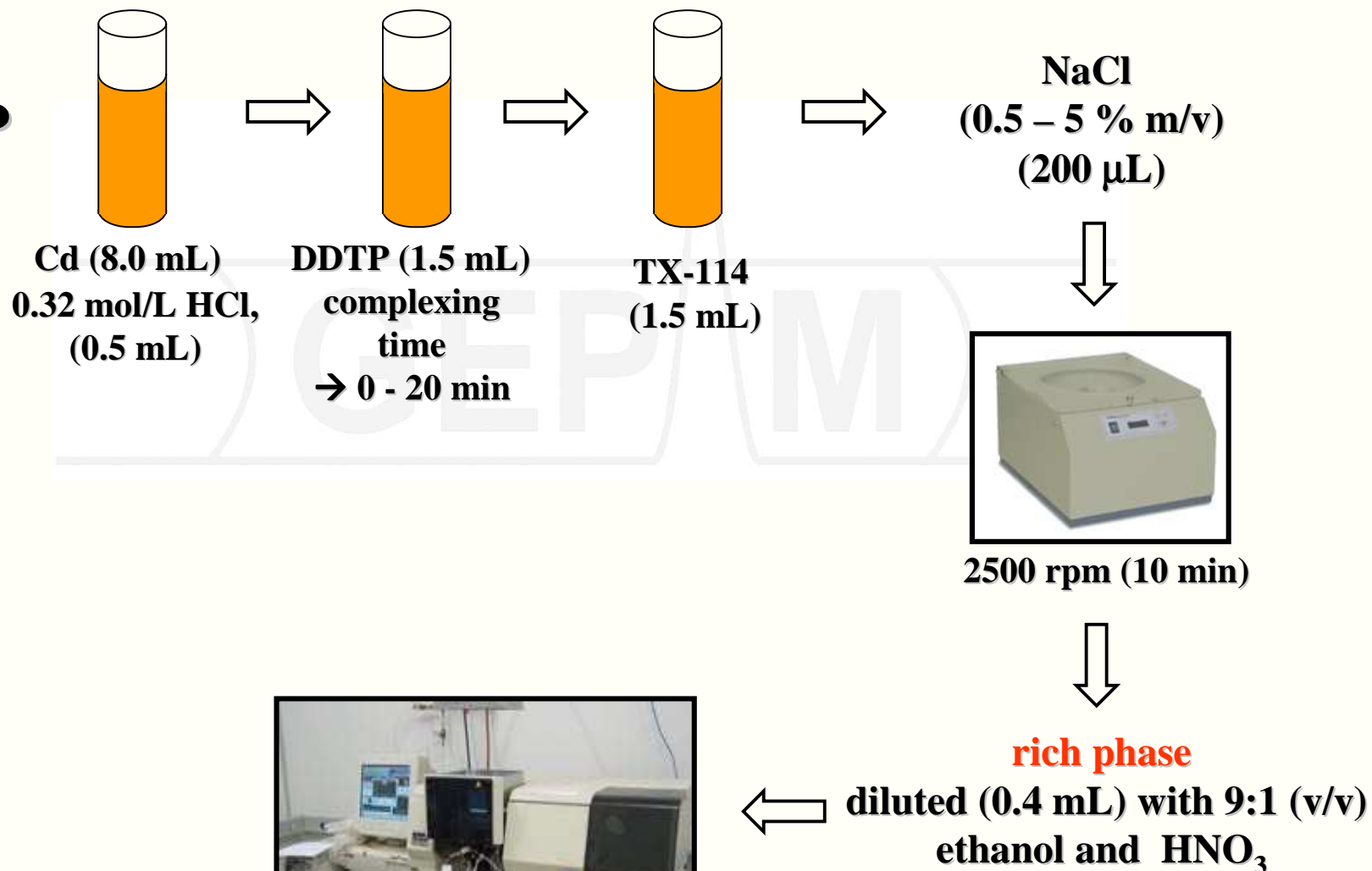
pH = 9



Cloud point extraction – CPE: Cd

Coelho, et al. *Talanta* 71(2007)353

DDTP



Cloud point extraction – CPE: Cd

Studied variables :

- complexing time
- nature of complexing agent
- complexing agent concentration
- surfactant concentration
- volume ratio between Cd and surfactant
- electrolyte concentration
- ethanol volume



Cloud point extraction – CPE: Cd

rich phase: Triton X-114 (m/v)



0.04%

0.08%

0.13%

0.44%

0.86%

0.44% - before
centrifuging



Cloud point extraction – CPE: Cd

Analytical performance

Linear equation	$Y = 0.0026 + 0.0018X$
r	0.997
Linear range ($\mu\text{g/L}$)	3 - 400
Limit of detection ($\mu\text{g/L}$)	0.9
Limit of quantification ($\mu\text{g/L}$)	2.9
Precision (100 $\mu\text{g/L}$)	4% (n=11)
CF	19



Cloud point extraction – CPE: Cd

Cd determination ($\mu\text{g/g}$) in tobacco samples by FAAS and ETAAS (n=4)

Sample	FAAS	ETAAS
1	0.546 ± 0.050	0.479 ± 0.038
2	0.437 ± 0.036	0.437 ± 0.013
3	0.353 ± 0.022	0.364 ± 0.019
4	0.229 ± 0.022	0.250 ± 0.025



Cloud point extraction – CPE: Cd

Sample	Cd added ($\mu\text{g L}^{-1}$)	Cd found ($\mu\text{g L}^{-1}$)	Recovery ^a (%)
Mineral water	0	--- ^b	-
	10	10.7 ± 1.8	107
	20	20.7 ± 4.8	104
Lake water	0	--- ^b	-
	10	11.4 ± 1.0	96
	20	22.0 ± 1.8	101
Physiological serum	0	--- ^b	-
	10	10.5 ± 1.5	105
	20	18.6 ± 0.2	93
Tobacco 1	0	10.9 ± 1.0	-
	4	15.3 ± 1.4	103
Tobacco 2	0	8.7 ± 0.7	-
	4	13.4 ± 1.6	106
Tobacco 3	0	7.1 ± 0.5	-
	4	11.5 ± 0.5	104
Tobacco 4	0	4.3 ± 0.4	-
	4	8.6 ± 0.6	103

^a recovery in fortified samples; ^b < LOQ



Cloud point extraction – CPE: Organics

Quina & Hinze, *Ind. Eng. Chem. Res.* 38(1999)4150

PAHs – river water

C₁₂E₄, 40 °C, HPLC-F (fluorescence)

Fungicides (captan, captafol) – river water

Triton X-114, 40 °C, HPLC-EC (electrochemical detector)

Fulvic acid – river water

Triton X-100, 90 °C, HPLC-UV (ionic pair)

Phenol – aqueous solution

PONPE-10, 70 °C, GC-FID



Cloud point extraction – CPE: Proteins

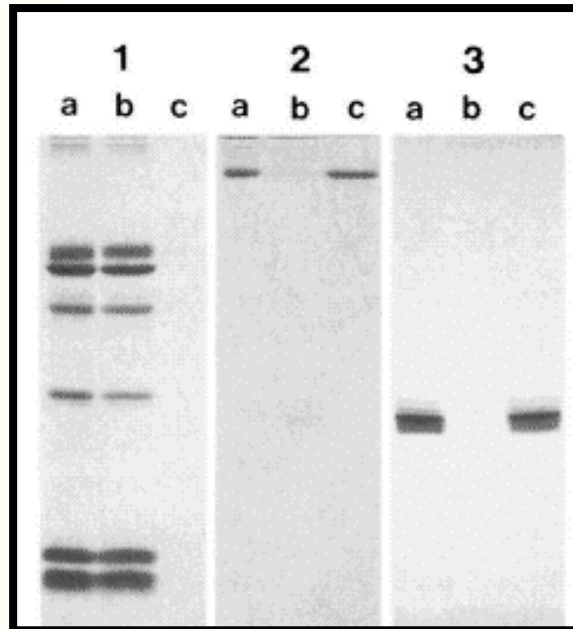
THE JOURNAL OF BIOLOGICAL CHEMISTRY
Vol. 256, No. 4, Issue of February 25, pp. 1604-1607, 1981
Printed in U.S.A.

Phase Separation of Integral Membrane Proteins in Triton X-114 Solution*

(Received for publication, March 31, 1980, and in revised form, October 6, 1980)

Clément Bordier‡

From the Biozentrum der Universität Basel, Klingelbergstr. 70, CH-4056 Basel, Switzerland



a = before phases separation

b = aqueous phase after separation

c = surfactant phase after separation

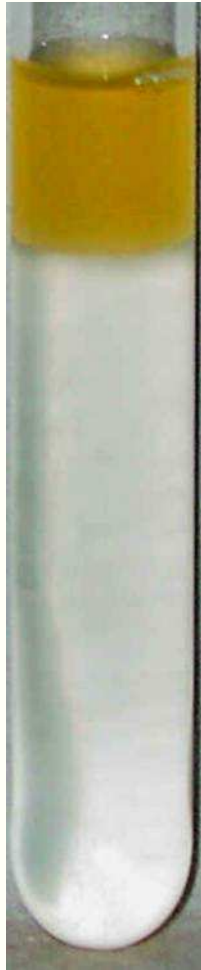
1. serum albumin, catalase, ovalbumin, concanavalin A, myoglobin and cytochrome c;
2. human erythrocyte acetylcholin-esterase; 3. bacteriorhodopsin



Cloud point extraction – CPE: Proteins

Extraction mechanisms

Nikas et al., *Macromolecules* 25(1992)4797



$$K_p = \exp\left\{- (\phi_t - \phi_b)(1 + R_p / R_0)^2\right\}$$

$$K_p = \exp\left\{- (\phi_t - \phi_b)(1 + R_p / R_0)^3\right\}$$

Φ_t = top fraction volume

Φ_b = bottom fraction volume

R_p = protein hydrodynamic radius

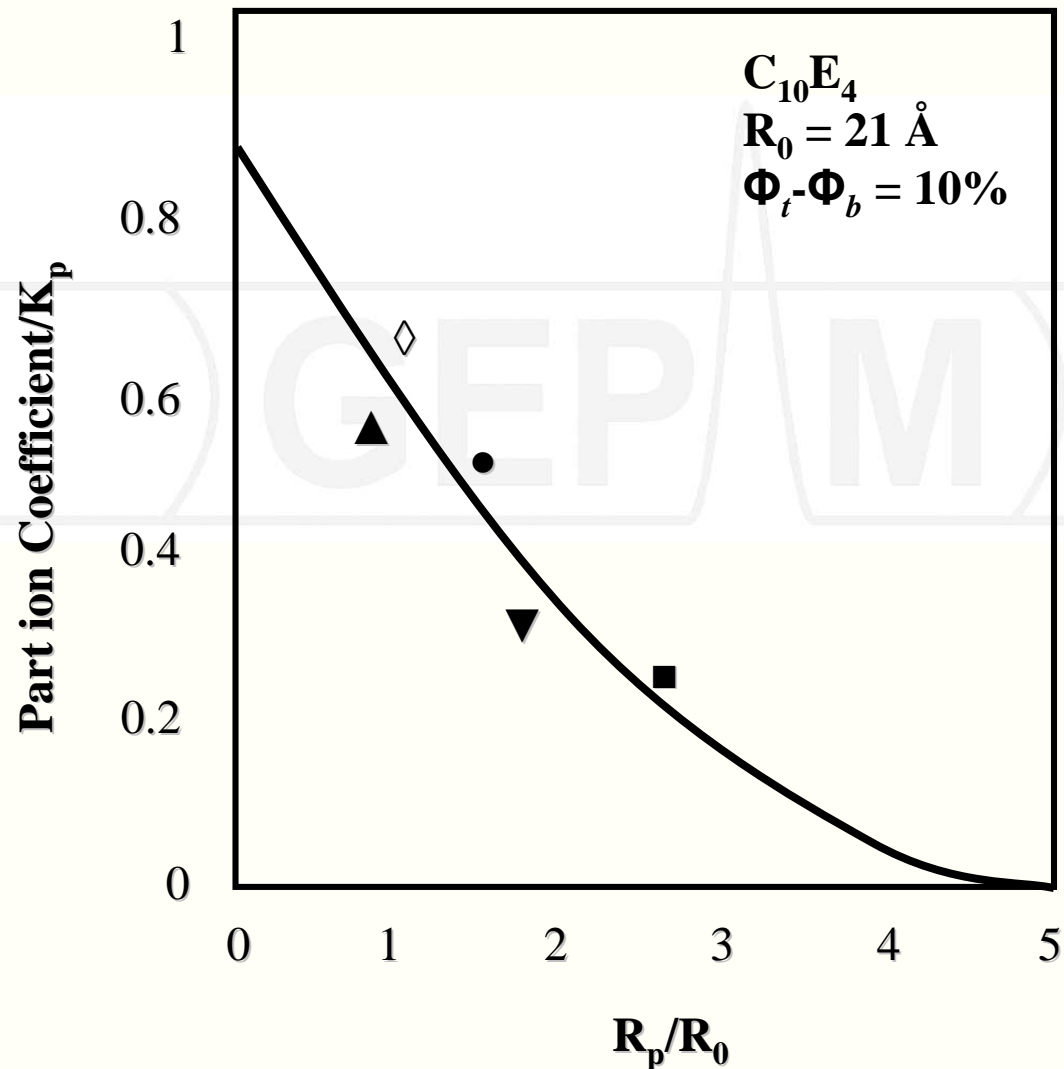
R_0 = transversal section radius of the cylindrical micelle (or spherical)



Cloud point extraction – CPE: Proteins

Extraction mechanisms

Tani et al., *Anal. Sci.* 14(1998)875



- ▲ $R_p = 19 \text{ \AA}$
Cytochrome C
- ◇ $R_p = 22 \text{ \AA}$
Soybean inhibitor
- $R_p = 29 \text{ \AA}$
Ovalbumin
- ▼ $R_p = 36 \text{ \AA}$
BSA
- $R_p = 52 \text{ \AA}$
Catalase



Cloud point extraction – CPE: Proteins

Tani et al., *J. Chromatogr. A* 780(1997)229

Bacteriorhodopsin, hemoglobin in serum

Triton X-114, 0.15 mol/L NaCl (pH 7.4), 30 °C

Cytochrome *b*₅, cytochrome *c*₁, Fe-S proteins
in bacteria

Triton X-114, 0.05-0.15 mol/L NaCl (pH 7.5), 30 °C

Pyruvate oxidase in *Escherichia Coli*

Triton X-114, 0.15 mol/L NaCl, (pH 6.0), 30 °C

Tyrosinase in mushroom

Triton X-114 (pH 7.3), 37 °C



Cloud point extraction – CPE

Proteins in milk

Lopes *et al.*, *Anal. Chim. Acta* 590(2007)166



NaCl_(s), 8 mL Triton X-114
(in KH₂PO₄/NaOH buffer) + milk



centrifugation
1780 g, 10 min



Optimized conditions:

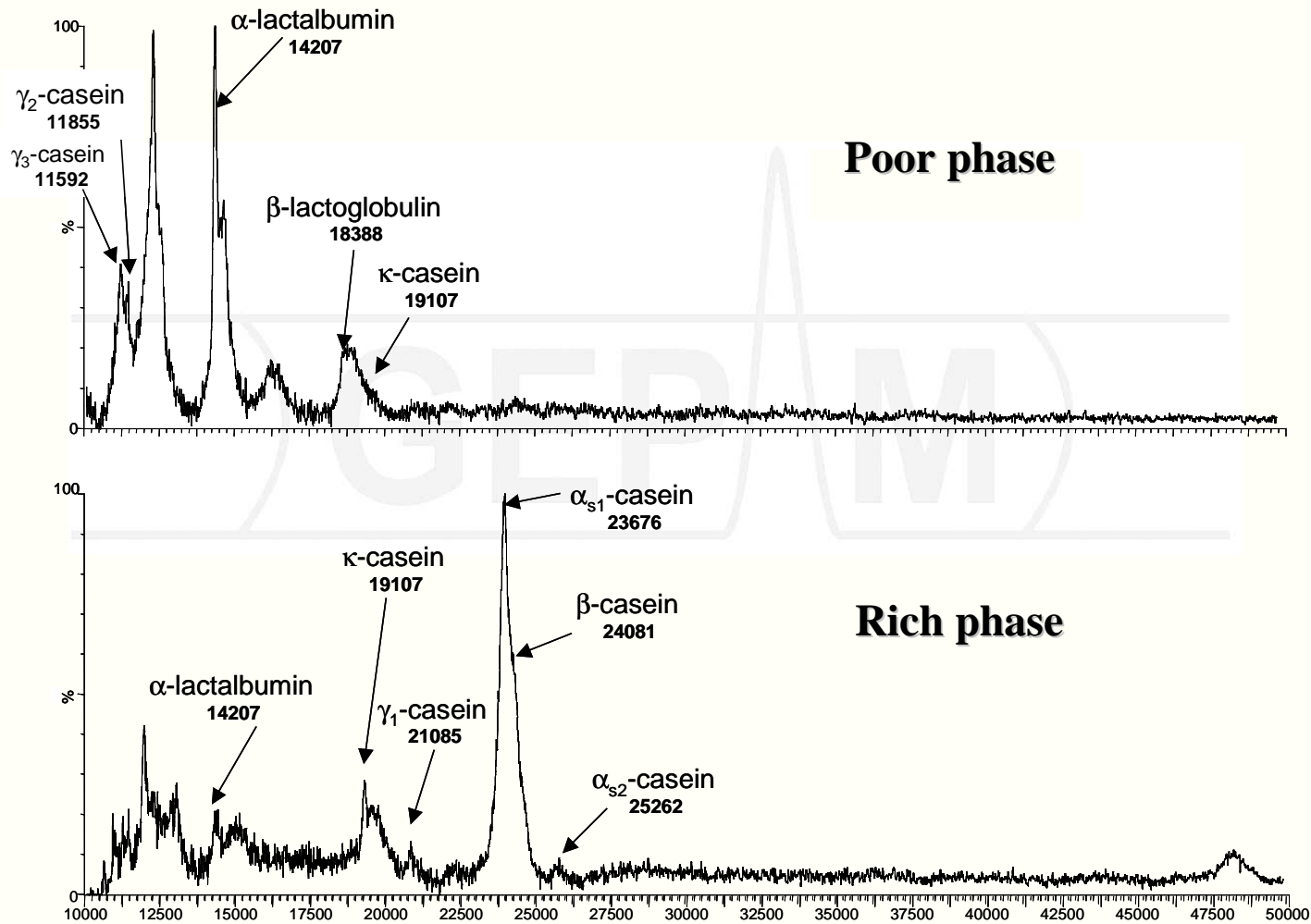
- 1% (w/v) Triton X-114;
- 50 μ L – sample volume
- 6% (w/v) NaCl,
- pH = 7.0

- ✓ Rich phase: surfactant removed using acetone
- ✓ Protein quantification: Bradford method



Cloud point extraction – CPE

Proteins in milk



Cloud point extraction – CPE

Proteins in milk

Phase	Protein	MM (Da)		Deviation (%)
		Literature	MALDI-MS	
Poor	γ_3 -Casein	11500	11592	0.8
	γ_2 -Casein	11800	11855	0.5
	α -Lactalbumin	14200	14207	0.05
	β -Lactoglobulin	18360	18388	0.2
	κ -Casein	19000	19107	0.6
Rich	α -Lactalbumin	14200	14207	0.05
	κ -Casein	19000	19107	0.6
	γ_1 -Casein	21000	21085	0.4
	α_{s1} -Casein	23600	23676	0.3
	β -Casein	24000	24081	0.3
	α_{s2} -Casein	25250	25262	0.05



Cloud point extraction – CPE

Some references

C. Bordier, *J. Biol. Chem.*, 236(1981)1604

K. Selber et al., *Process Biochem.*, 39(2004)889

T. Minuth et al., *Biotechnol. Bioeng.*, 55(1997)339

G. L. McIntire, *Critical Reviews in Anal. Chem.*, 21(1990)257

M. A. Bezerra et al., *Appl. Spectrosc. Reviews*, 40(2005)269



Cloud point extraction – CPE

Conclusions

good alternative
for extracting metals
and bio-molecules

biocompatible
system

CPE

good CF

safe, fast and low cost procedure



OUTLINE

CPE

→ Cd

→ Proteins

US/MAWE

→ Inorganic

→ Organic

IVFP

→ Catechol

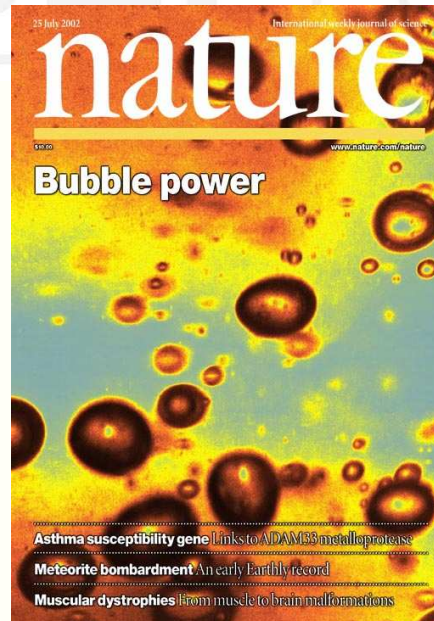
Miscellaneous

→ Metalloproteins

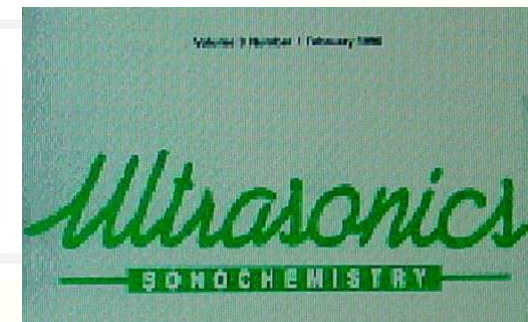
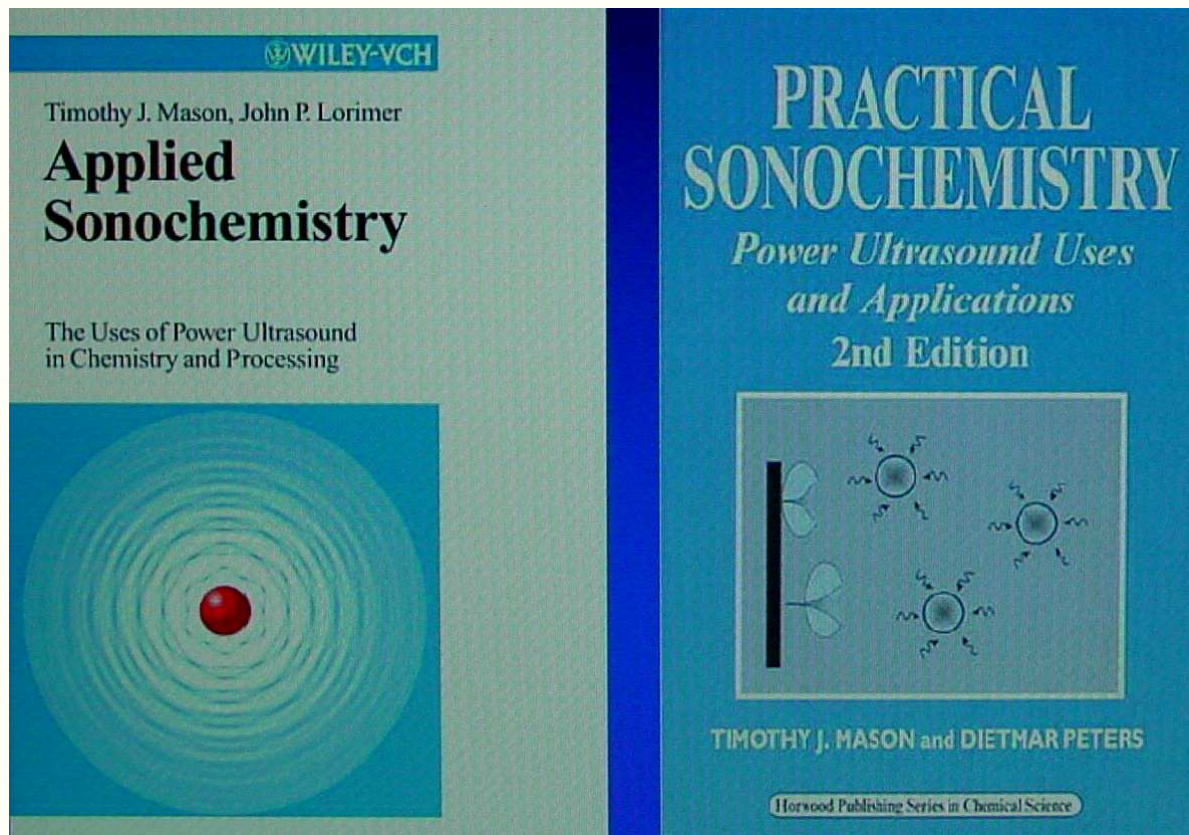
from trace elements to metalloproteins



- **Sound:** propagation of mechanical waves through successive compression and expansion cycles in the medium
- **Ultrasound:** sound presenting frequency higher than 16 kHz
 - **Sonochemistry:** study of ultrasonic waves influence on chemical systems

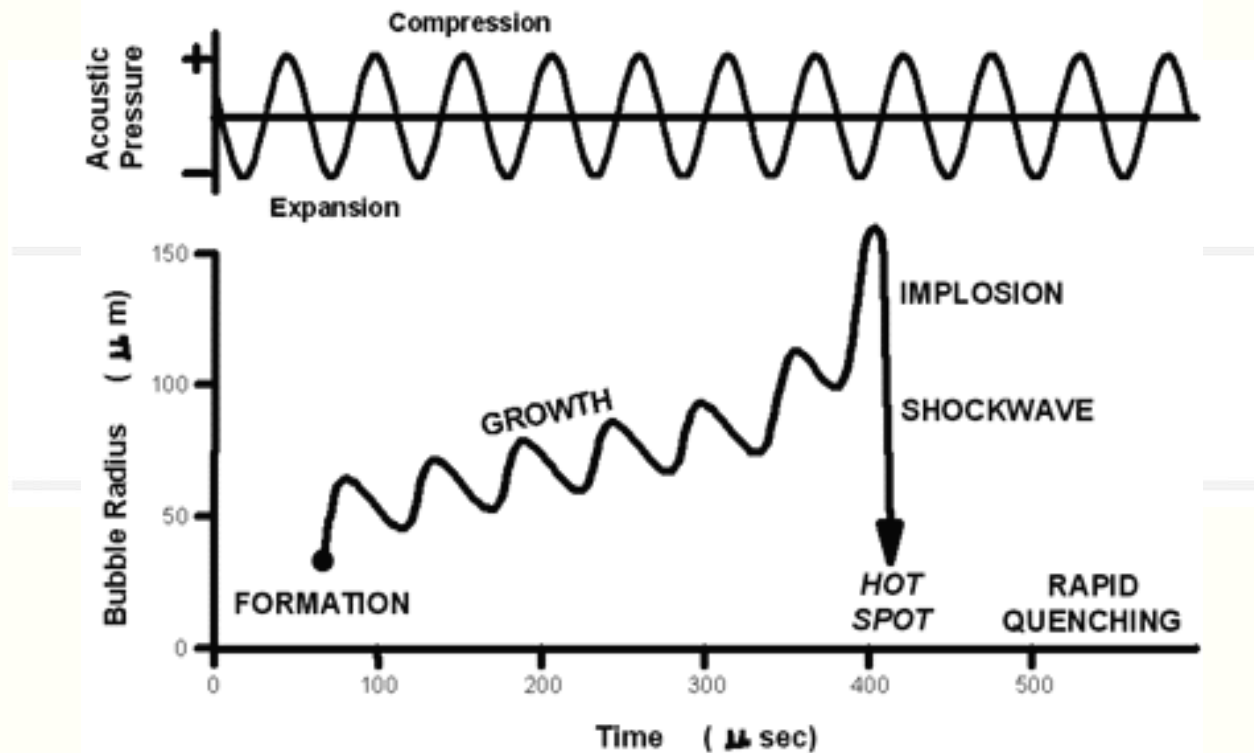


SONOCHEMISTRY



CAVITATION

Process of formation, growth and collapse of micro bubbles in liquids



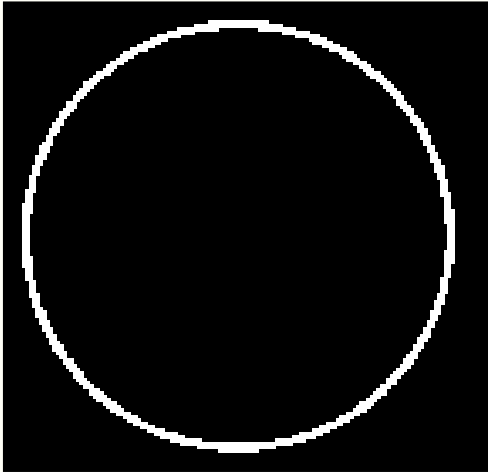
...a way to concentrate the ultrasound energy to a chemically useful form



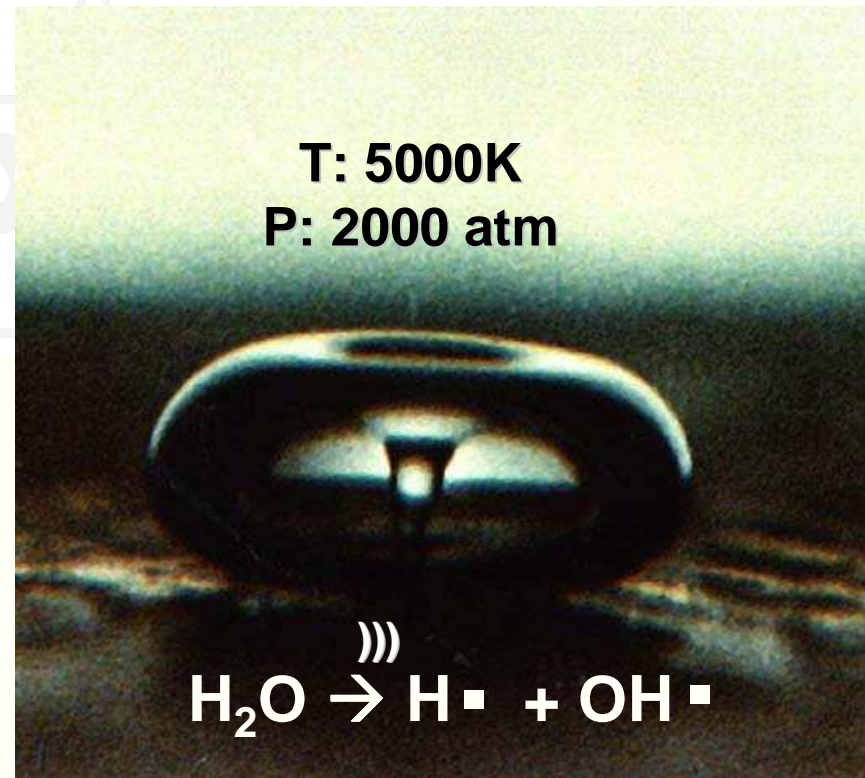
CAVITATION



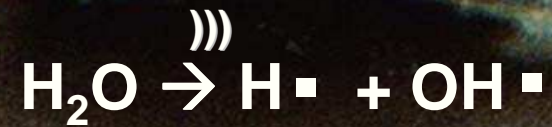
Marco A. Z. Arruda



adapted: www.scs.uiuc.edu/~suslick/britannica.html

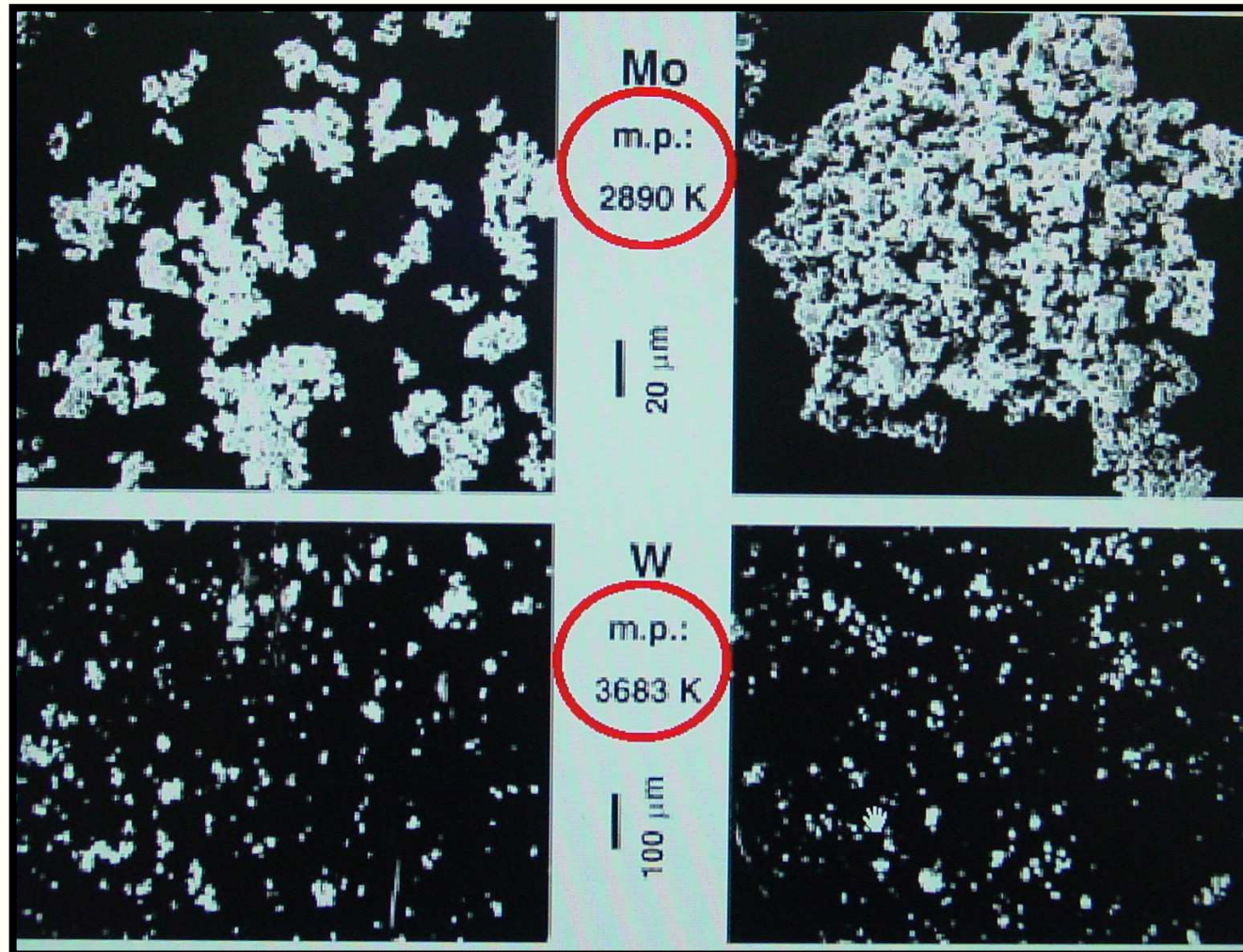


T: 5000K
P: 2000 atm



CAVITATION

Cavitation effect and micro-jet effect:
fusion of Mo particles after sonication



without ultrasound

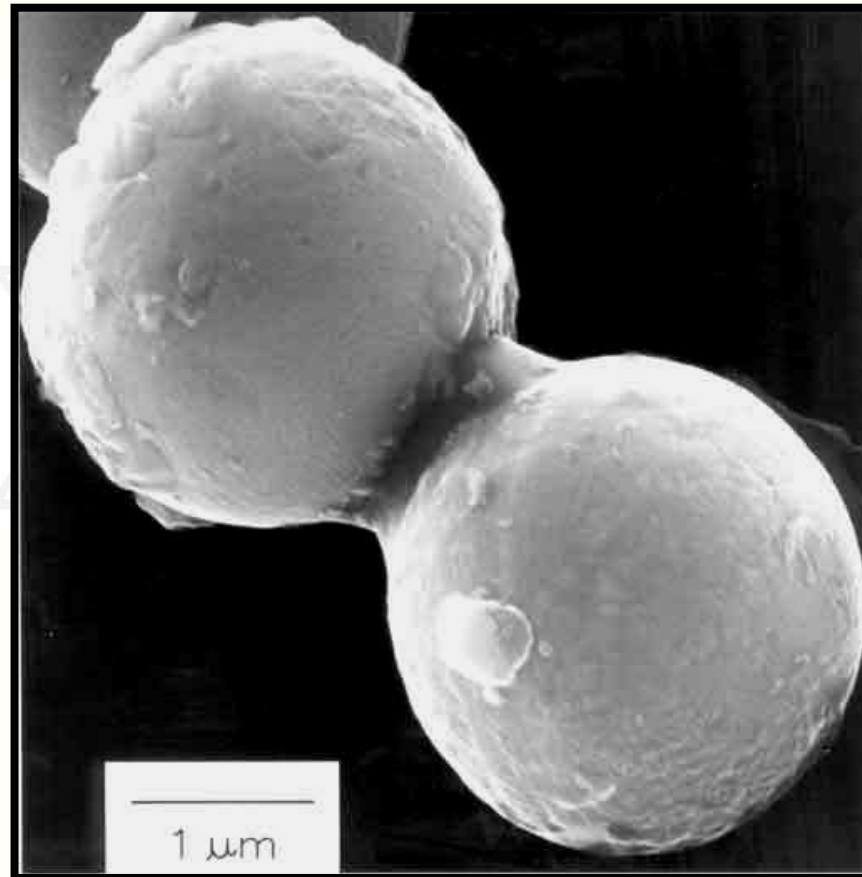
with 30 min ultrasound



CAVITATION

Cavitation effect and micro-jet effect:
fusion of Zn particles after sonication

20 kHz
50 W cm⁻²

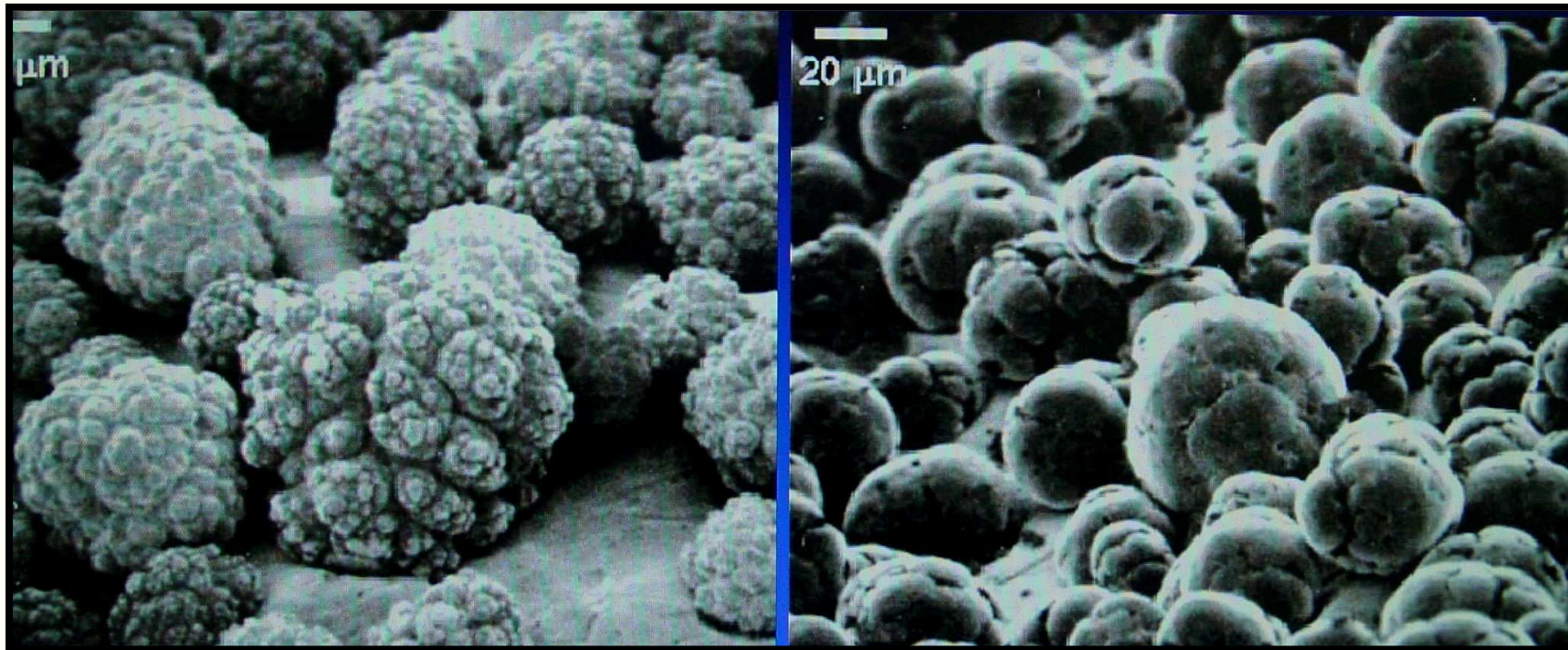


m.p. 419.5 °C



CAVITATION

Cavitation effect and micro-jet effect:
morphology and particle sizes changes → nickel oxide



CAVITATION

Factors affecting

- **Dissolved gas**
- **Irradiation frequency**
- **Temperature**
- **Viscosity and surface tension**
- **External pressure**
- **Presence of particles in solution**

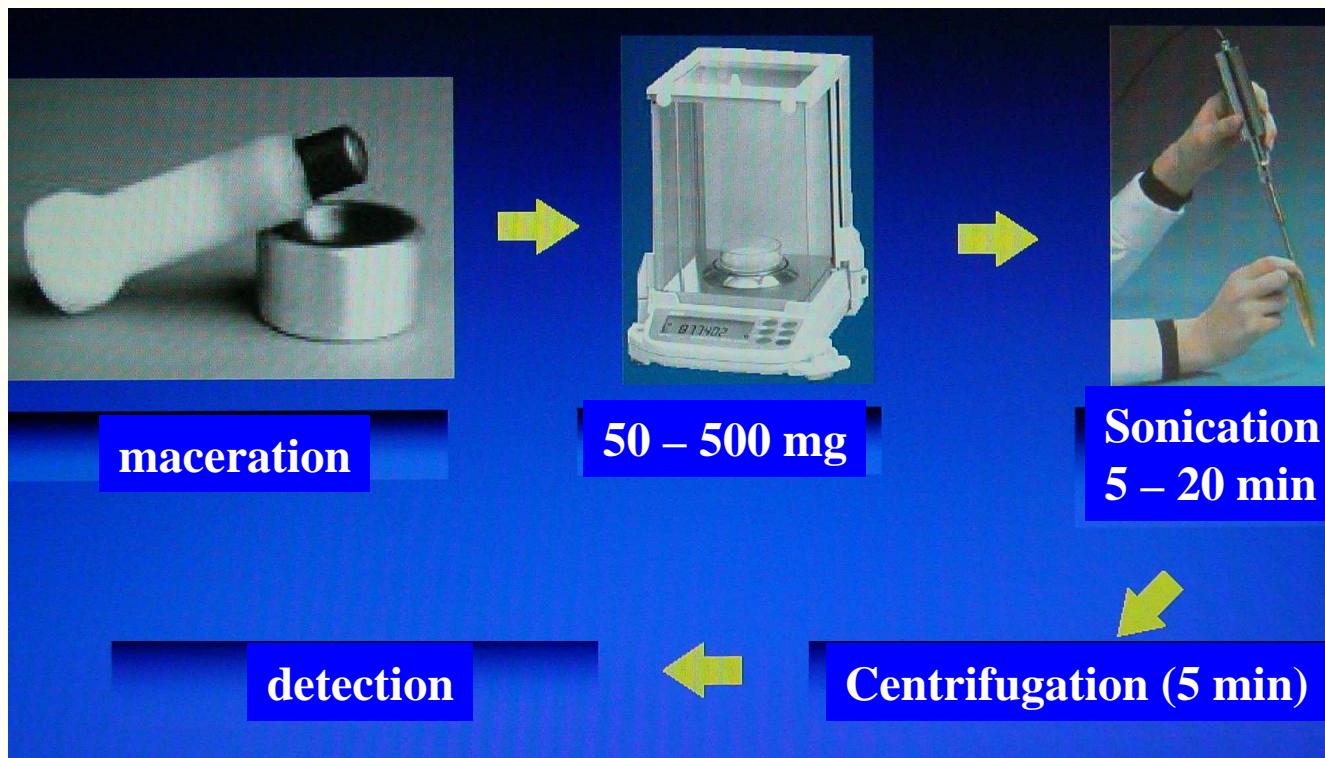


CAVITATION - Sonochemistry

- **Organic synthesis**
- **Polymers degradation**
- **Polymerization**
- **Sonoluminescence**
- **Sonolysis (free-radicals formation)**
- **Sonogels formation**
- **Catalysts preparation**
- **Sample preparation**



CAVITATION - Sonochemistry



CAVITATION - Sonochemistry

Commercial available instruments
ultrasonic probes



The image displays three different configurations of SONICS ultrasonic probes. The leftmost setup shows a control unit with a keypad and a digital display connected to a probe in a flask containing a yellow liquid. The middle setup shows a probe in a flask inside a white sound containment box. The rightmost setup shows a smaller control unit with a digital display and a probe.

500 and 700 W (20 kHz)
250 μ L – 1 L

box for sound
contention

130 W, 20 kHz
150 μ L – 150 mL

 SONICS
Newtown, CT, USA

<http://www.sonicsandmaterials.com>



CAVITATION - Sonochemistry

Commercial available instruments
ultrasonic baths



30 – 80 W
1 – 25 L

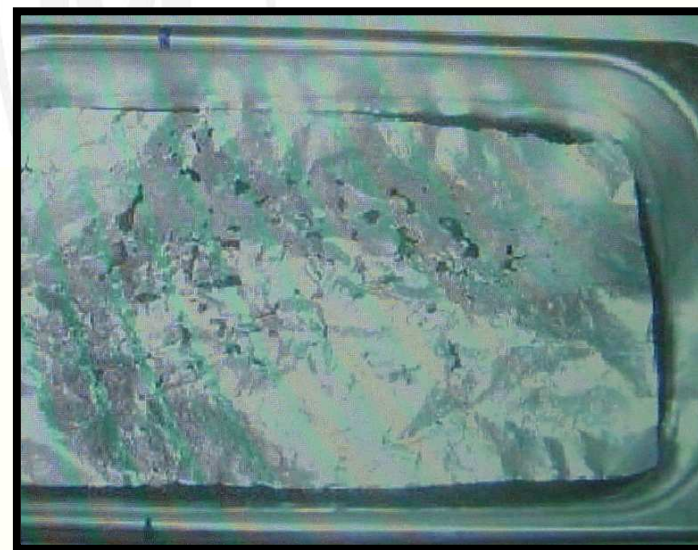
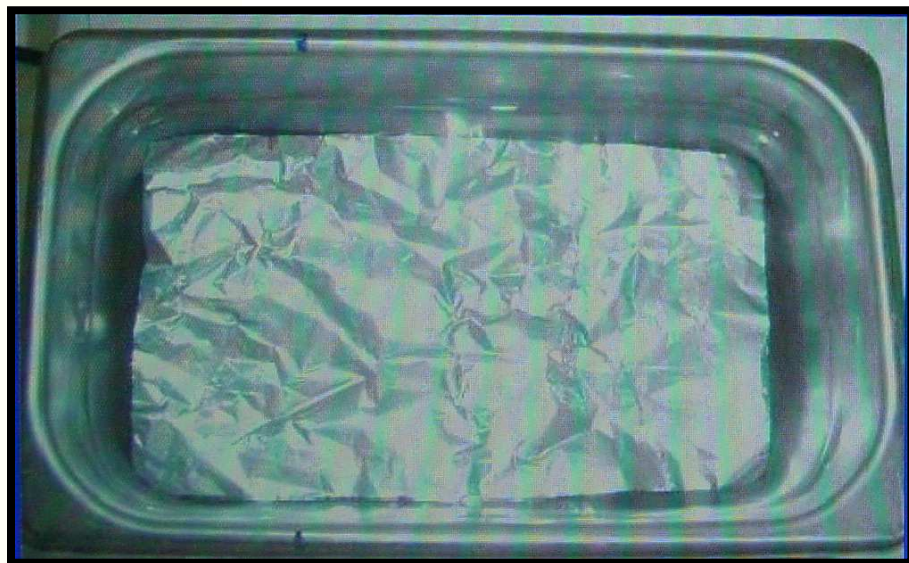


500 W
350 mL



CAVITATION - Sonochemistry

Energy distribution
ultrasonic baths



CAVITATION - Sonochemistry

Energy distribution – ultrasonic baths

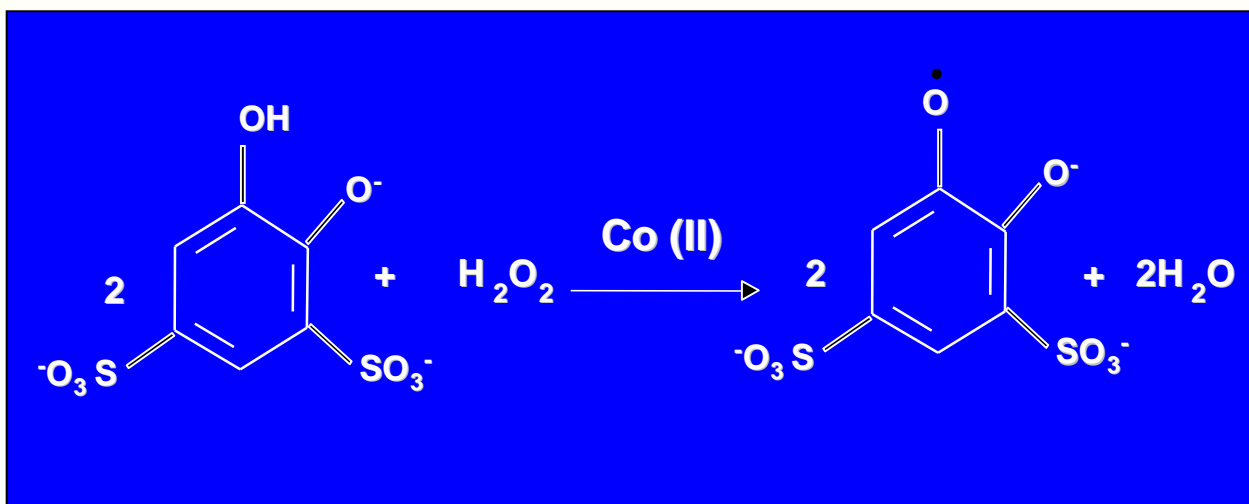
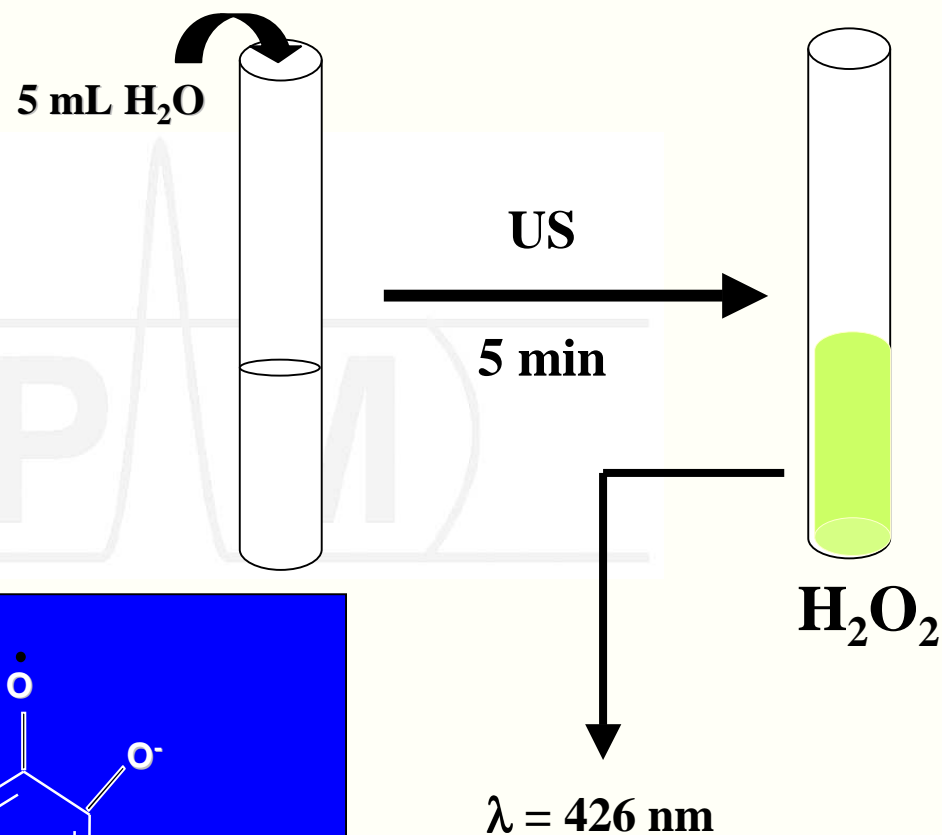
Nascentes *et al.*, *J. Braz. Chem. Soc.*, 12(2001)57

- H_2O_2 method

- H_2O_2 formation

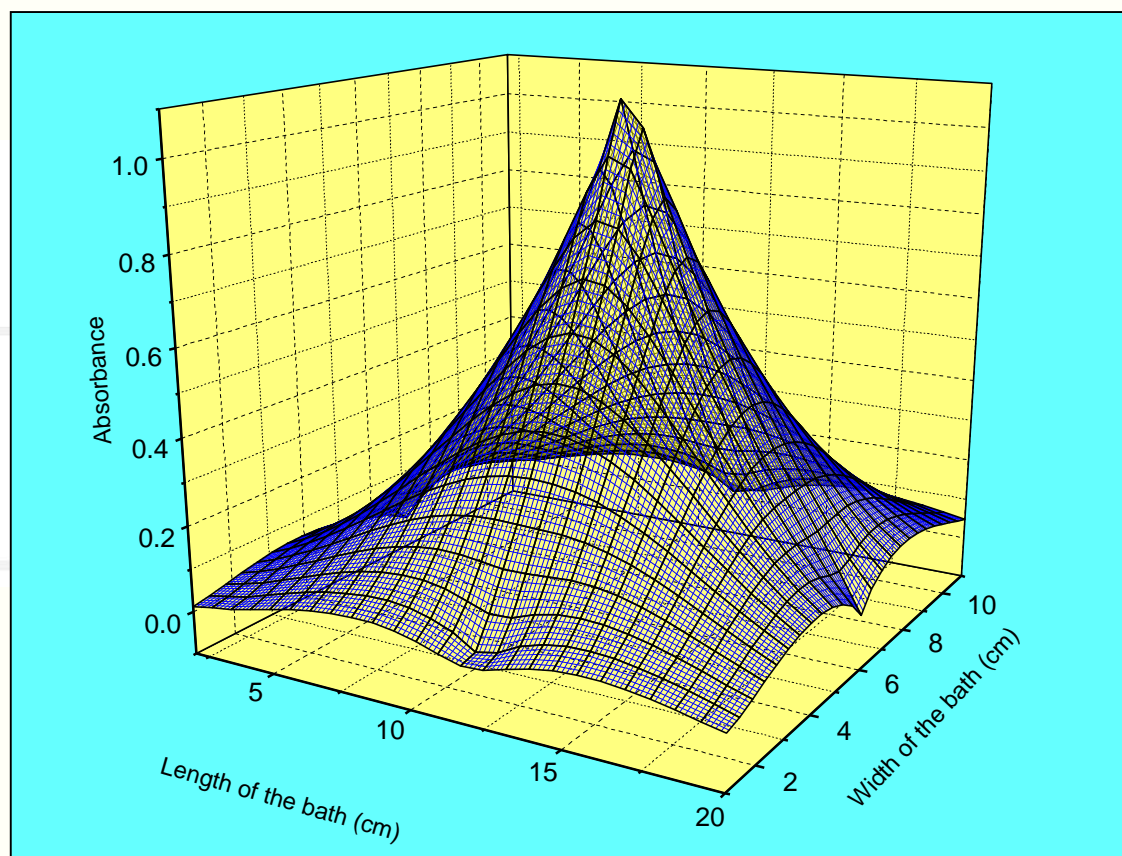


- Tiron oxidation by H_2O_2



CAVITATION - Sonochemistry

Energy distribution – ultrasonic baths



Point 5 → highest H_2O_2 concentration

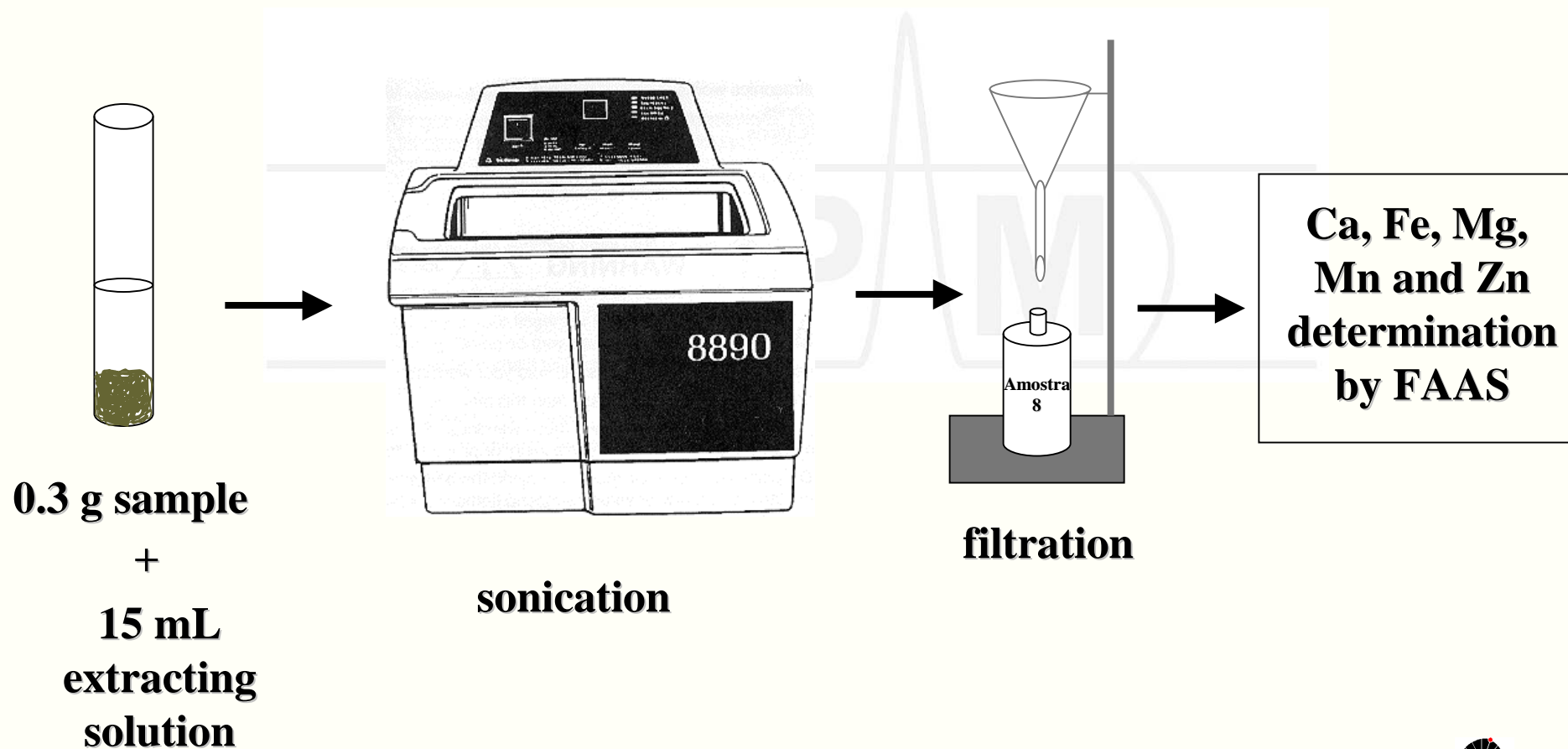
Others: < LOD



CAVITATION - Sonochemistry

Sample preparation

Nascentes et al., *Microchem. J.* 69(2001)37



CAVITATION - Sonochemistry



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Sample preparation

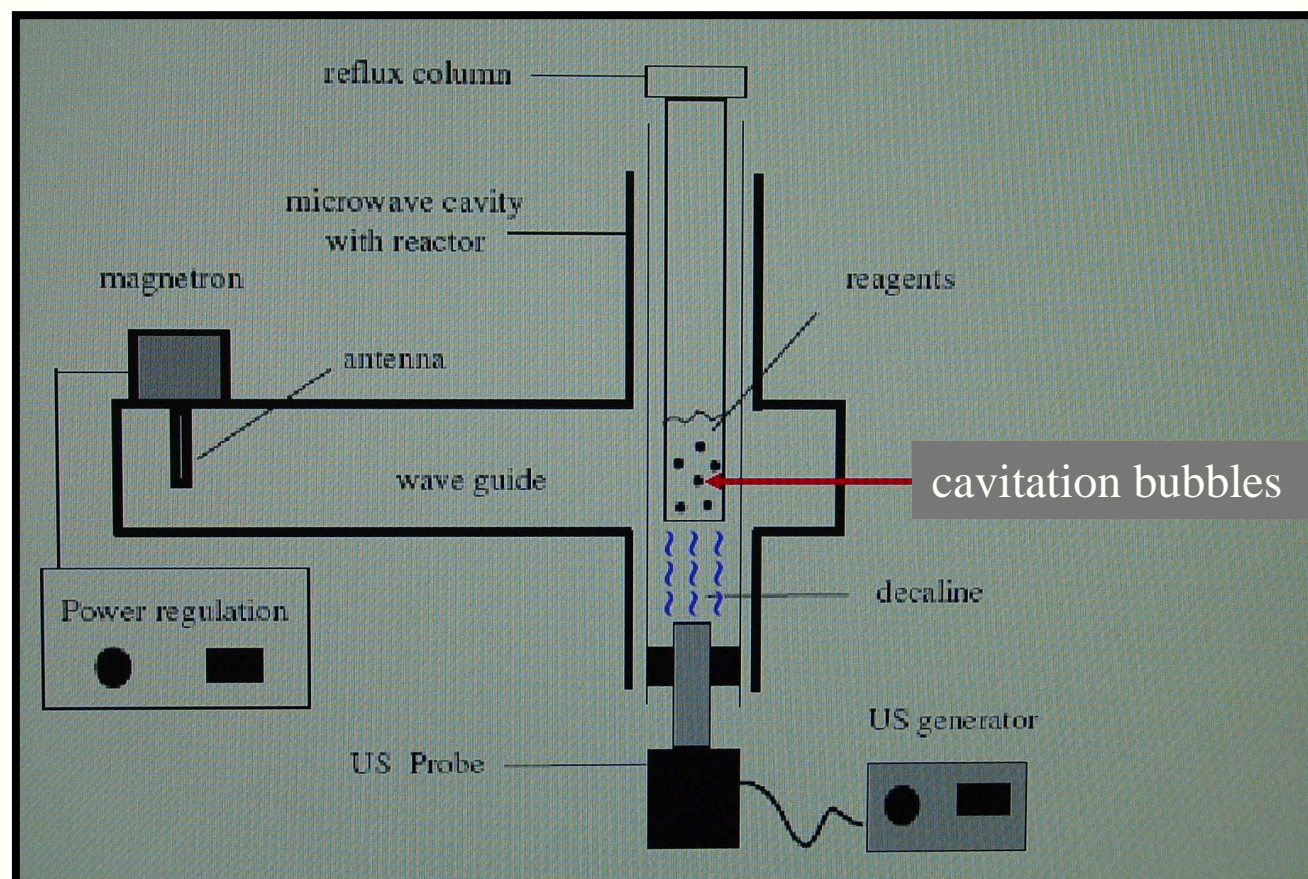
Samples	Wet digestion ($\mu\text{g g}^{-1}$)			
	Ca	Mg	Mn	Zn
Lettuce	1.69±0.06	0.280±0.009	177±7	119±5
Cabbage	0.88±0.05	0.180±0.008	31±1	26±1
Cavalinha	2.64±0.07	0.200±0.005	80±1	36±1
Spinach ^a	---	---	---	---
Apple leaves ^b	---	---	---	---
Ultrasound extraction ($\mu\text{g g}^{-1}$)				
	Ca	Mg	Mn	Zn
Lettuce	1.68±0.03	0.280±0.012	166±2	112±2
Cabbage	0.89±0.04	0.180±0.010	32±1	26±1
Cavalinha	2.68±0.02	0.190±0.008	75±3	33±1
Spinach ^a	1.10±0.05 ^c	0.850±0.022	72±3	74±4
Apple leaves ^b	1.60±0.14	0.250±0.007	53±1	11±1

^aNIST 1570a: 1.53±0.05; 0.9; 79±2; 82±8 $\mu\text{g g}^{-1}$ for Ca, Mg, Mn and Zn
^bNIST 1515: 1.51±0.02; 0.270±0.012; 54±2; 12.5±0.4 $\mu\text{g g}^{-1}$ for Ca, Mg, Mn and Zn
^cno statistical differences at 99.5% confidence level

CAVITATION - Sonochemistry

Sample preparation

Chemat et al., *Ultrasonics Sonochem.* 11(2004)5-8



CAVITATION - Sonochemistry



Marco A. Z. Arruda

Sample preparation

Chemat *et al.*, *Ultrasonics Sonochem.* 11(2004)5-8

Efficiency of simultaneous MW-US irradiation on the digestion time

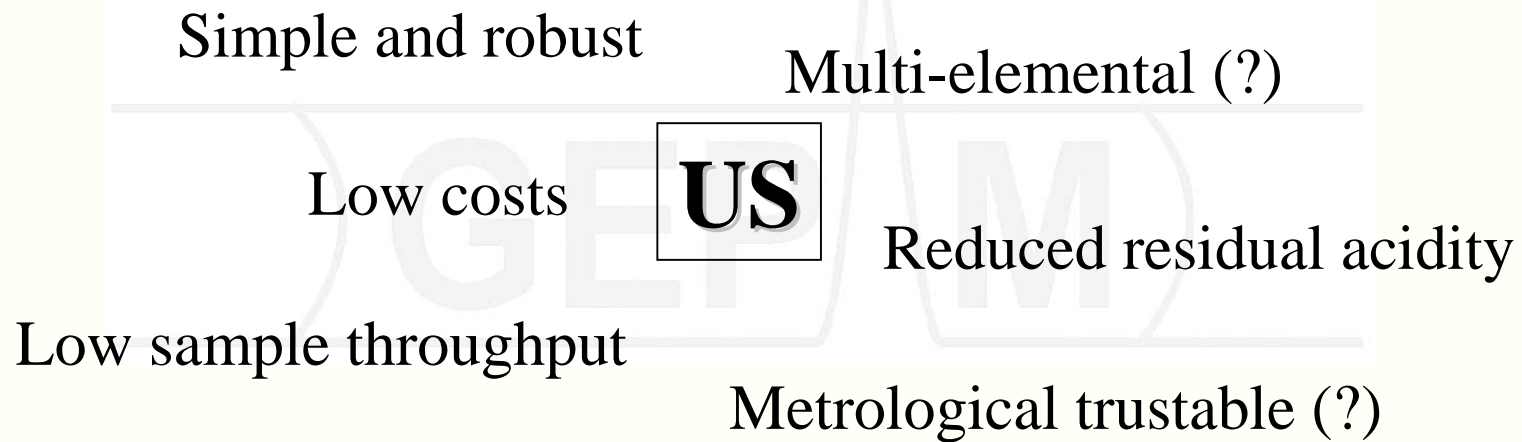
Heating system	Digestion time of sunflower oil (min)	Digestion time of sesame oil (min)
(1) Microwave	40	50
(2) Classical	50	70
(3) Microwave + US	24	35

Comparative Kjeldahl nitrogen determination in food and agricultural products between MW and classical standards methods

Food products	Classical Kjeldahl		MW Kjeldahl		MW-US Kjeldahl	
	%N	t (min)	%N	t (min)	%N	t (min)
Cow's milk	0.5	180	0.48	30	0.45	10
Rice	1.04	180	1.18	30	1.1	10
Corn	1.12	180	1.09	30	1.19	10
Flour	1.57	180	1.72	30	1.62	10
Beef	2.67	180	2.81	30	2.91	10
Corned beef	3.69	180	3.61	30	3.53	10
Chick pea	3.20	180	3.35	30	3.55	10
Powdered milk	4.75	180	4.82	30	4.63	10

Ultrasound – US

Conclusions



Extraction success → interactions between analyte-sample



Microwave assisted water extraction-MAWE

✓ Advantages:

- ✓ No cost and abundant substance
- ✓ Changes in its polarity by changing T
- ✓ Clean extraction methods

✓ Limitations:

- ✓ Low significance as extracting agent when not assisted by auxiliary energies

✓ Alternative extraction when assisted by:

- ✓ High temperature – high pressure
- ✓ Microwave radiation
- ✓ Ultrasound



Microwave assisted water extraction-MAWE

water $\leftarrow \rightarrow$ microwaves interaction

- ✓ High dielectric constant compound necessary for producing heating when MW is used (presence of water!!)
- ✓ $\gggg T \rightarrow <$ viscosity. MW penetrates into sample destroying macrostructure of the matrix
- ✓ T and P produce a decrease on polarity with increases on analytes solubility. Process completely different to conventional extraction \rightarrow solvent diffusion through the sample and analytes are removed from matrix by solubilization



Microwave assisted water extraction-MAWE

water properties

- ✓ Polarity modified by temperature
- ✓ P and room temperature:
 - ✓ Dielectric constant (80) – extremely polar solvent, favoring the solubility of high polarity compounds
- ✓ High T and high P (water in liquid phase)
 - ✓ Lower dielectric constant, favoring the solubility of compounds presenting low polarity
 - ✓ Example: water (at 250°C)/ $\epsilon = 27$
 - ✓ ethanol/ $\epsilon=24$; methanol/ $\epsilon=33$ (at room T)
 - ✓ Water at high T contributes to the non-polar compounds solubility increases: benzo(e)pyrene solubility increases 25 million times (water → from room T to 350°C)
 - ✓ Extraction varying from ionic analytes to non-polar compounds



Microwave assisted water extraction-MAWE

5092

J. Agric. Food Chem. 2001, 49, 5092–5097

Focused Microwave Assistance for Extracting Some Pesticide Residues from Strawberries into Water before Their Determination by SPME/HPLC/DAD

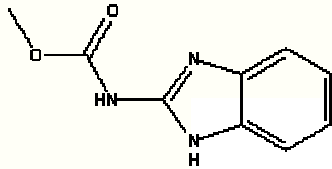
Clara Falqui-Cao, Zhi Wang, Louise Urruty, Jean-Jacques Pommier,[†] and Michel Montury*

Equipe Périgourdine de Chimie Appliquée, Laboratoire de Physico et Toxico Chimie des Systèmes Naturels UMR 5472, Université Bordeaux 1/CNRS, BP 1043, 24001 Périgueux Cedex, France and Centre Interrégional de Recherche et d'Expérimentation de la Fraise, 24130 Prigonrieux, France

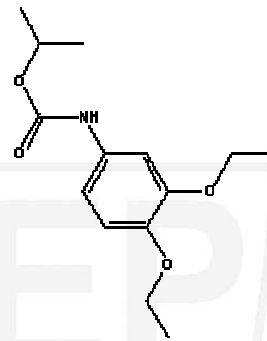
A novel and simple method for the determination of some pesticide residues in strawberries using both focused microwave-assisted extraction (FMAE) and solid-phase micro extraction (SPME), coupled with high-performance liquid chromatography (HPLC), has been developed. The pesticides were first extracted from strawberries with water and the assistance of focused microwaves at 30 W for 7 min. Then, an aliquot of the resulting aqueous extract was subjected to SPME with a 60- μ m thick poly(dimethylsiloxane)/divinylbenzene (PDMS/DVB) fiber for 45 min at room temperature, with the solution being stirred at 1000 rpm. The extracted pesticides on the SPME fiber were desorbed into the SPME/HPLC interface for quantitative analysis with a diode array detector (DAD). The whole sample pretreatment procedure before chromatographic analysis did not use any organic solvents or involve any blending or centrifugation steps. The five compounds (carbendazim, diethofencarb, azoxystrobin, napropamide, and bupirimate) were chosen because they cannot be analyzed easily by GC. The efficiency of this relatively fast procedure was comparable to that of previously reported methods, with detection limits at low μ g/kg levels and linear responses in the range from 0.05 to 1 mg/kg of pesticide in strawberries, with RSDs between 3 and 7.3%, depending on the analyte. In all but one case results obtained by this method for field-incurred samples were comparable to those obtained with traditional methods.



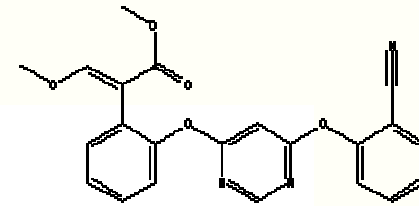
Microwave assisted water extraction-MAWE



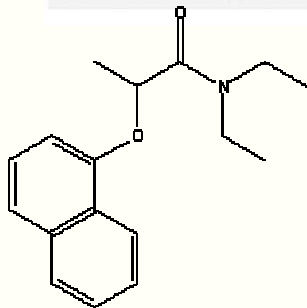
Carbendazim
[C₉H₉N₃O₂]



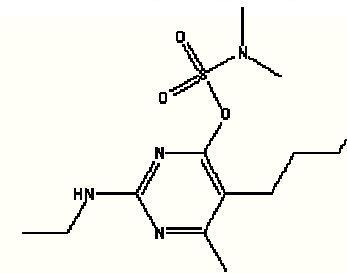
Diethofencarb
[C₁₄H₂₁N₄O₄]



Azoxystrobin
[C₂₂H₁₇N₃O₅]



Napropamide
[C₁₇H₂₁N₁O₂]



Bupirimate
[C₁₃H₂₄N₄O₃S]



Microwave assisted water extraction-MAWE

- ✓ Extraction:

- ✓ FMAE

- ✓ SPME – direct analytes transference to the PDMS/DVB fiber
via pure water

Matrix processing and centrifugation are avoided

- ✓ Quantification: HPLC



Microwave assisted water extraction-MAWE

✓ Samples:

- ✓ synthetic (contamination by adding the analytes)
- ✓ real

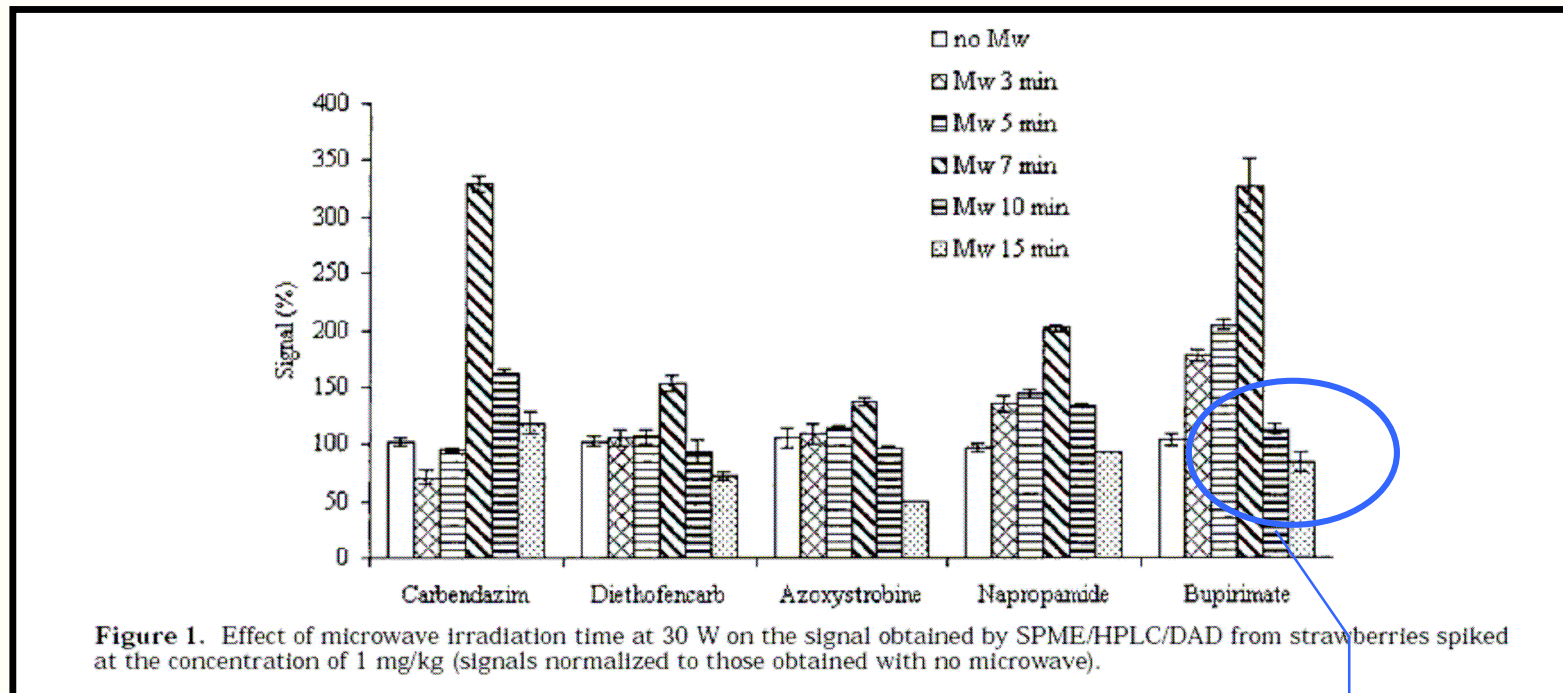
✓ Optimization:

- ✓ SPME – through literature and manufacturer information
- ✓ FMAE – time of fiber exposition (at a fixed power)



Microwave assisted water extraction-MAWE

Extraction efficiency

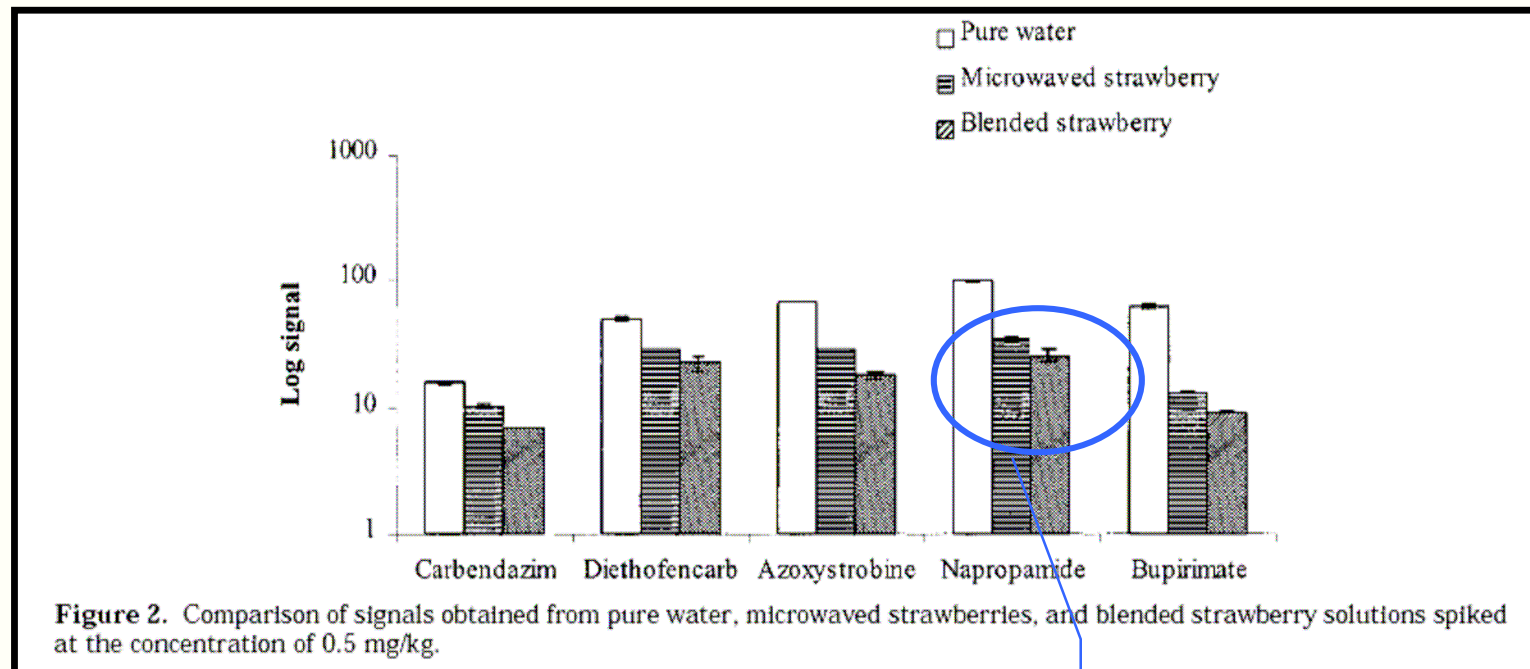


↑ Power
↑ T
→ analytes degradation



Microwave assisted water extraction-MAWE

Recoveries



Matrix effect:

- lower recoveries

Extraction efficiency:

- FMAE ↑



Microwave assisted water extraction-MAWE

Results

pesticide	experimental assay		commercial sample	
	SPME	traditional method	SPME	traditional method
Carbendazim	3.7	3.5		
	1.9	1.7		
	0.04	0.04		
	0.09	0.06		
Diethofencarb	1.8	2.07	0.66	0.73
	0.99	1.22	0.03	0.03
			0.03	0.27
Azoxystrobin	2.9	n.a.		
	0.38	n.a.		
Napropamide	< LOD	< LOD		
Bupirimate	< LOD	< LOD	0.09	0.09
			0.09	0.07
			0.19	0.21
			< LOD	0.03
			0.13	0.15

Table 2. Comparison of Concentrations Obtained by Using SPME and Traditional Methods from Field-Incurred Samples Coming from Experimental Assays and Commercial Production (n.a., not analyzed; LOD, limit of detection)

Table 1. Calibration Curve, Relative Standard Deviation (RSD), Limit of Detection (LOD), Limit of Quantification (LOQ), and French Maximum Residue Limit (MRL) Corresponding to the 5 Pesticides Analyzed

pesticide	λ	regression equation	r	RDS	LOD (mg/kg)	LOQ	MRL (mg/kg)
Carbendazim	205	$y = 1093x$	0.9994	5.8	0.022	0.074	0.100
Diethofencarb	205	$y = 3069x$	0.9977	7.3	0.018	0.060	0.500
Azoxystrobin	205	$y = 3149x$	0.9981	5.8	0.016	0.053	^a
Napropamide	205	$y = 4639x$	0.9964	3.0	0.013	0.067	0.100
Bupirimate	240	$y = 2331x$	0.9977	4.1	0.017	0.044	0.500

^a Registration process for strawberry in progress.



Microwave assisted water extraction-MAWE

Conclusions

No organic solvent for extraction

ϵ similar to organic solvents

→ Water

MAWE

low cost,
abundant,
high purity material

Non-polar compounds extraction

(almost) no matrix modification

Speciation studies??



Microwave assisted water extraction-MAWE

Recommended literature

Morales-Muñoz *et al.*, *Anal. Chim. Acta* 557(2006)278



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Analytica Chimica Acta 557 (2006) 278–286

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Pure and modified water assisted by auxiliary energies: An environmental friendly extractant for sample preparation

S. Morales-Muñoz, J.L. Luque-García, M.D. Luque de Castro *

Department of Analytical Chemistry, Marie Curie Building, Campus of Rabanales, University of Córdoba, E-14071 Córdoba, Spain

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Available online 15 November 2005

Abstract

The growing trend both to avoid the use of organic solvents and reduce the time needed to extract pollutants from environmental solid samples has led to the use of water assisted by auxiliary energies for leaching. The most common of the auxiliary energies used are high pressure–high temperature, microwaves and ultrasounds. One of the most interesting aspects of the use of water in combination with auxiliary energies is the possibility of coupling extraction with other steps of the analytical process, thus enabling partial or total automation of the analytical process, so it is expedited. The addition of reagents (such as surfactants, acids, etc.) to water enlarges its field of application and provides an additional way of shortening the leaching time; thus allowing the establishment of environmental friendly methods. A review about the potential of water as extractant and its main applications so far is here presented.

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Keywords: Water; Auxiliary energies; Sample preparation



Marco A. Z. Arruda

OUTLINE

CPE

→ Cd

→ Proteins

US/MIA/VE

→ Inorganic

→ Organic

MIP

→ Catechol

Miscellaneous

→ Metalloproteins

from trace elements to metalloproteins



Biomimetic Receptors

Molecularly Imprinted Polymers (MIP)

Synthetic polymers presenting selectivity to a target molecule

Science tries to imitate those natural recognition sites

Antigen/**Antibody**

Enzyme/**Substrate**

Drug/**Receptor**



MIPs: Application to the Analytical Chemistry

Samples



Drugs, Enzymes,
Hormones, Proteins,
Amino acids, Sugars,
Agro-toxics, etc.

Techniques



Chromatography
(as stationary phase),
Capillary electrophoresis,
SPE, SPME

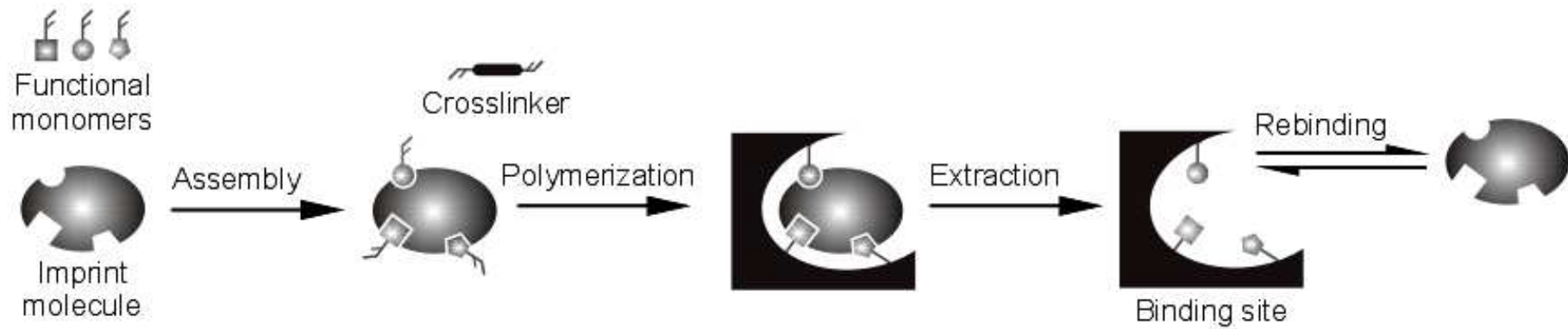


MIP – Synthesis

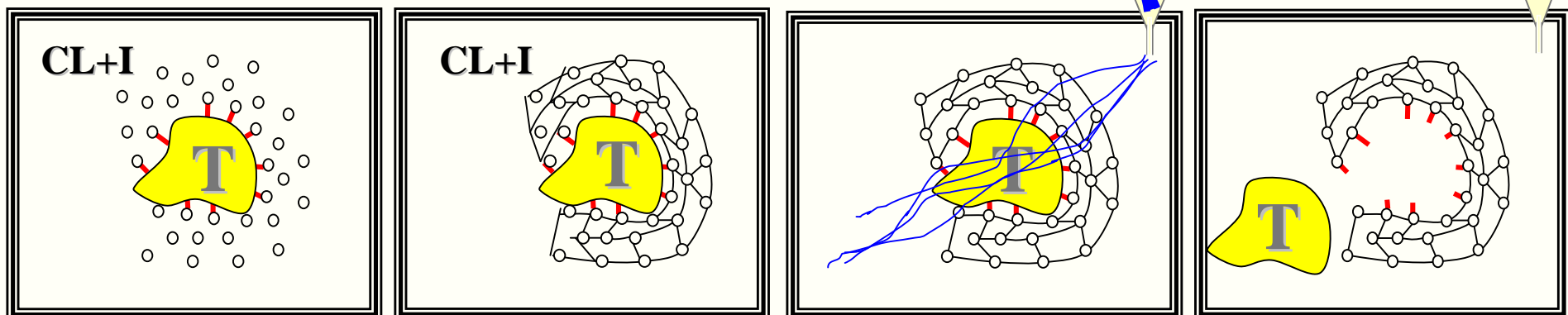


Marco A. Z. Arruda

Medium (solvent, water); Template (IM); Functional monomer (FM); Cross-linker (CL); Initiator (I)



ethanol/acetic acid



MIP – Reagents

Imprinted molecule – IM

- has specific groups to promote the bond to the FM
- no presence of polymerizable groups (insertion to the polymeric net)
- no groups that increase or decrease the polymerization reaction (ex. tyol groups)
- stability (T *ca.* 60°C)



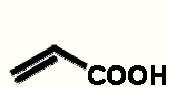
MIP – Reagents

Functional monomer – FM

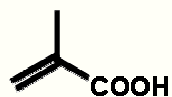
- Complementarities between FM and IM.
- if FM is proton donor (acidic character) → IM must be proton acceptor (basic character) and *vice-versa*
- Commonly used: methacrylic acid (for basic IM) and 4-vinylpyridine (for acidic IM)
- Concentration: interaction between FM and IM is based on equilibrium processes. \gg [FM] than [IM] - *ca.* 4:1 [FM]:[IM] → formation of a great number of specific recognizing sites
- Possibility of combining different FM → polymerization cocktail



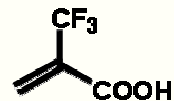
MIP – Reagents



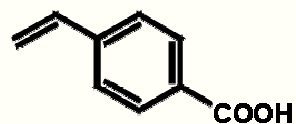
acrylic acid



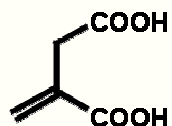
methacrylic acid



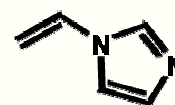
trifluoro-methacrylic acid



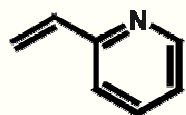
4-vinylbenzoic acid



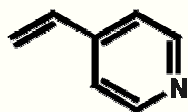
itaconic acid



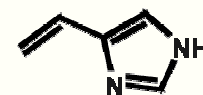
1-vinylimidazole



2-vinylpyridine



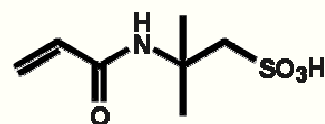
4-vinylpyridine



4(5)-vinylimidazole



4-vinylbenzyl-iminodiacetic acid



**2-acrylamido-2-methyl-
1-propane sulphonic acid**

FM



MIP – Reagents

Cross-linker – CL

-Function:

- *morphology control of the polymeric matrix

- *stabilizes the bond sites

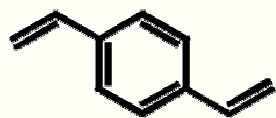
- *promote mechanical stability to the polymer

- necessary at higher proportions in the MIP synthesis → to access the porous of the polymer, guarantying mechanical stability

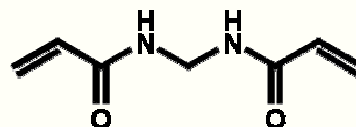
- Ethylene glycol dimethacrylate → commonly used



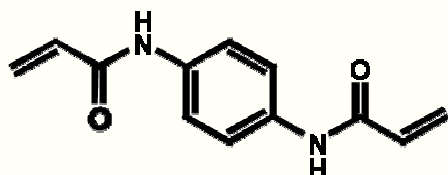
MIP – Reagents



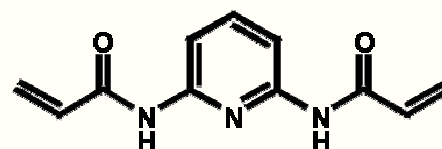
4-divinylbenzene



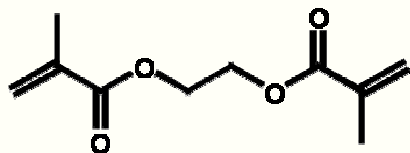
N,N'-methylene-bisacrylamide



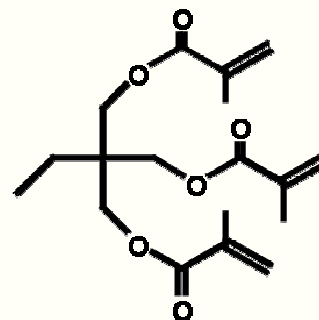
N,N'-phenylene-bisacrylamide



2,6-bisacrylamidopyridine



ethylene glycol dimethacrylate



trimethylolpropane
trimethacrylate

Cross-linker



MIP – Reagents

Solvent

-Function:

- * reagents solubilization in the synthesis

- * contribute to best porous formation

- * contribute to macro-porous formation in the polymer →

large surface area

-does not interfere in the complex FM-IM formation (probability to form low selective sites and in less quantities)

-majority of the interactions between FM-IM is due to the electrostatic forces and hydrogen bounds → solvents should present non polar characteristics and have lower dielectric constant (e.g. chloroform and toluene)

-solubility problems: uses acetonitrile. MIP presenting lower selectivity can be formed



MIP – Reagents

Initiator - I

- Function: creates free radicals for polymerization
- external stimulus: T, UV radiation
- check its influence on others reagents (T increase without its degradation?, etc.)
- commonly used: 2,2'-azobisisobutyronitrile

Caution:

- synthesis processed at 60°C as maximum T (azobisisobutyronitrile)
- it must be carried out in an oxygen-free atmosphere



MIP – Synthesis: Reagents

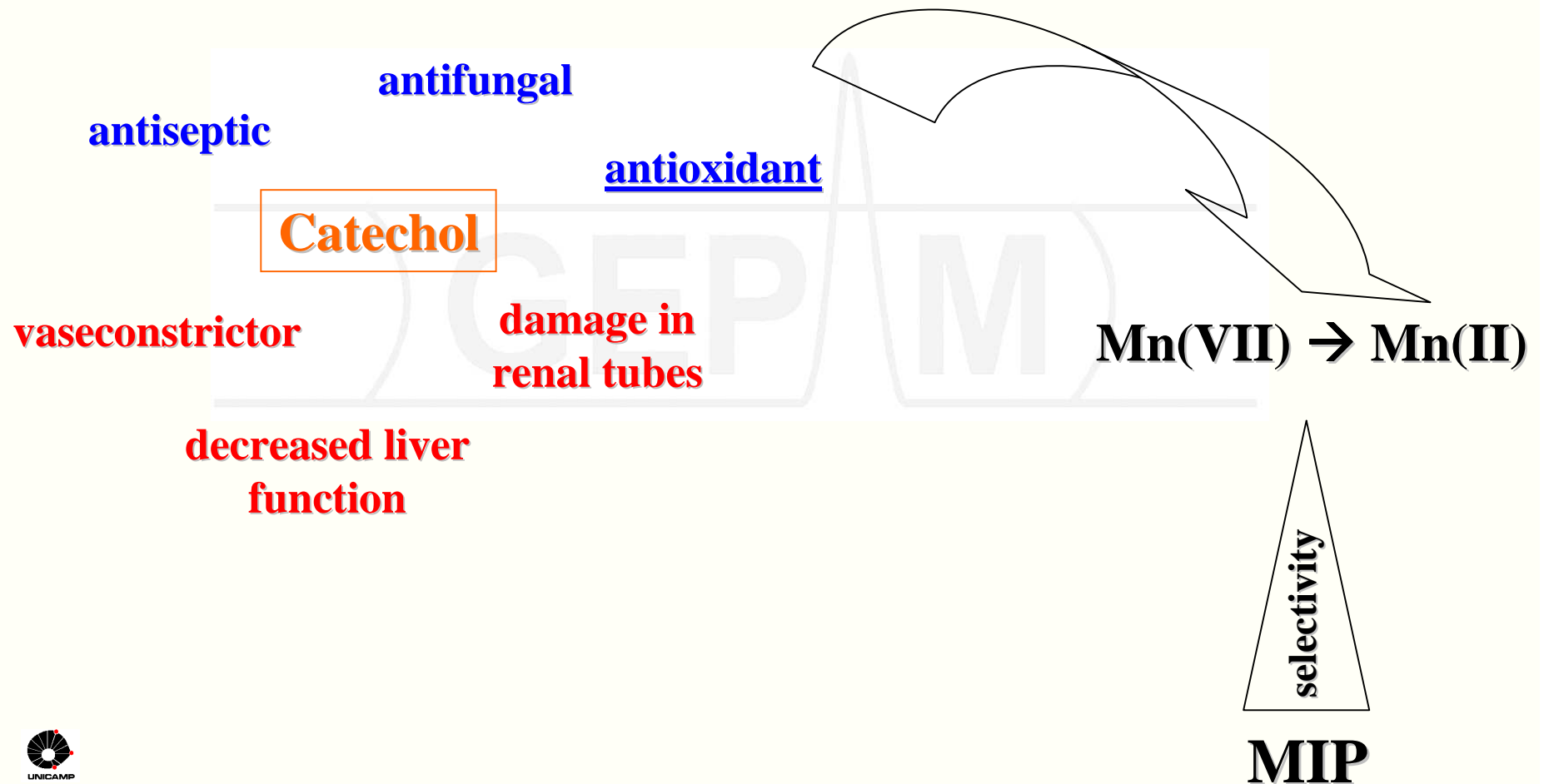
Medium	Chloroform/acetonitrile
Functional monomer	Methacrylic acid (MA) used for basic molecules 4 vinylpyridine (4VP) used for acidic molecules
Cross-linker	Ethylene glycol dimethacrylate (EGDA).
Initiator	Azobisisobutyronitrile



MIP – Applications

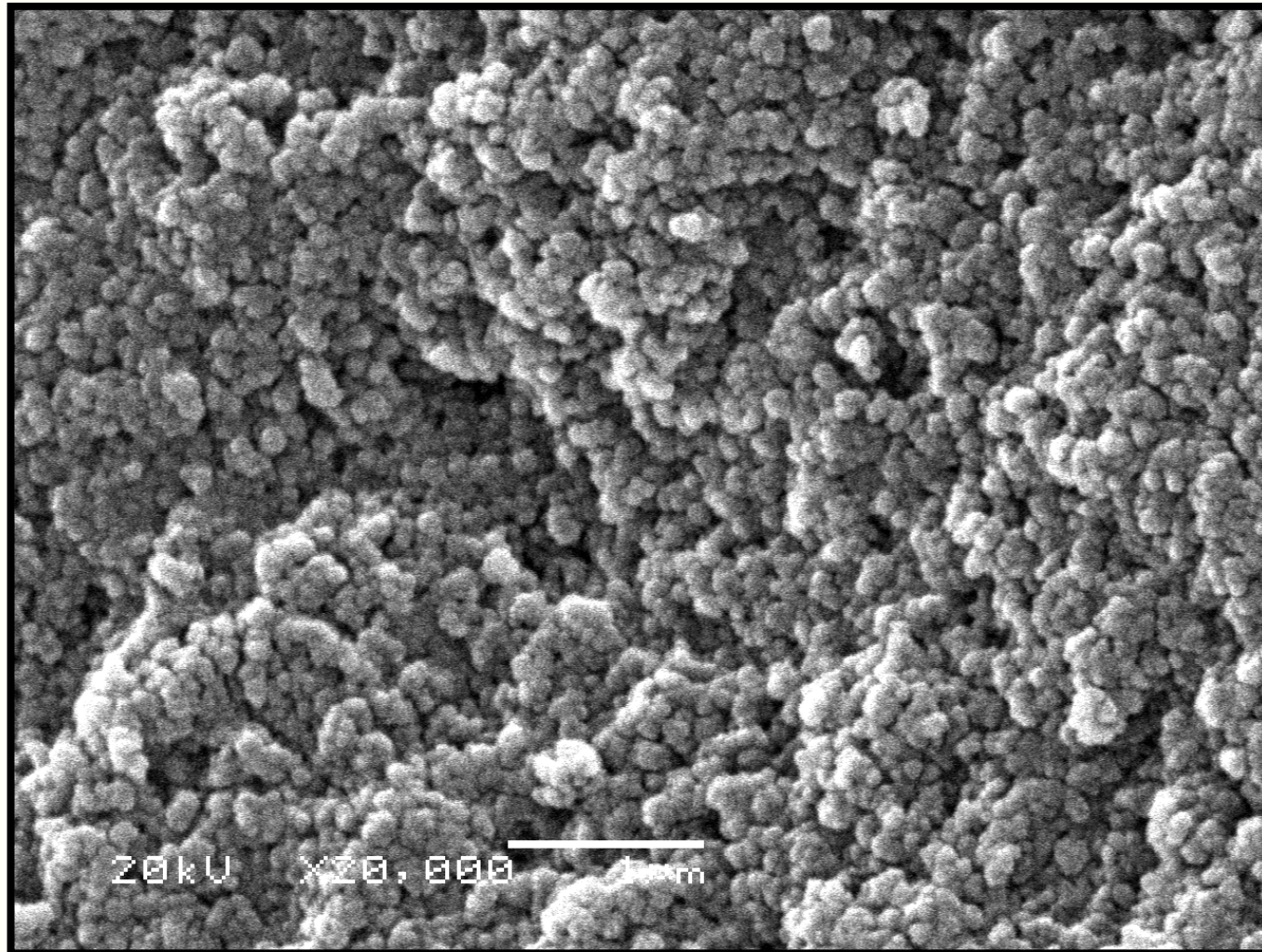
On-line molecularly imprinted solid-phase extraction for the selective spectrophotometric determination of catechol

Figueiredo *et al.*, *Microchem. J.* 85(2007)290



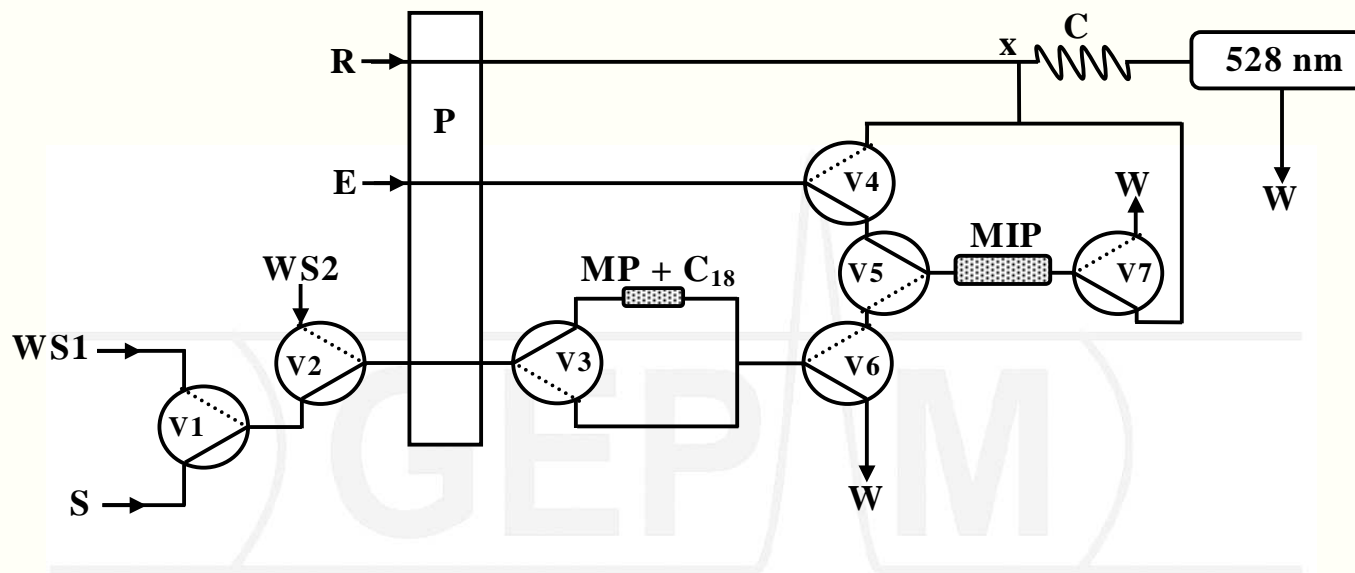
MIP – characterization

Tarley et al., *Talanta* 69(2006)259



MIP – Applications

Flow system



R: KMnO_4 0.010% (m/v)

E: HNO_3 1 mol/L

WS1: washing solution (0.01 mol/L HNO_3 + 2 % v/v ACN)

WS2: washing solution (ACN 50% v/v)

MP+C₁₈: 70 mg (70% polymer + 30% C₁₈)

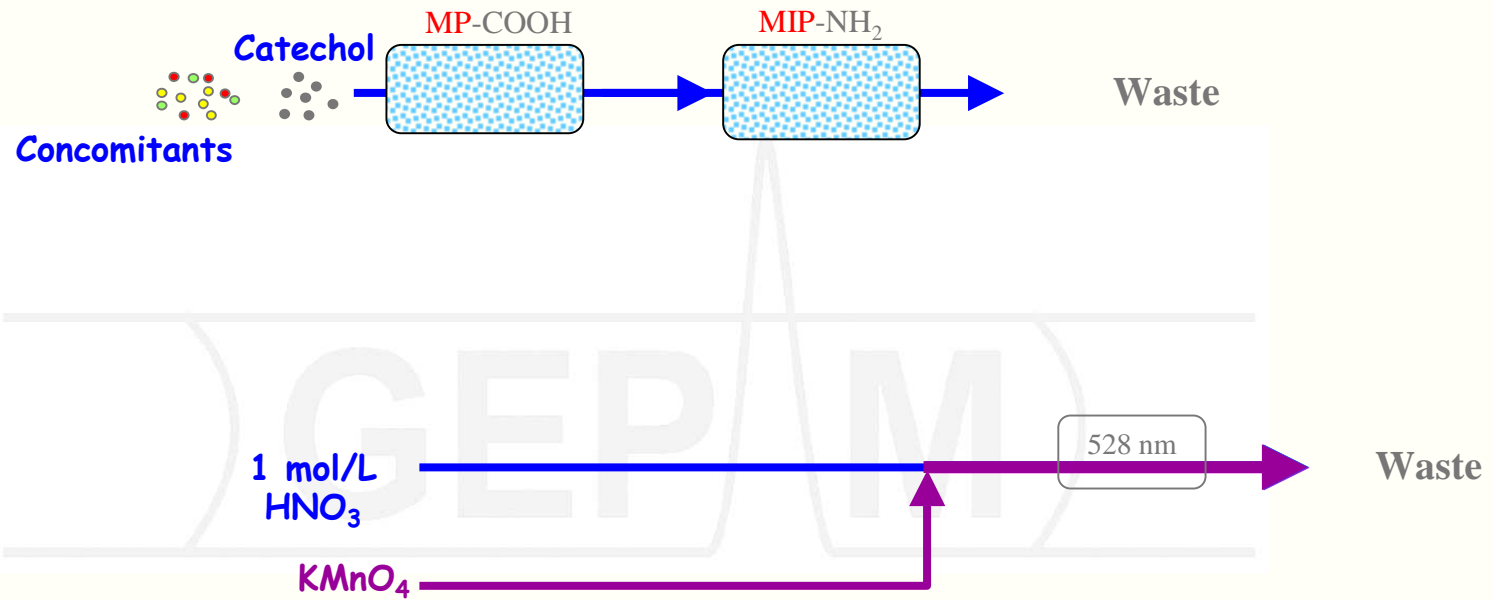
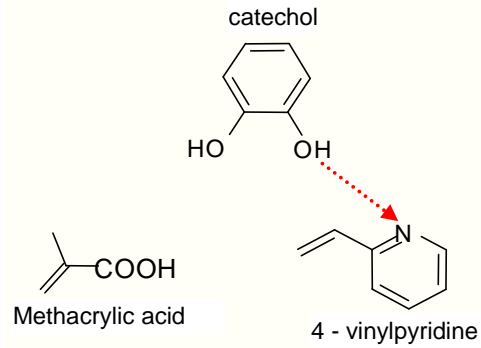
MIP: 70 mg

C: coil 20 cm



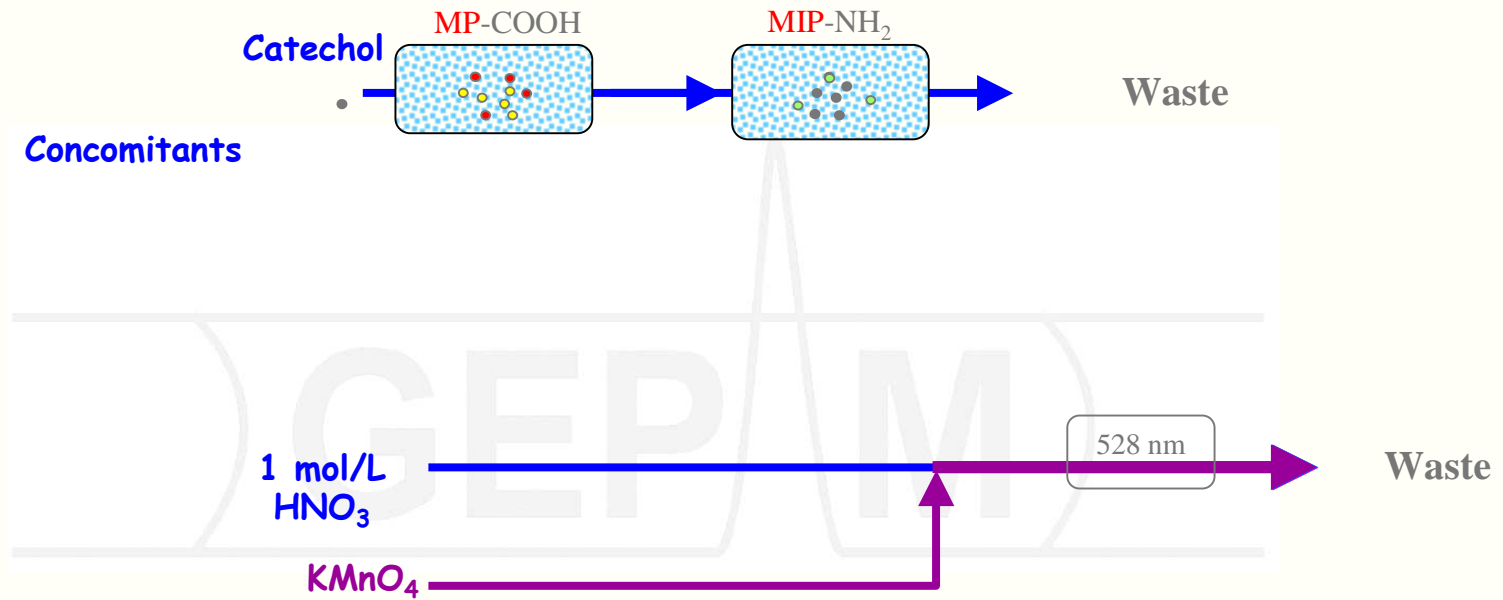
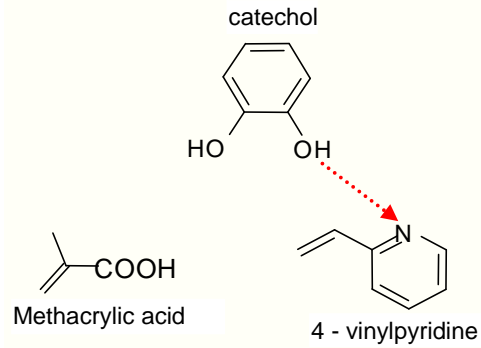
MIP – Applications

Flow system



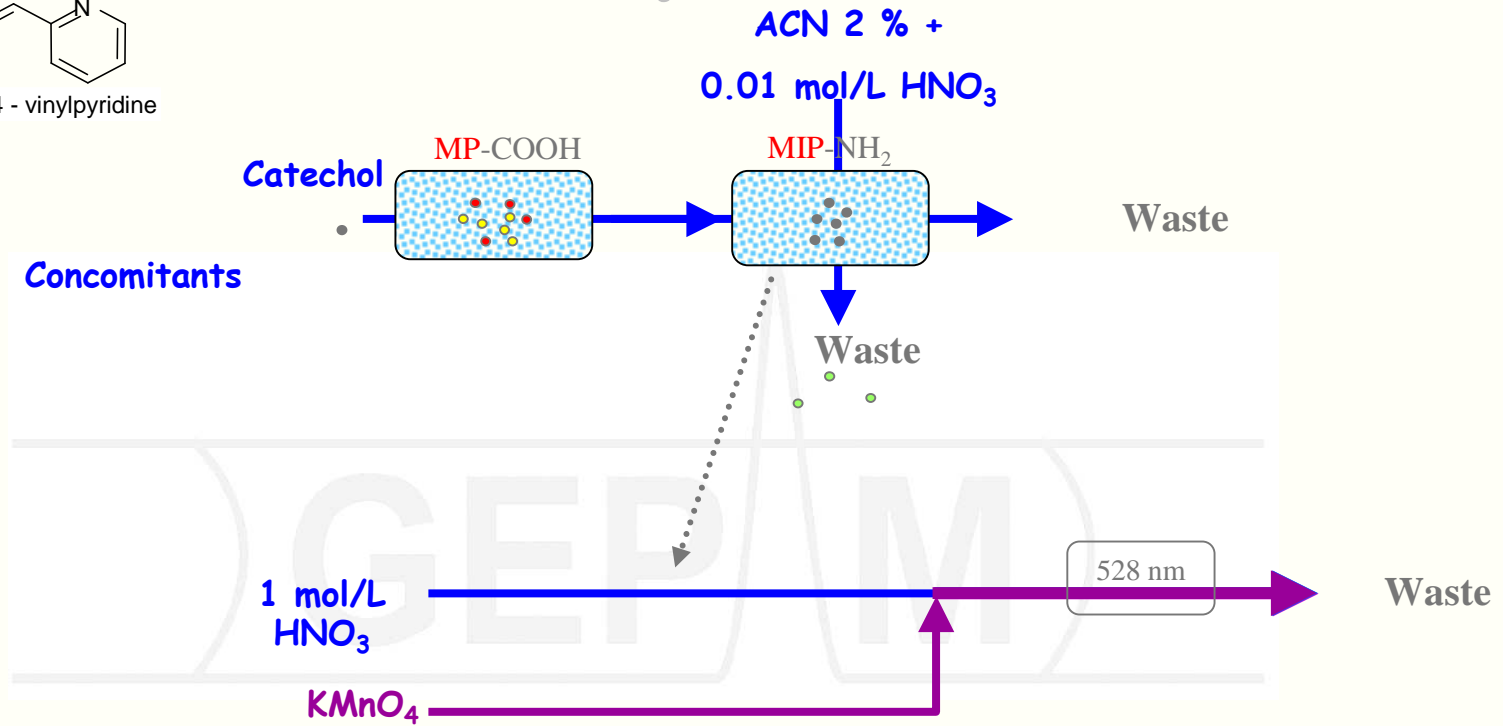
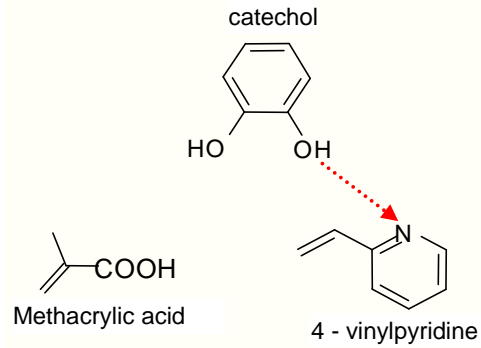
MIP – Applications

Flow system



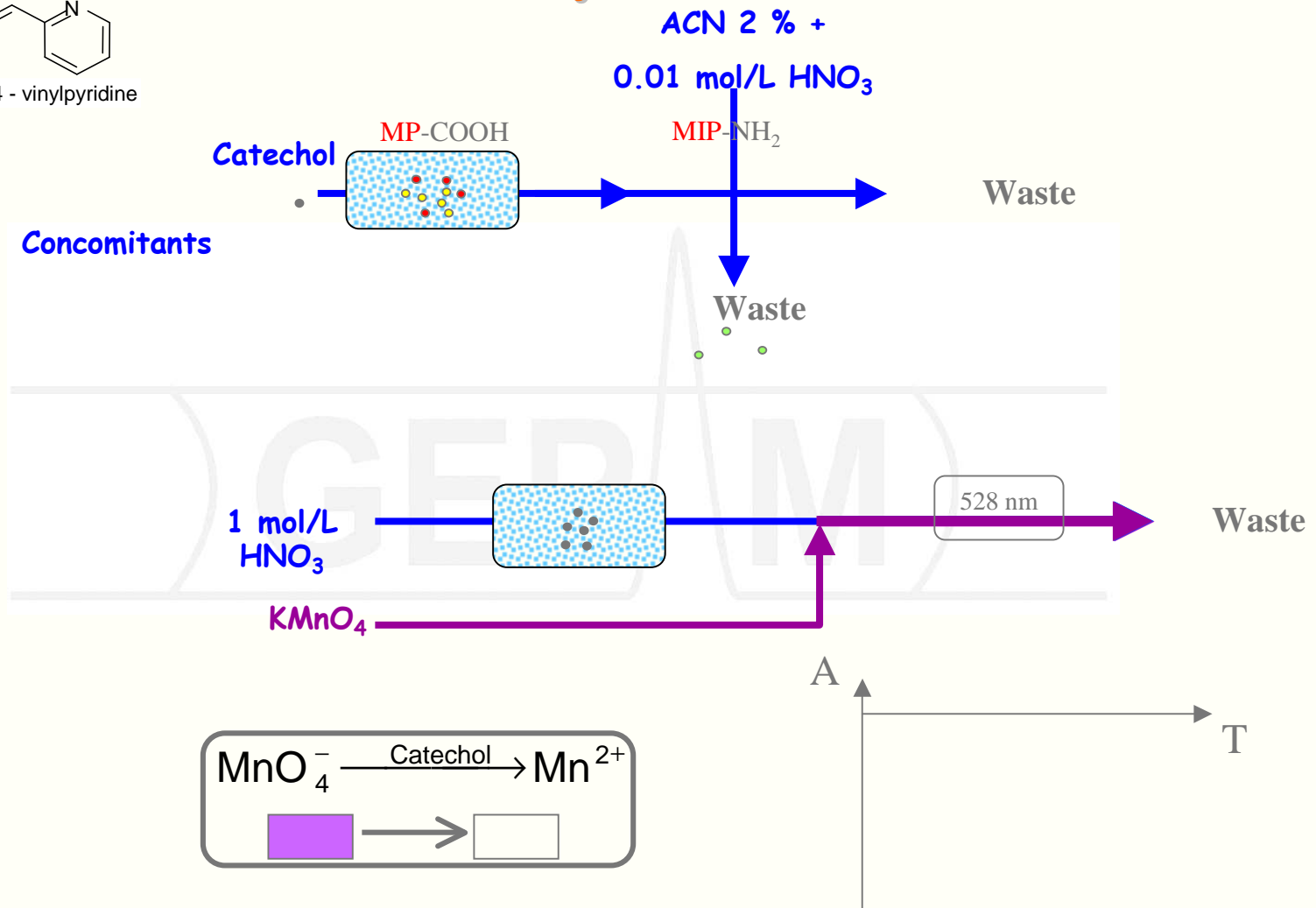
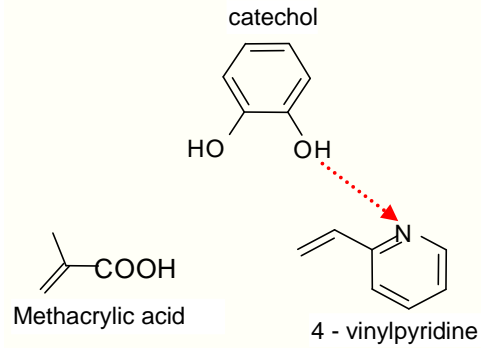
MIP – Applications

Flow system



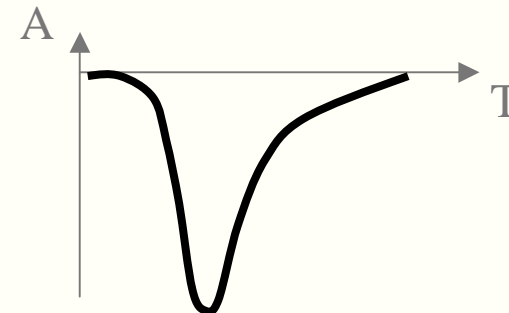
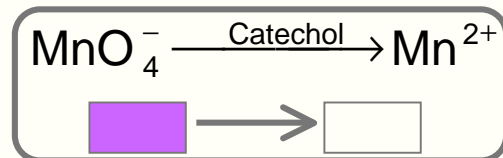
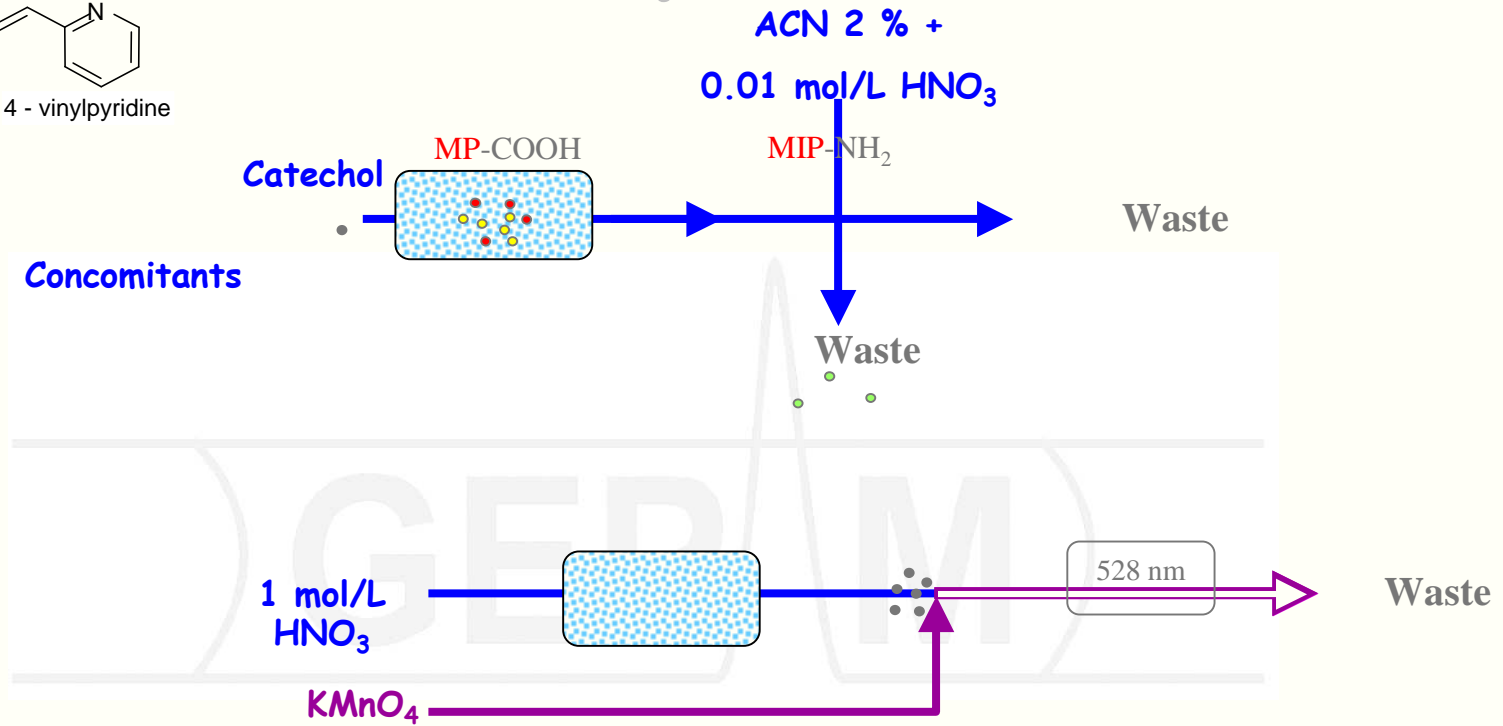
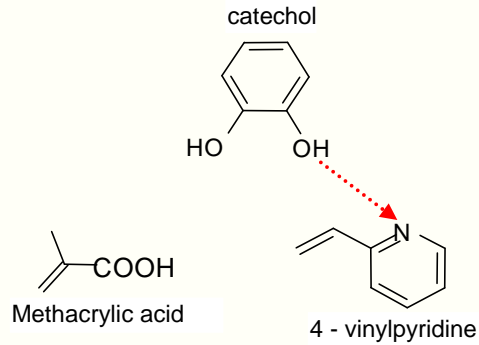
MIP – Applications

Flow system

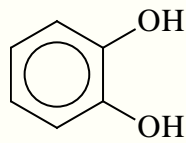


MIP – Applications

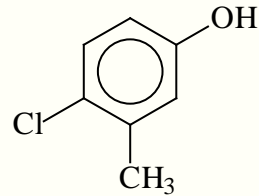
Flow system



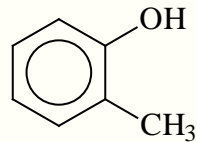
Concomitant evaluation



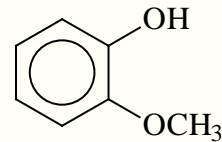
Catechol



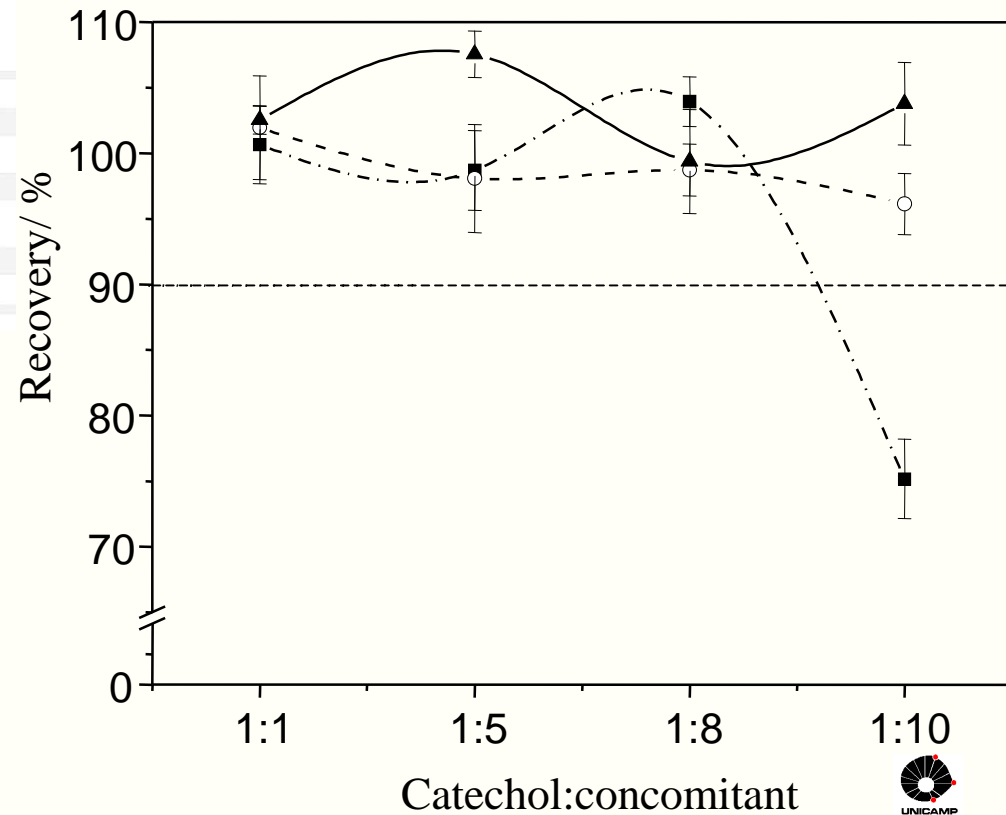
4-chloro-3-methylphenol



2-cresol



2-methoxyphenol



Figures of merit

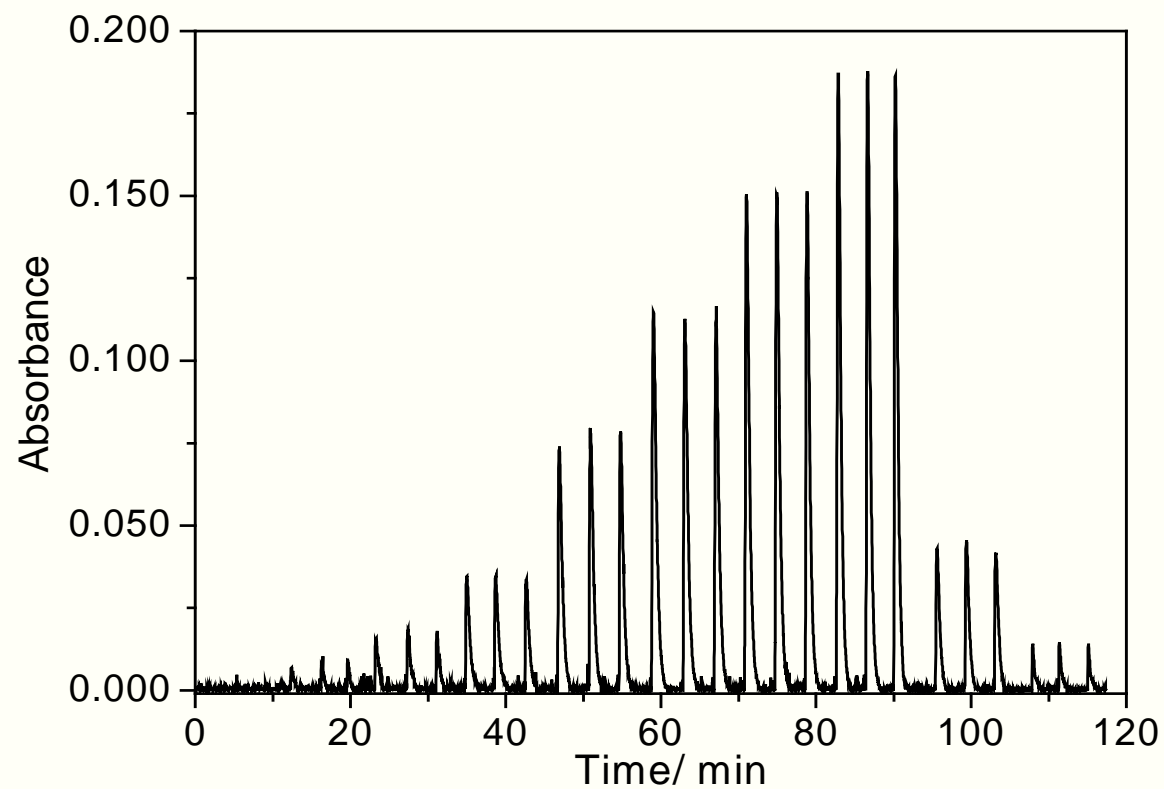
Linear range: 3.00 – 100 $\mu\text{mol/L}$

($r > 0.999$; $n=7$)

Precision: 3% RSD ($n=10$)

LOQ: 2.7 $\mu\text{mol/L}$

LOD: 0.8 $\mu\text{mol/L}$



Results

Sample	Addition ($\mu\text{mol L}^{-1}$)	FI-MISPE	FI-MISPE recovery/%	HPLC	
Guaraná	A	0	< LOQ ¹	-	< LOQ ²
		10	10.5 \pm 0.5	105	10.4 \pm 0.5
		20	20.6 \pm 0.6	103	21.3 \pm 0.2
	B	0	2.8 \pm 0.4	-	< LOQ
		10	11.9 \pm 0.2	90	12.0 \pm 0.1
		20	23.2 \pm 0.4	102	24.0 \pm 0.3
Mate tea	0	7.7 \pm 0.5	-	7.4 \pm 0.2	
	10	18.9 \pm 0.2	112	17.0 \pm 0.1	
	20	29.5 \pm 0.4	109	27.4 \pm 0.2	
Tap water	0	< LOQ	-	< LOQ	
	10	9.5 \pm 0.4	95	10.3 \pm 0.8	
	20	19.6 \pm 0.2	98	20.0 \pm 0.6	
¹ LOQ = 2.7 $\mu\text{mol L}^{-1}$ y ² LOQ = 4.2 $\mu\text{mol L}^{-1}$					



Molecularly imprinted polymers – MIP

Conclusions

Alternative for separating
those compounds presenting
similar structure

Good analytical performance

MIP

Use of non-selective
spectrophotometric
reactions



OUTLINE

CPE

→ Cd

→ Proteins

US/MIA/VE

→ Inorganic

→ Organic

IVFP

→ Catechol

from trace elements to metalloproteins

Miscellaneous

→ Metalloproteins



Some definitions

METALLOME: Set of metals or metalloids inside a cell or tissue – *ionome* and *metalloproteome*

Ionome: Metal (free) contained in a cell

Metalloproteome: Set of complexed metals with proteins in a cell

METALLOMICS: Study (quali and quantitative) of the metallome

HARAGUCHI, 2002



Examples of metalloenzymes (and metalloproteins)

Element	Enzymes/Proteins
Selenium	<i>Glutathione peroxidase</i> . An enzyme that catalyzes the reduction of peroxides and protects the cells from oxidative damage
Chromium	<i>Transferrin</i> (plasma protein). Transports Cr(III) throughout the body in the blood cells
Copper	<i>Ceruloplasmin</i> (human serum protein), <i>ascorbate oxidase</i> (plants and bacteria), <i>plastocyanin</i> (higher plants and cyanobacteria), <i>superoxide dismutase</i> , <i>tyrosinase</i> , <i>cytochrome oxidase</i> and <i>hemocuprein</i> (animals).
Lead	Around 95% of total Lead in human blood is bound to erythrocytes. In most erythrocytes, the lead is bound within the cell in <i>haemoglobin</i> . In extracellular fluids lead is bound to <i>albumin</i> and some high molecular weight proteins (<i>globulins</i>).
Zinc	Zinc is a constituent in more than 200 enzymes and proteins. The principal examples are <i>insulin</i> and <i>carboxypeptidase A</i> .
Manganese	This metal is found in a variety of enzymes such as <i>pyruvate carboxylase</i> and <i>oxalacetate decarboxylase</i> . It can be found in proteins such as <i>glutamine synthetase</i> , <i>β-globulin</i> and <i>albumin</i> .
Iron	Proteins containing iron are classified into two categories. The first is haeme, where this metal is chelated by <i>porphyrin</i> (a water-insoluble ligand). This class is constituted by <i>haemoglobin</i> , <i>myoglobin</i> and <i>cytochromes</i> . The second is formed by non-haeme iron. The principal examples are <i>transferrin</i> , <i>ferritin</i> , <i>ovotransferrin</i> , <i>casein</i> , <i>hemosiderin</i> and <i>albumin</i> .



DECIPHERING METALLOMICS...

- I) How the element (metal or metalloid) is distributed in cellular compartments of a cell

- II) Its coordination environment; in which biomolecule is complexed or what is the bioligand involved in its complexation

- III) Individual concentrations of the metallic species



Classification on the metallomic information

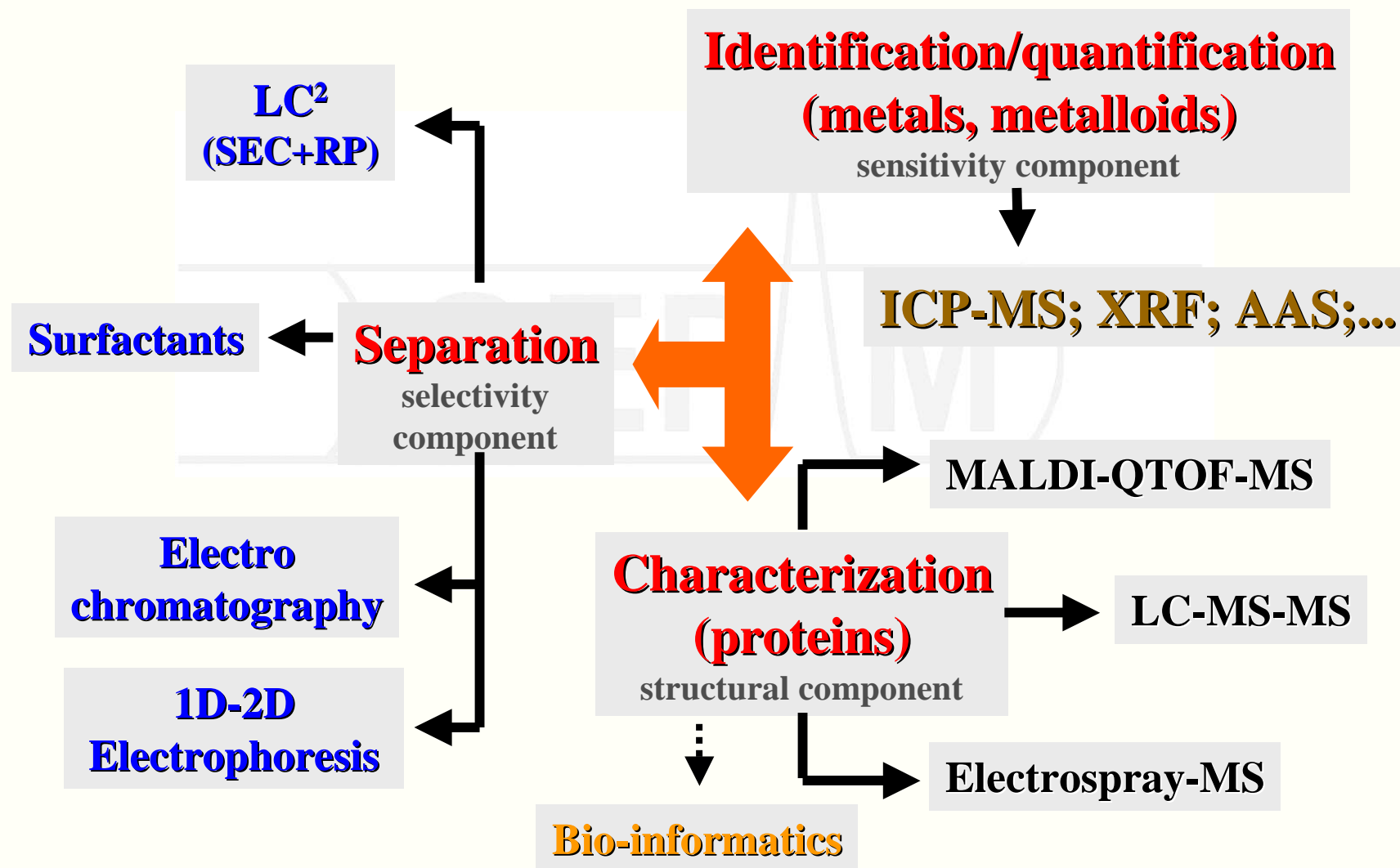
Qualitative metallomics: individual species identification

Quantitative metallomics: concentration determination of the species

Comparative metallomics: monitoring of metallome alterations related to an organism, under influence of external stimulus.



Strategies on metalloproteomics studies



Strategies on metalloproteomics studies

Should the sample preparation be the same one employed to proteomics studies?

To be or not to be...it is the question

Evaluation of metalloprotein extraction procedures in
phytotherapeutic medicines

Aesculus hippocastanum L. – Horse Chestnut

Magalhães & Arruda, *Talanta* 71(2007)1958



Protein extraction procedures

1 shaken (water) → centrifugation → 25°C


2 shaken (buffer) → centrifugation → 25°C



3 shaken → dialysis (42h) → centrifugation → 25°C



4  (water) → centrifugation → 25°C

5  (buffer) → centrifugation → 25°C

6  (water) → dialysis → centrifugation → 25°C

7 shaken (water) →  → centrifugation → 25°C



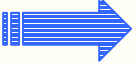




























8  (water) →  → centrifugation → 25°C

9  (buffer) →  → centrifugation → 25°C

10 as in procedure 8 → 40°C

11 as in procedure 9 → 40°C



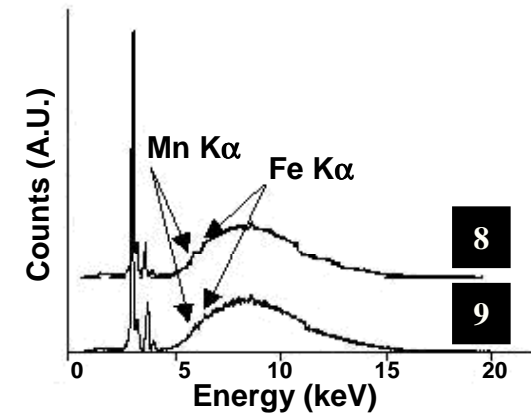
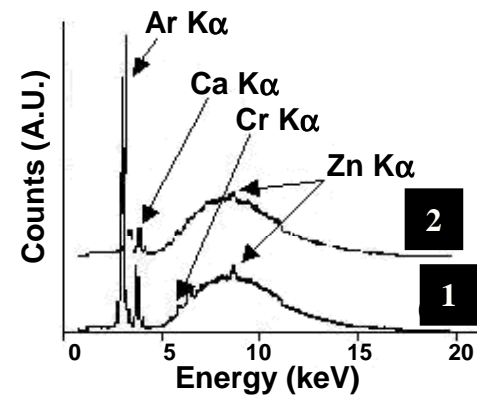
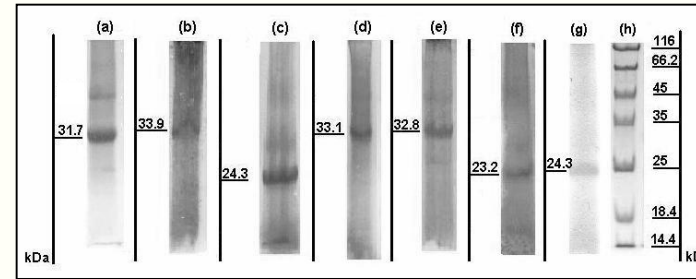
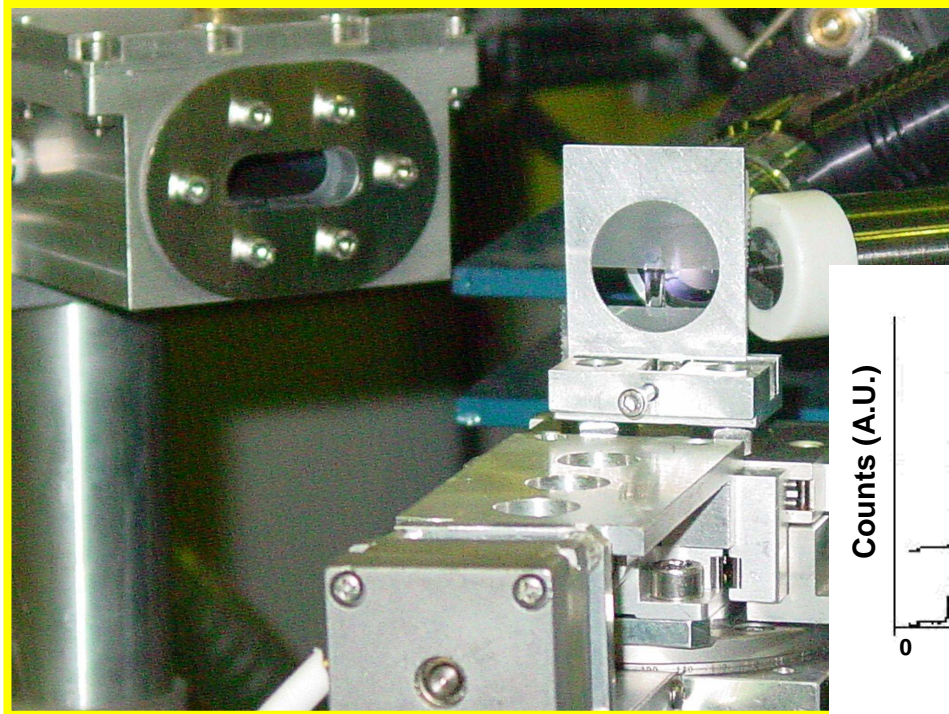
- | | | | | | | | |
|-----------|--|---|---|--|----------------|---|--------------------------|
| 1 | shaken (water) |  | centrifugation |  | 25°C | mg/g
1.81±0.06 | |
| 2 | shaken (buffer) |  | centrifugation |  | 25°C | 1.51±0.06 | |
| 3 | shaken |  | dialysis (42h) |  | centrifugation |  | 25°C
1.22±0.04 |
| 4 |  (water) |  | centrifugation |  | 25°C | 3.03±0.08 | |
| 5 |  (buffer) |  | centrifugation |  | 25°C | 3.6±0.2 | |
| 6 |  (water) |  | dialysis |  | centrifugation |  | 25°C
1.74±0.08 |
| 7 | shaken (water) |  |  |  | centrifugation |  | 25°C
2.51±0.04 |
| 8 |  (water) |  |  |  | centrifugation |  | 25°C
3.38±0.09 |
| 9 |  (buffer) |  |  |  | centrifugation |  | 25°C
5.5±0.1 |
| 10 | as in procedure 8 → 40°C | | | | | 4.8±0.2 | |
| 11 | as in procedure 9 → 40°C | | | | | 3.8±0.2 | |



Mapping the metals... (qualitative metalloproteomics)







Marco A. Z. Arruda














1 shaken (water) \Rightarrow centrifugation \Rightarrow 25°C **1.81±0.06**

2 shaken (buffer) \Rightarrow centrifugation \Rightarrow 25°C **1.51±0.06**

8  (water) \Rightarrow  centrifugation \Rightarrow 25°C **3.38±0.09**

9  (buffer) \Rightarrow  centrifugation \Rightarrow 25°C **5.5±0.1**

Determining the metals... (quantitative metalloproteomics)

- 4  (water)  centrifugation  25°C **3.03±0.08**
- 5  (buffer)  centrifugation  25°C **3.6±0.2**
- 9  (buffer)    centrifugation  25°C **5.5±0.1**

Procedure	Ca ($\mu\text{g g}^{-1}$)	Cr ($\mu\text{g g}^{-1}$)	Fe ($\mu\text{g g}^{-1}$)	Mn ($\mu\text{g g}^{-1}$)
1	12.5±0.5	4.98 ±0.05	35 ±6	2.7 ±0.4
2	9.8 ±0.7	1.6 ±0.7	38 ±2	1.9 ±0.2
4	6.4 ±0.3	4.00 ±0.02	40 ±7	15 ±1
5	11.4 ±0.8	1.5 ±0.4	37 ±7	12 ±2
8	0.74 ±1	< LOD	23 ±9	< LOD
9	< LOD	1.3 ±0.1	38 ±3	1.1 ±0.4

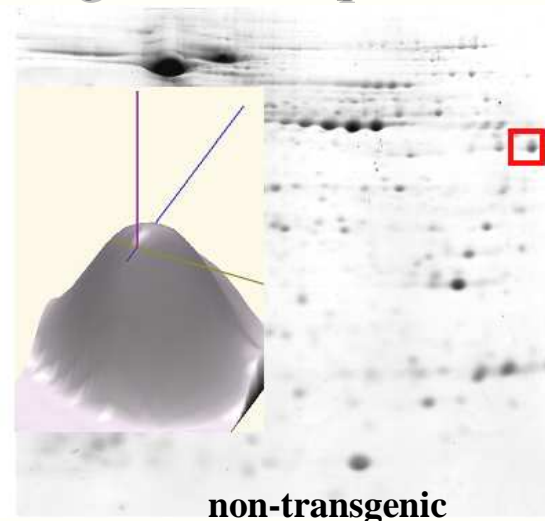
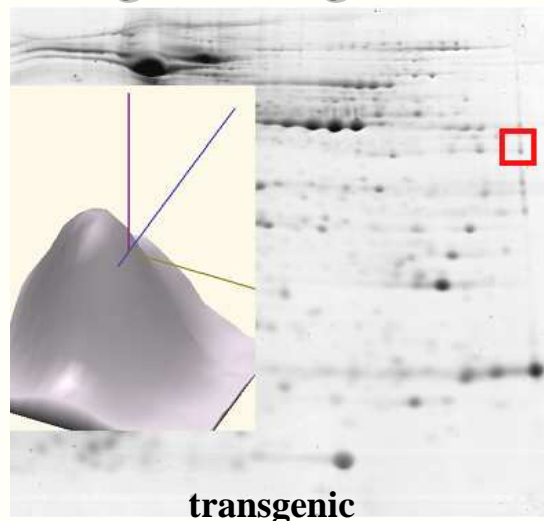
LOD: 0.05 $\mu\text{g g}^{-1}$ Ca; 0.054 $\mu\text{g g}^{-1}$ Cr; 0.99 $\mu\text{g g}^{-1}$ Fe; 0.06 $\mu\text{g g}^{-1}$ Mn



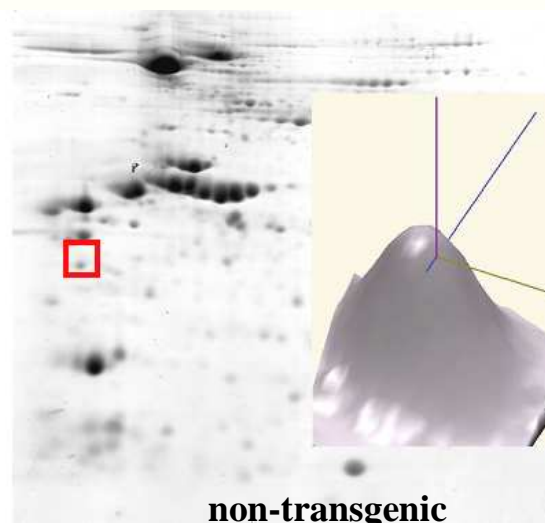
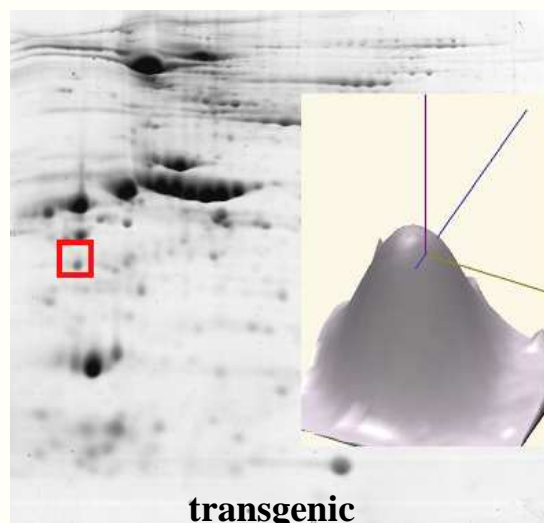
Comparative metallomics

Sussulini *et al.*, *J. Anal. At. Spectrom.*, 2007, DOI: 10.1039/b706684h

Transgenic organisms: changes in the proteome!!



54.50 kDa
pI: 6.83



27.03 kDa
pI: 4.37

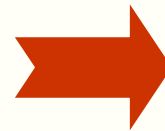
Hypothesis: also metallome changes??



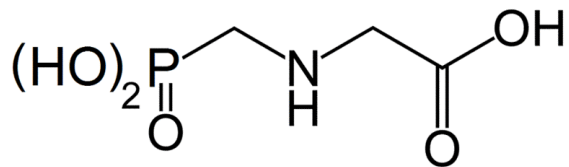
Comparative metallomics

Transgenic soybean [*Glycine max* (L.) Merrill]: *Roundup Ready*[®]

Action of glyphosate: inhibits the EPSP synthase enzyme, which participates on aromatic amino acids synthesis in plants



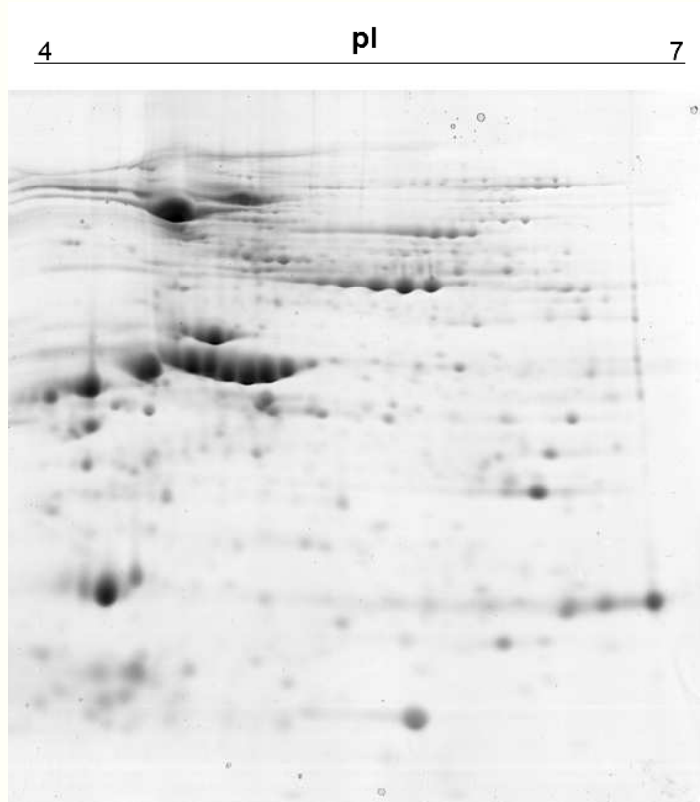
Soybean genetic modification: based on inserting the CP4 EPSPS gene from *Agrobacterium* sp. strain CP4 that provides CP4 EPSPS (EC 2.5.1.19) protein (tolerant to glyphosate)



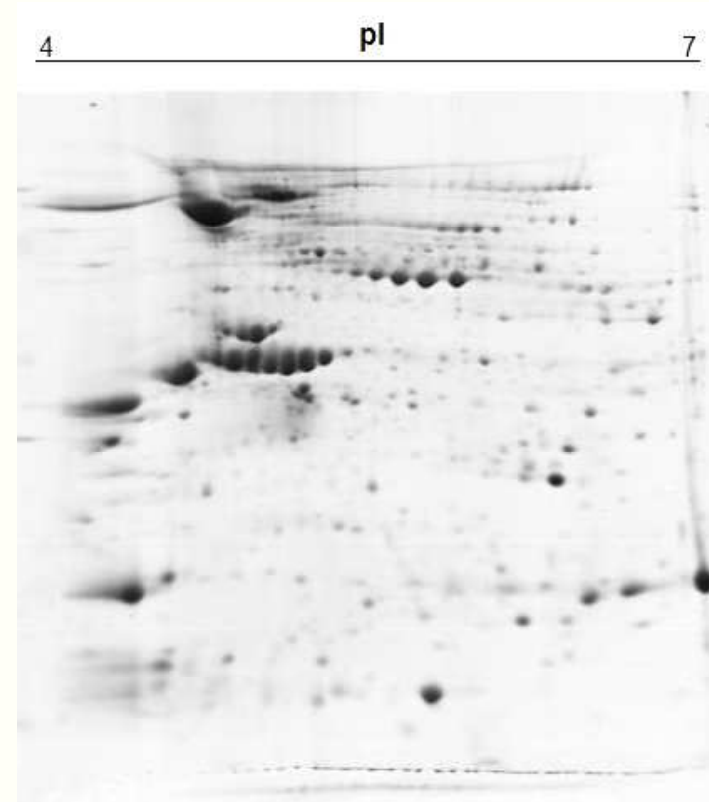
Non-selective herbicide:
non-transgenic plants are
also exterminated



Comparative metallomics



transgenic

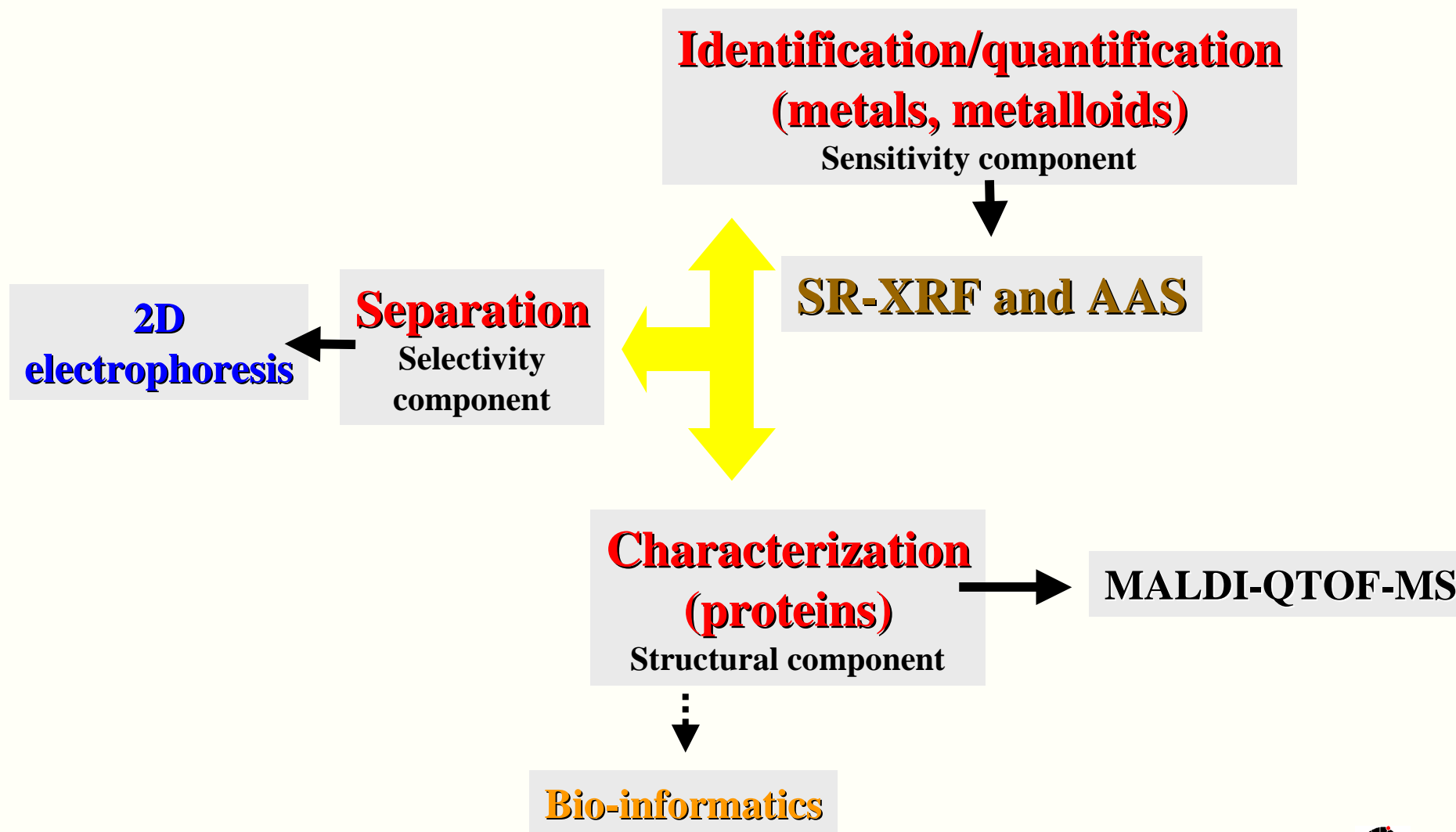


non-transgenic

Transgenic soybean: 408 ± 27
Non-transgenic soybean: 397 ± 26
match: ca. 70% (n=3) and ca. 40%

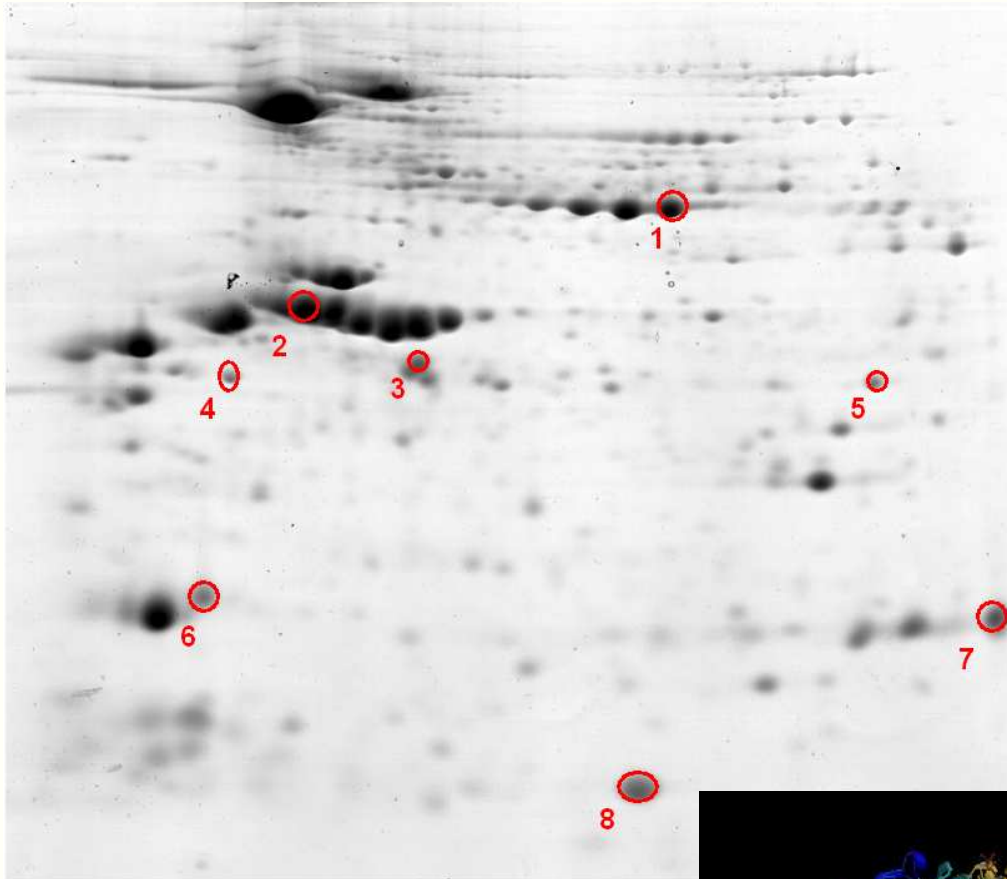


Comparative metallomics - Strategy



Comparative metallomics

Selectivity and Structural components



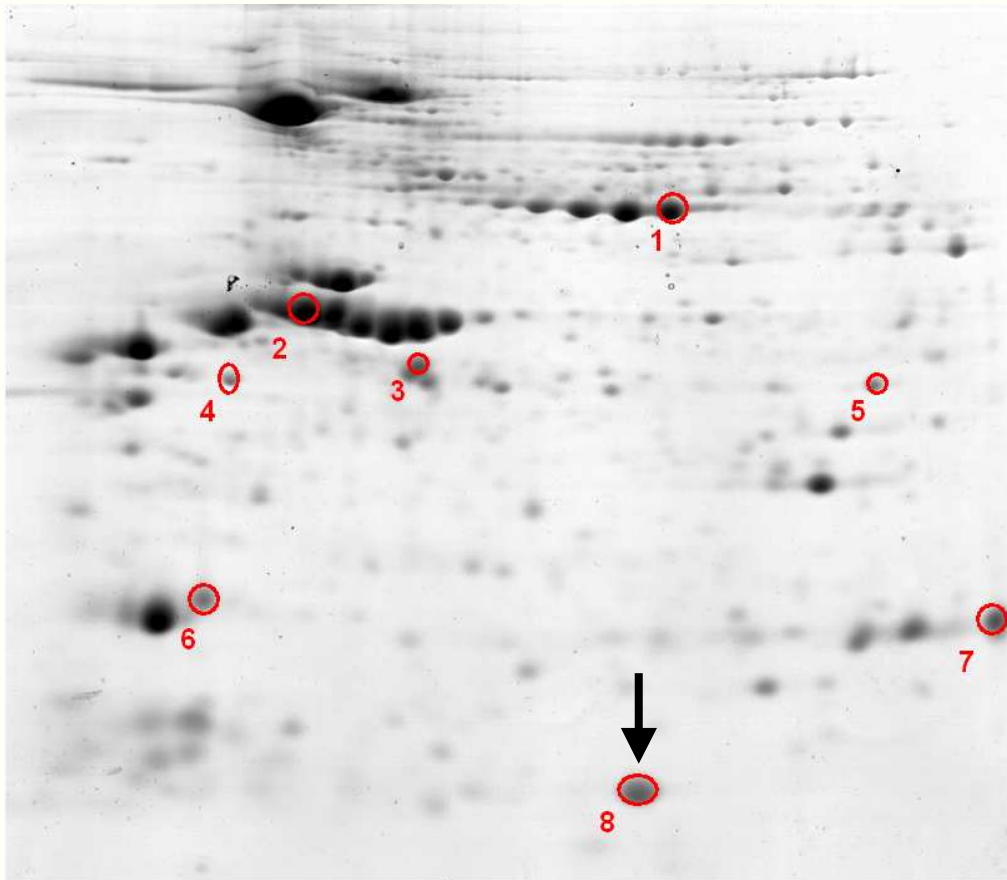
- 1: β -conglycinin β -chain precursor
 - 2: Glycinin G2
 - 3: Soybean agglutinin
 - 4: Seed maturation protein PM 25
 - 5: not identified
 - 6: not identified
 - 7: Glycinin chain A2B1a precursor
 - 8: Glycinin G4 (precursor)
- MALDI-QTOF MS

transgenic



Comparative metallomics

Sensitivity component



LA-ICP-MS



Extracted from Agilent CD

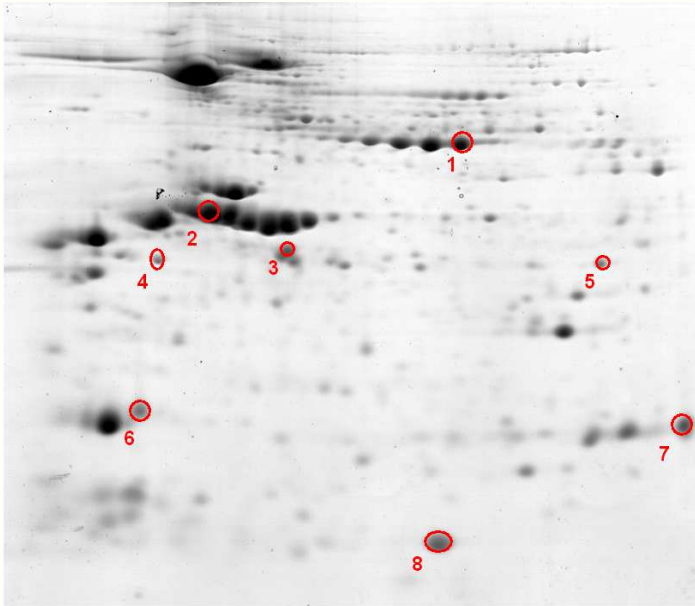
Binet et al., *Anal. Biochem.* 318(2003)30
Escherichia coli (Cd/Zn) \rightarrow \gg [metalloproteins]
under stress conditions



Marco A. Z. Arruda

Comparative metallomics

Sensitivity component



gel decomposition



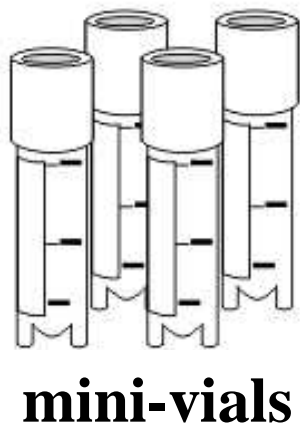
mini-vials



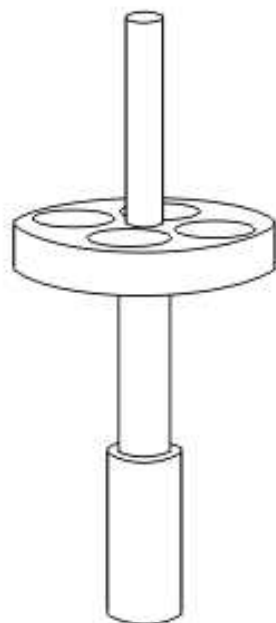
Comparative metallomics

Sensitivity component

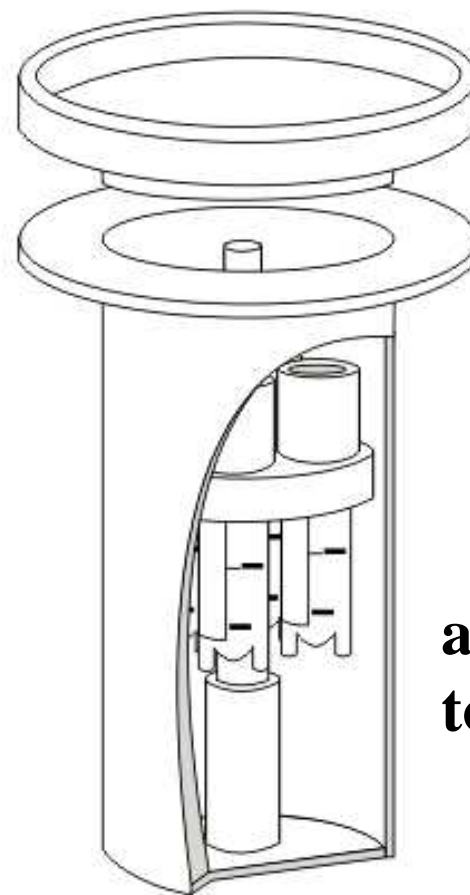
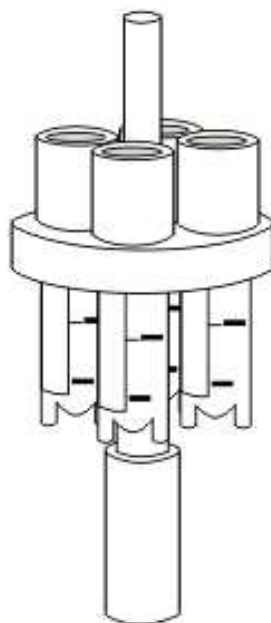
Sussulini et al., *Anal. Biochem.* 361(2007)146



support



system

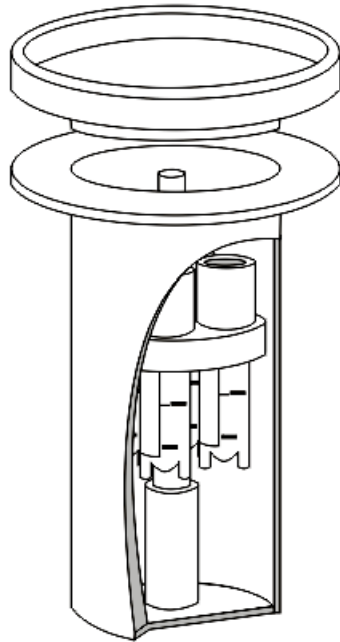


**adaptation
to the MW
vessel**



Comparative metallomics

Sensitivity component



1 ^a step	300W - 1 min
2 ^a step	0W - 1 min
3 ^a step	500W - 30 s
4 ^a step	0W - 2 min
5 ^a step	500W - 1 min
6 ^a step	0W - 2 min
7 ^a step	800W - 30 s
8 ^a step	0W - 2 min
9 ^a step	800W - 30 s
10 ^a step	0W - 2 min
11 ^a step	800W - 30 s
12 ^a step	0W - 2 min
13 ^a step	500W - 1 min

Time: 16 min (2x)

Reagents: 200 μL HNO_3 + 150 μL H_2O_2

Mass: 1 – 2.5 mg

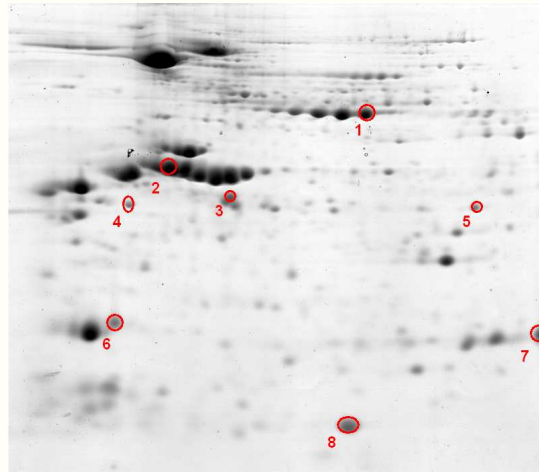
residual C : < 1%



Comparative metallomics

Sensitivity component – SR-XRF

Spot	TS	N-TS
1	Fe(+)	Fe
2	ND	Ca, Fe
3	Ca(+), Cu(+)	Ca, Cu
4	Ca	ND
5	Ca(+), Cu(+), Fe(+), Zn(+)	Ca, Cu, Fe, Zn
6	Ca, Fe	Ca(+), Fe(+), Mn, Ni
7	Ca	Ca(+), Fe
8	ND	Fe



Comparative metallomics

Sensitivity component – AAS

Spot	Ca ^a (mg g ⁻¹)		Cu ^b (µg g ⁻¹)		Fe ^b (µg g ⁻¹)	
	TS	N-TS	TS	N-TS	TS	N-TS
1	ND ^c	ND	ND	ND	< LOQ ^d	< LOQ
2	< LOQ	< LOQ	ND	ND	212 ± 31	348 ± 49
3	2.6 ± 0.2	2.2 ± 0.4	1.5 ± 0.2	2.8 ± 0.2	ND	ND
4	< LOQ	< LOQ	ND	ND	ND	ND
5	17 ± 2	3.6 ± 0.5	1.6 ± 0.1	< LQ	691 ± 78	447 ± 54
6	3.5 ± 0.2	15 ± 2	ND	ND	663 ± 79	869 ± 93
7	< LOQ	< LOQ	ND	ND	< LOQ	< LOQ
8	ND	ND	ND	ND	< LOQ	< LOQ

a: FAAS; b: ETAAS; c: not determined; d: limit of quantification

Proteins of spots 1,2,7,8 → do not present metal in their structures;

Spot 2: Fe(II) micronutrient → better nutrient preservation??

Protein of spot 3 → involved in the metallic ions coordination from a bond site with carbohydrate [Ca(II), Mn(II) and others transition metallic ions]



Conclusions

Different results in terms of
protein and/or metal extraction

Decisive to preserve metals
in the protein structure

Extraction procedure
Extraction medium

Must be carefully chosen:
proteomics and/or metalloproteomics



References

N. Jakubowski, R. Lobinski, L. Moens,

J. Anal. At. Spectrom., 19 (2004) 1

H. Haraguchi,

J. Anal. At. Spectrom., 19 (2004) 5

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Anal. Bioanal. Chem., 378 (2004) 54

J. S. Garcia, C. S. Magalhães, M. A. Z. Arruda,

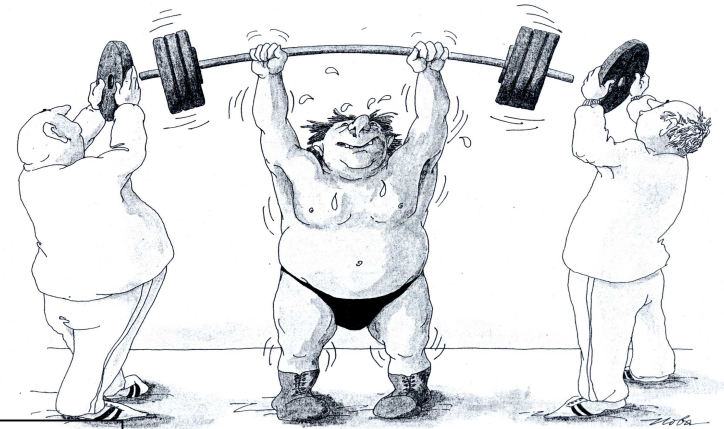
Talanta, 69 (2006) 1



General Conclusions



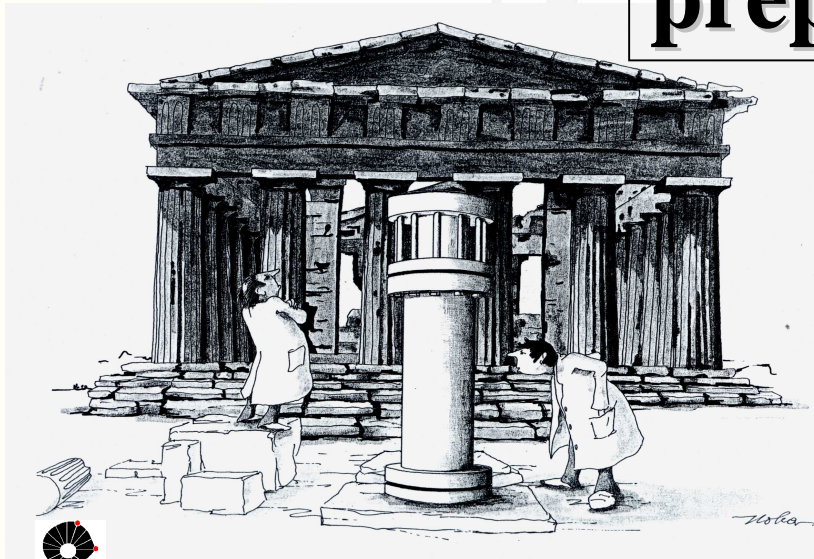
in-deep view



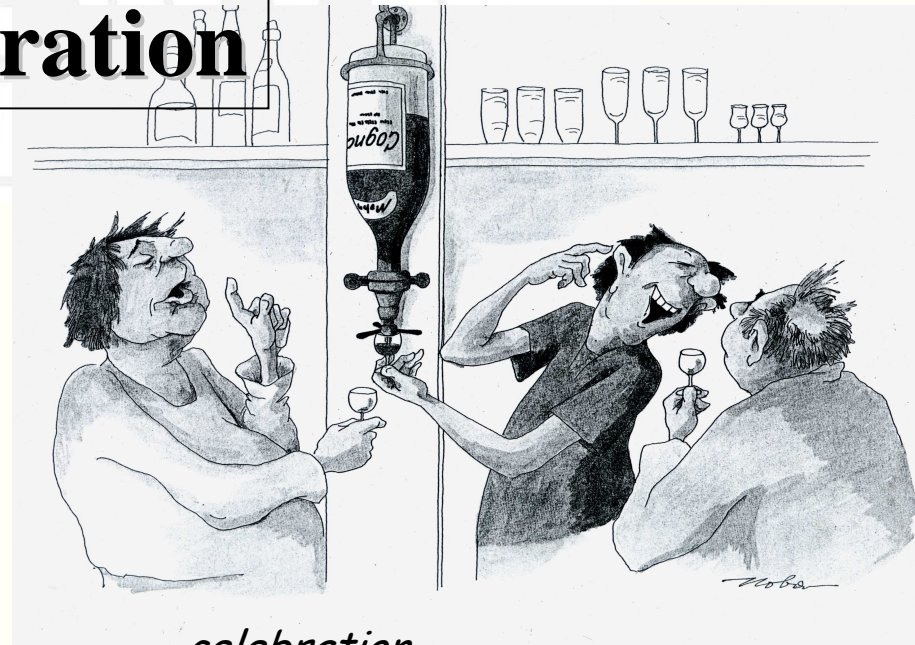
ability

knowledge

Sample preparation



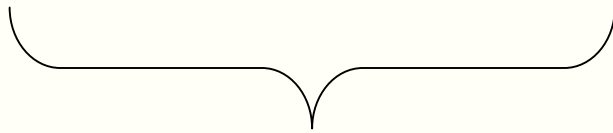
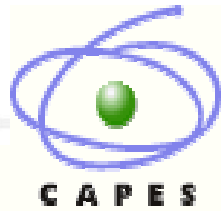
first stages



celebration...



Special thanks...



UNICAMP

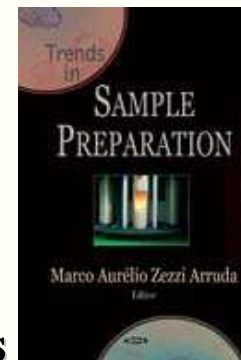


...and even more



Marco A. Z. Arruda

Trends in Sample Preparation



Günter Knapp: Preface

Pedro V. Oliveira: Micro sampling for solid and slurries analytical methods

Joaquim Nóbrega: Microwave-assisted procedure for sample preparation: new developments

Érico Flores: Trends in sample preparation using combustion techniques

Patricia Smichowski: Sample preparation of atmospheric aerosols for elemental analysis and fractionation studies

Fabio Augusto: Extraction and pre-concentration techniques for chromatographic analysis

José M. Pingarrón: Strategies in sample preparation for applications in analytical electrochemistry

Elias A. G. Zagatto: In-line sample preparation in flow analysis

Miguel Valcárcel: The role of vanguard-rearguard strategies in sample preparation in routine analytical laboratories

Marco A. Z. Arruda: Strategies for sample preparation focusing biomolecules determination/characterization

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) GEP M)

