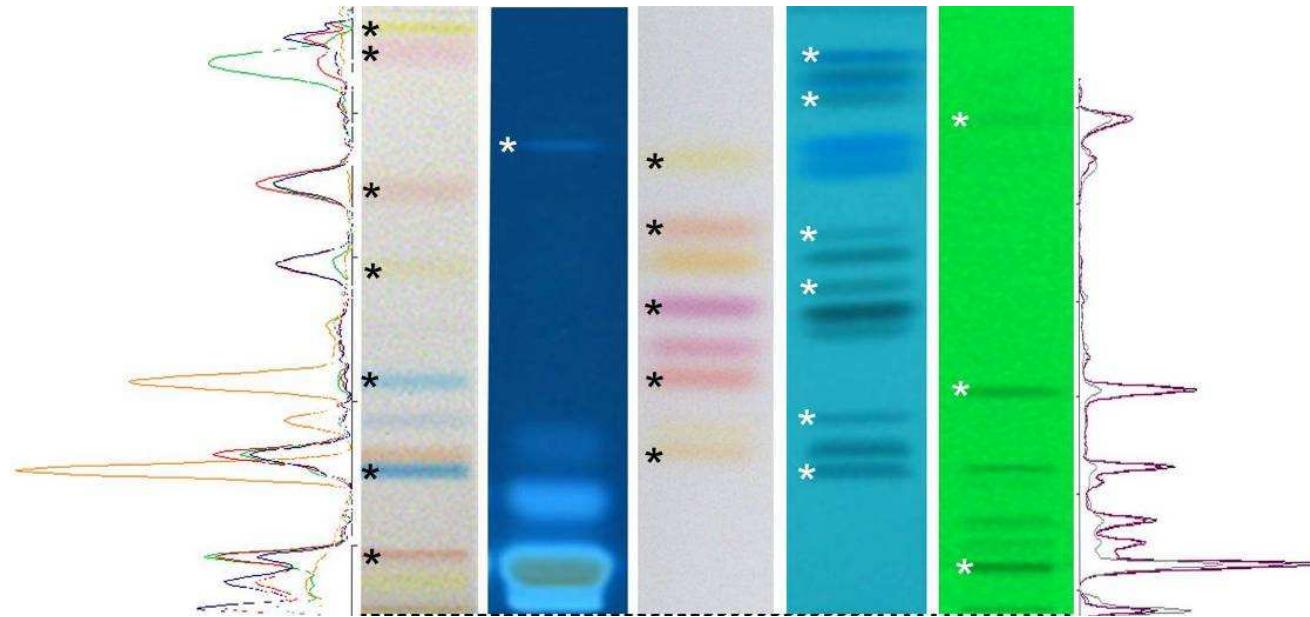


# Hyphenations in HPTLC with UV/Vis/FLD, MS, FTIR, NMR and bioassays



apl. Prof. Dr. habil. Gertrud Morlock  
Institute of Nutritional Science, Justus-Liebig-University of Gießen  
Institute of Food Chemistry, University of Hohenheim, Stuttgart

# Hyphenation

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- 1980: term hyphenation by Hirschfeld
- comprises the different approaches to combine mainly spectrometers with chromatographic systems to get further information about the sample
- **hyphen** (-) symbolizes this attempt of combination, which did not reach its stage of full maturity so far
- **slash** (/) is found for hyphenated methods at a mature state

Nobody takes care about it!

- 2007: term “hypernation” (super-hyphenation) by Wilson and Brinkman  
→ to place all of the required spectrometers into a single system  
so that all of the spectroscopic information is obtained in a single run

# Hyphenation

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Problems associated with column-based hyphenations

- Capital cost and strategies for dealing with the large amounts of data produced by such systems.
- Complexity of instrumentation increases → difficult to operate in routine
- A single eluent (→ optimal for all detectors) is difficult to obtain.
- Differences in sensitivity are challenging.

Less challenging in HPTLC-based hyphenations

- Open system is highly adaptive to different sensitivities
- Cost-effective by modular instrumentation
- Generating less data due to targeted access to points-of-care
- Directly accessible for the respective optimal solvent

# Hyphenation

Poster 3a

→ The main difference

HPLC: sample in solvent; after separation → sample in waste

HPTLC: solvent evaporated; after separation → sample on plate

Journal of Chromatography A, 1217 (2010) 6600–6609



Contents lists available at ScienceDirect

Journal of Chromatography A

journal homepage: [www.elsevier.com/locate/chroma](http://www.elsevier.com/locate/chroma)



Review

## Hyphenations in planar chromatography

Gertrud Morlock\*, Wolfgang Schwack

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- HPTLC-UV/Vis/FLD-MS [13,14],
- HPTLC-UV/Vis/FLD-bioactivity-HRMS [15],
- HPTLC-UV-FTIR [16,17],
- HPTLC-UV/Vis/FLD-FTIR ATR [18],
- TLC-Vis-SERS [12].

### ARTICLE INFO

#### Article history:

Available online 20 May 2010

#### Keywords:

Mass spectrometry  
High-performance thin-layer chromatography  
Effect-directed analysis  
Bioassays  
Cost-effective analysis  
High-throughput system

### ABSTRACT

This review is focused on planar chromatography and especially on its most important subcategory high-performance thin-layer chromatography (HPTLC). The image-giving format of the open, planar stationary phase and the post-chromatographic evaporation of the mobile phase ease the performance of various kinds of hyphenations and even super-hyphenations. Examples in the field of natural product search, food and lipid analysis are demonstrated, which point out the hyphenation with effect-directed analysis (EDA) and mass spectrometry and illustrate the efficiency gain. Depending on the task at hand, hyphenations can readily be selected as required to reach the relevant information about the sample, and at the same time, information is obtained for many samples in parallel. The flexibility and the unrivalled features through the planar format valuably assist separation scientists.

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# Content

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## Hyphenations with

1. UV/VIS/FLD/derivatizations
2. MS
3. FTIR
4. NMR
5. Bioassays

# Hyphenation with MS

Trends in Analytical Chemistry, Vol. 29, No. 10, 2010, 1157-1171

Trends

## Coupling of planar chromatography to mass spectrometry

Gertrud Morlock, Wolfgang Schwack

Coupling of planar chromatography to mass spectrometry (MS) and especially ambient MS is a relatively new field of great interest. The direct sample access at ambient conditions and the feasibility to obtain mass spectra free of contamination within a minute or even within seconds greatly contributes to the progress of planar chromatography. Targeted recording of mass spectra on zones of interest is performed after evaluation of the chromatogram, thus providing high efficiency. Reported approaches for coupling are divided into elution-based and desorption-based techniques. Devices of both categories are commercially available. As a consequence of increasing importance, a rethink of the terminology of liquid chromatography with MS has to be considered.  
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**Keywords:** Ambient mass spectrometry; Cost-effective analysis; Coupling to mass spectrometry; Desorption-based technique; Elution-based technique; High-performance thin-layer chromatography; HPTLC-MS; Planar chromatography; Thin-layer chromatography

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Institute of Food Chemistry,  
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Stuttgart,  
Germany

### 1. Introduction

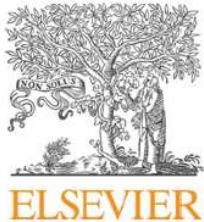
Planar chromatography comprises all chromatographic techniques that have an open planar stationary phase present as or on a plane [1]. Therein, simple thin-layer chromatography (TLC) is the most widespread chromatographic technique; whereas high-performance TLC (HPTLC) is considered as the most efficient and powerful planar chromatographic technique, with optimized coating material (lower particle size and narrower particle-size distribution) combined with advanced instrumentation for most of the steps of the chromatographic process [2]. Paper chromatography is not used very much at

the MS by means of a specially-built inlet probe [6].

Since that time, the spectrometry market has continued to grow and column chromatography has been brought forward, but coupling of an open planar system with MS required more effort than column-derived techniques. Although reviews about TLC-MS were regularly reported by Busch [7–10] or Wilson [11–13], it was not until the past decade that it attracted interest because of several successful approaches and the invention of ion sources working under ambient conditions and atmospheric pressure, which enormously eased the introduction of a planar object.

# Hyphenation with MS

Journal of Chromatography A, 1218 (2011) 2700–2711



Contents lists available at ScienceDirect

## Journal of Chromatography A

journal homepage: [www.elsevier.com/locate/chroma](http://www.elsevier.com/locate/chroma)



Review

## Thin layer chromatography/mass spectrometry

Sy-Chyi Cheng<sup>a</sup>, Min-Zong Huang<sup>b</sup>, Jentai Shiea<sup>b,c,d,\*</sup>

<sup>a</sup> Institute of Forensic Medicine, Ministry of Justice, Taipei, Taiwan

<sup>b</sup> Department of Chemistry, National Sun Yat-Sen University, Kaohsiung 80424, Taiwan

<sup>c</sup> National Sun Yat-Sen University–Kaohsiung Medical University Joint Research Center, Kaohsiung 80424, Taiwan

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### ARTICLE INFO

*Article history:*

Available online 2 February 2011

*Keywords:*

TLC-MS

Ambient ionization

Vacuum-based ionization

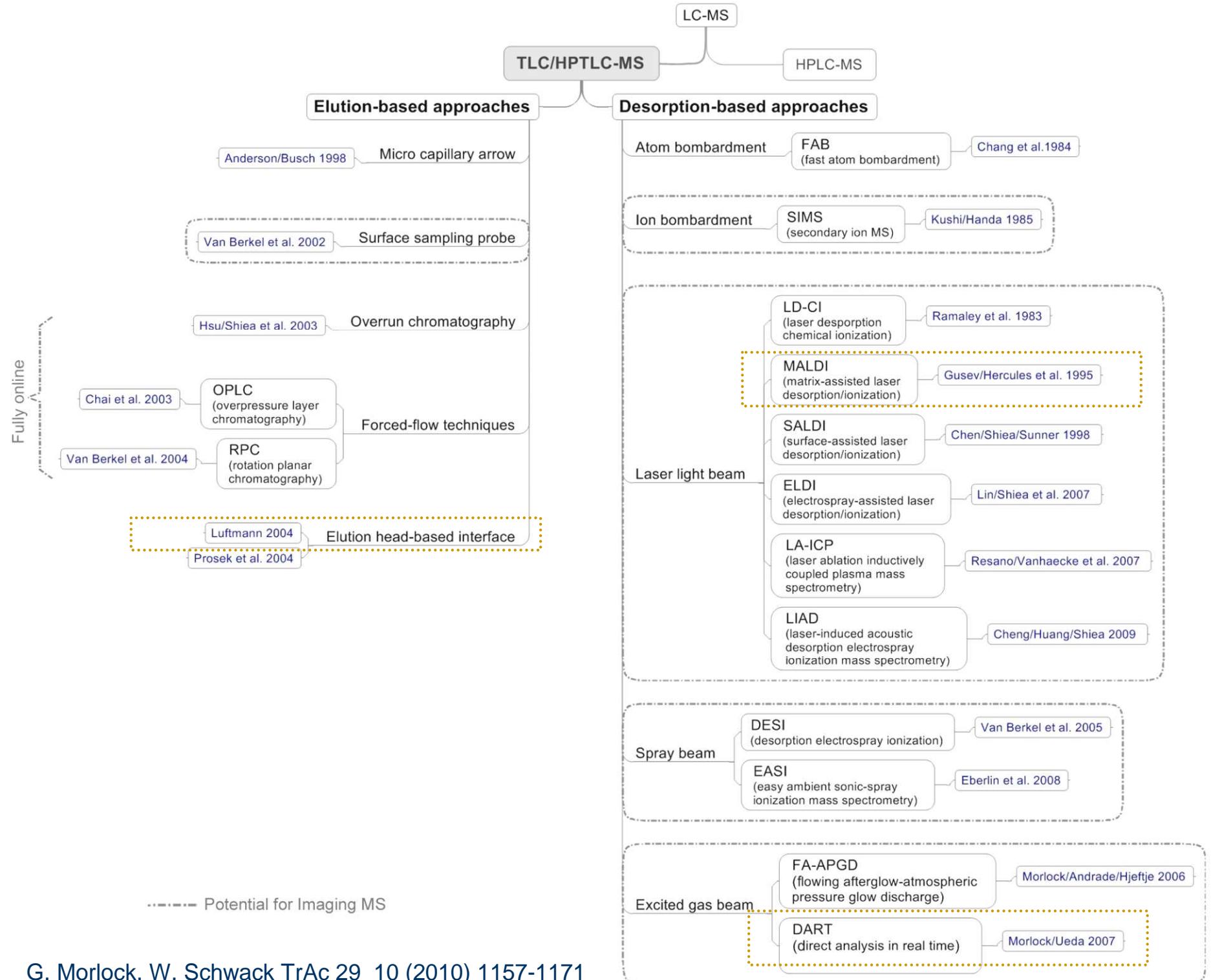
Desorption/ionization

Direct coupling

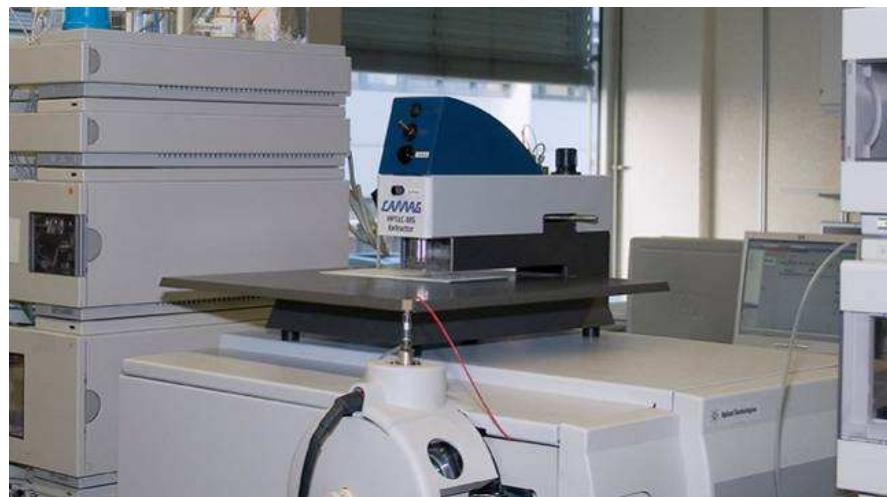
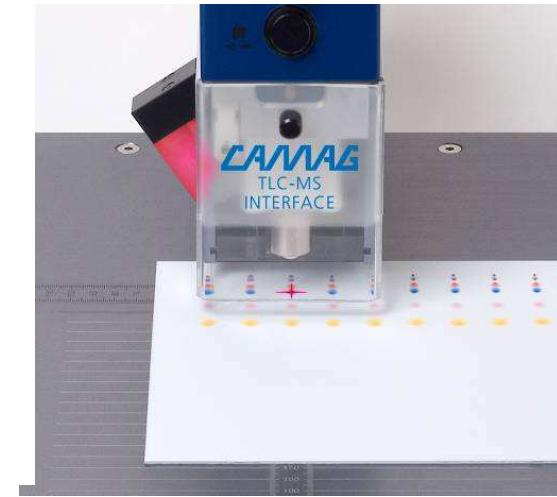
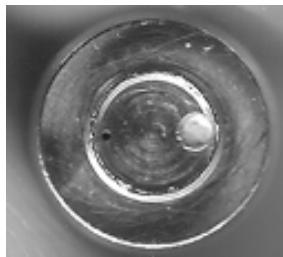
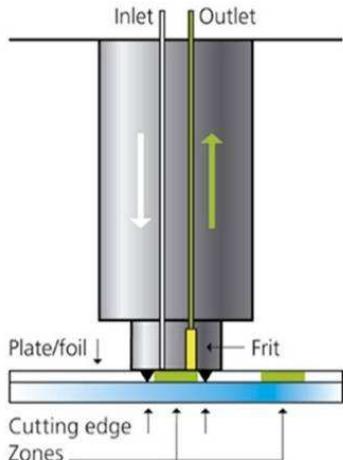
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### ABSTRACT

Thin layer chromatography (TLC)—a simple, cost-effective, and easy-to-operate planar chromatographic technique—has been used in general chemistry laboratories for several decades to routinely separate chemical and biochemical compounds. Traditionally, chemical and optical methods are employed to visualize the analyte spots on the TLC plate. Because direct identification and structural characterization of the analytes on the TLC plate through these methods are not possible, there has been long-held interest in the development of interfaces that allow TLC to be combined with mass spectrometry (MS)—one of the most efficient analytical tools for structural elucidation. So far, many different TLC-MS techniques have been reported in the literature: some are commercially available. According to differences in their



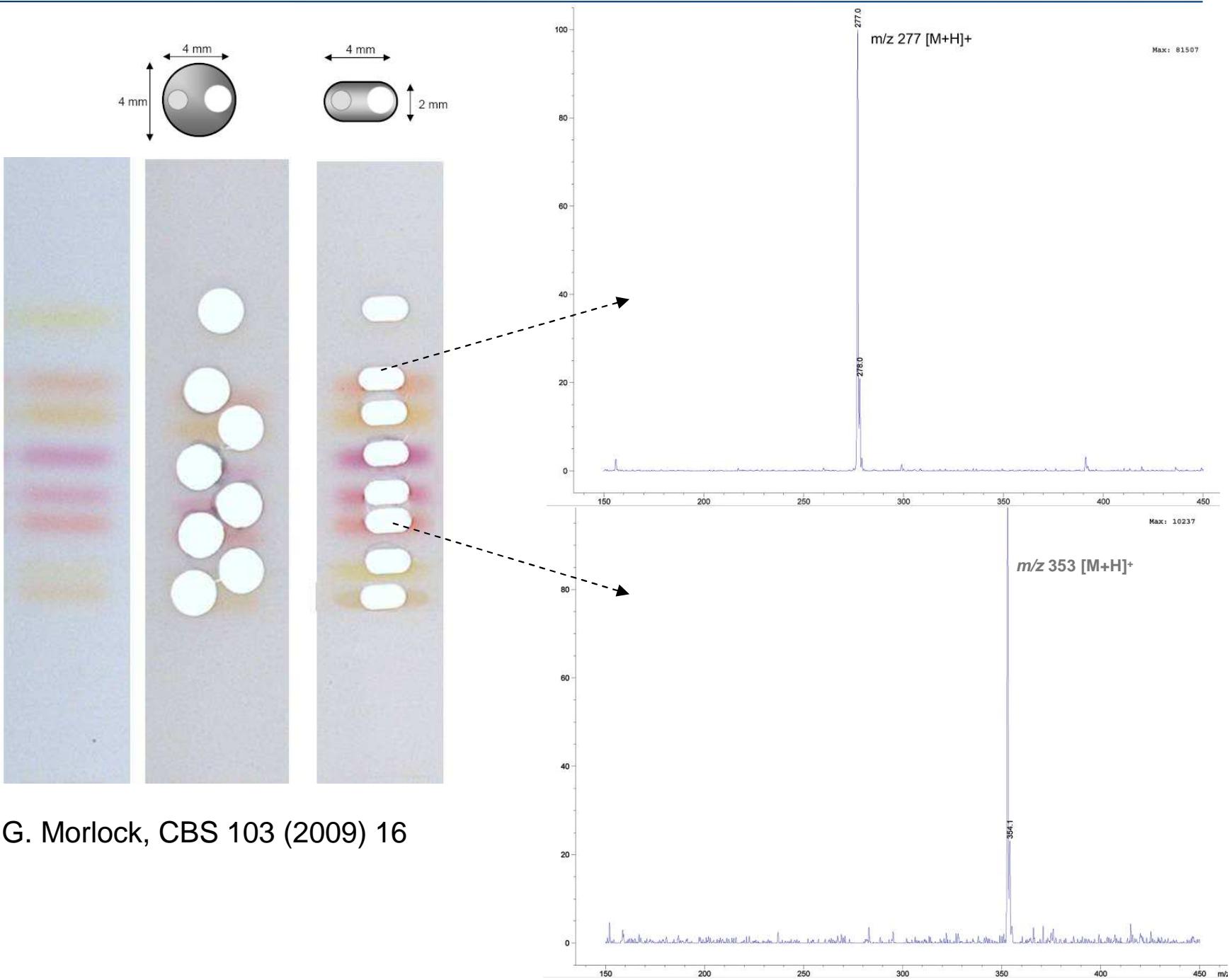
# Elution head-based HPTLC-MS → TLC-MS Interface



H. Luftmann, Anal Bioanal Chem 378 (2004) 964-968

A. Alpmann, G. Morlock, Anal Bioanal Chem 386 (2006) 1543-1551

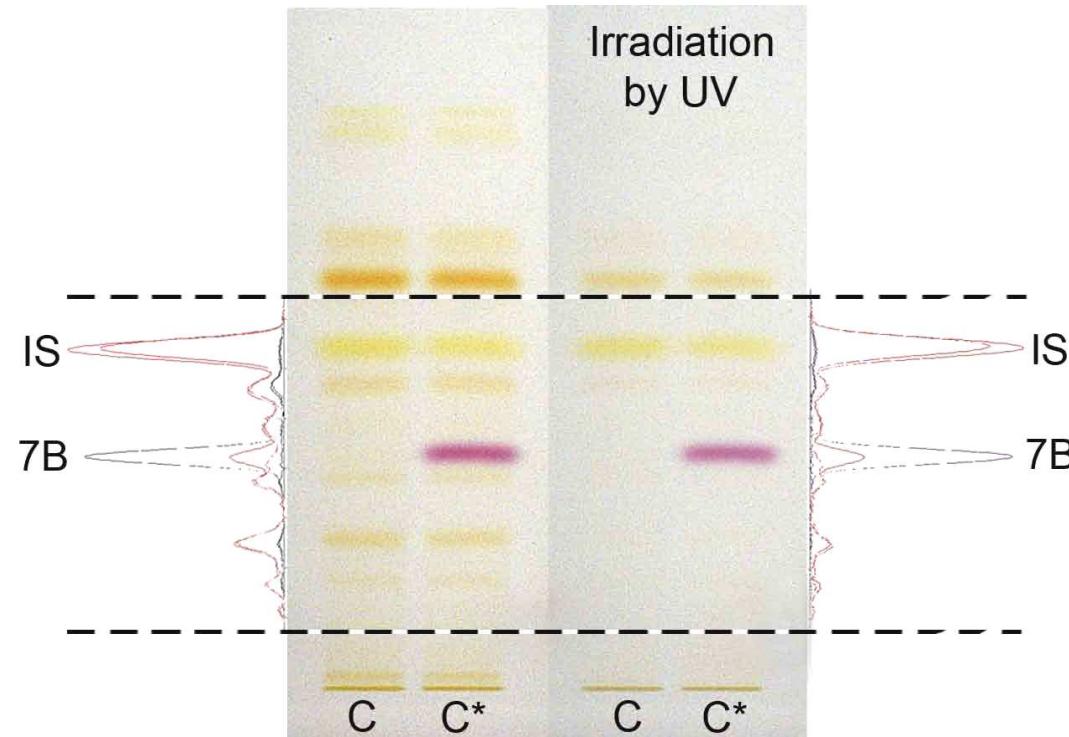
# Elution head-based HPTLC-MS



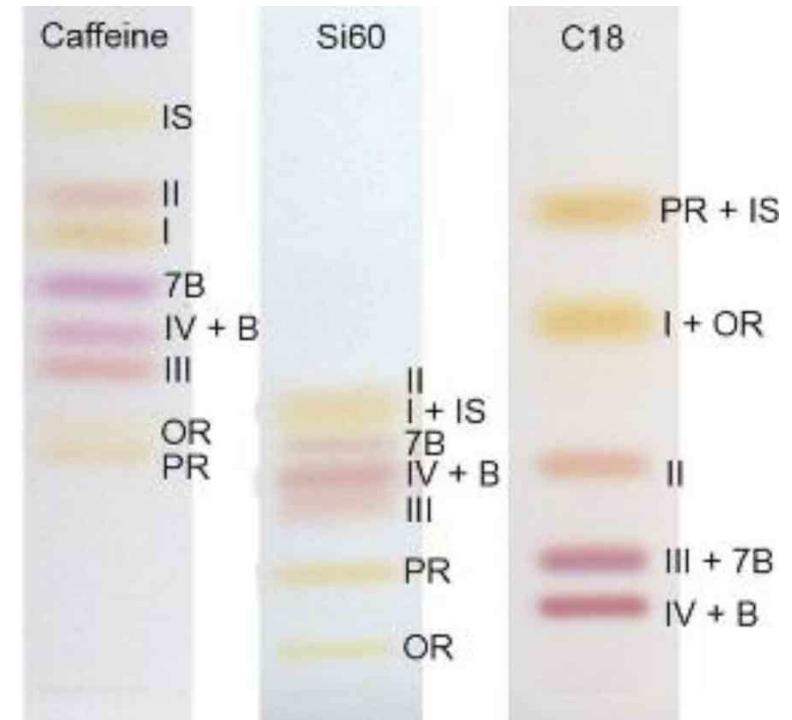
G. Morlock, CBS 103 (2009) 16

# Falsification of food color with unauthorized Sudan dyes

Reactions on the plate



The utmost selectivity change

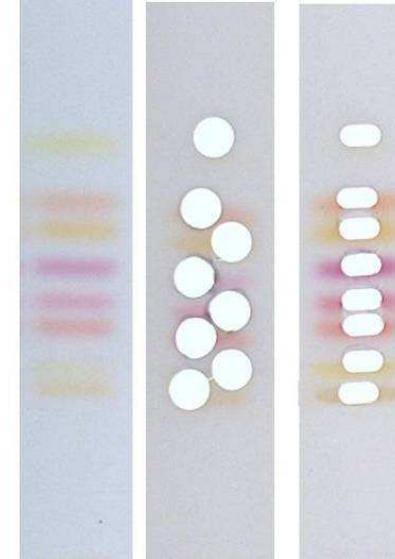
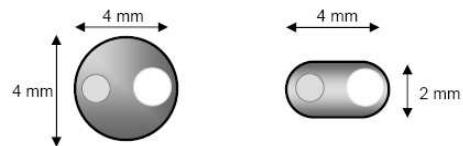


E. Pellissier, W. Schwack, CBS 103 (2009) 13-15

# Elution head

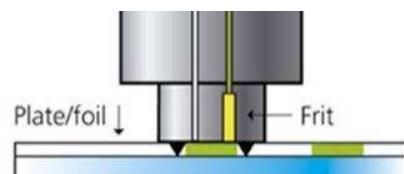
## Cutting edge geometry

- U. Jautz, G. Morlock, J Planar Chromatogr 21 (2008) 367
- G. Morlock , CBS 103 (2009) 16



## Cutting edge height

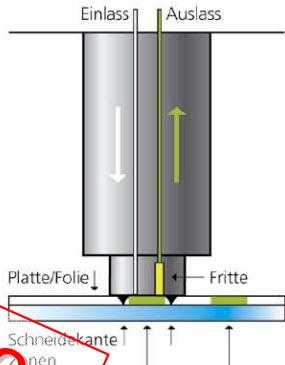
- 0.2 mm for standard layers → CAMAG Bibliography Service CBS 102 (2009)
- 0.1 mm for extra thin layers → U. Jautz, G. Morlock, Anal Bioanal Chem 387 (2007) 1083
- 0.5 mm for preparative layers → E. Dytkiewitz, G. Morlock, J AOAC Int 91 (2008) 1237



# Performance data → TLC-MS Interface

- highly reliable hyphenation
- highly targeted

Poster 3C



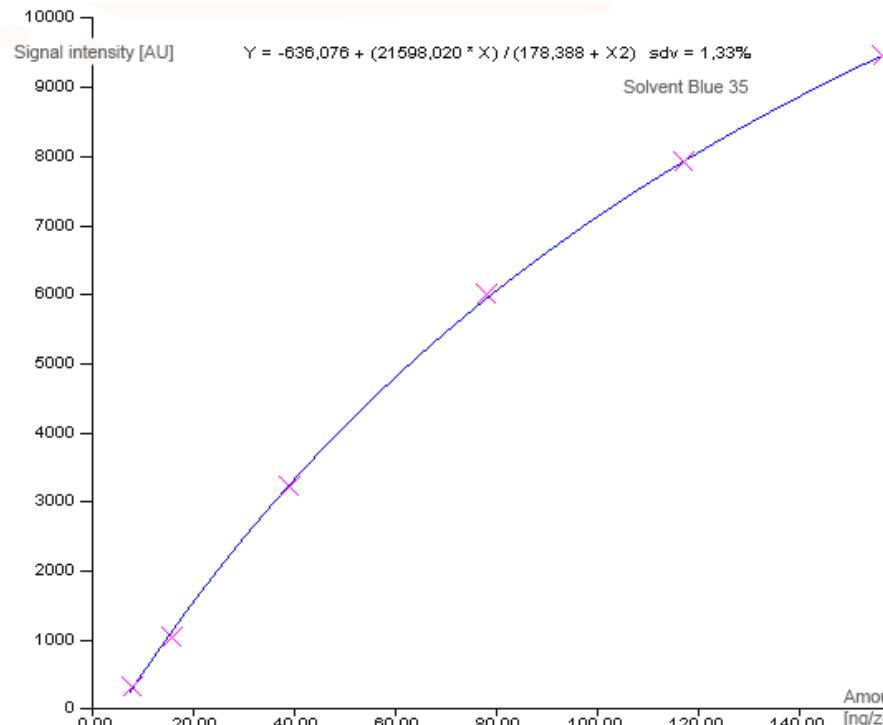
HPTLC-ESI-MS (SIM, peak area)	$hR_F$ - value	Linearity		Precision	
		Calibration range (ng/band)	Determination coefficient	Conc. (ng/band)	%RSD, $n = 5$
Dimethyl Yellow	65	12 – 234	0.9943	1125	8.1
Oracet Red G	50	2 – 39	0.9950	189	11.0
Solvent Blue 35	41	10 – 52	0.9931	750	4.6
Sudan Red G	27	6 – 117	0.9984	564	8.8
Solvent Blue 22	17	21 – 78	0.9976	750	3.8
Oracet Violet 2R	4	8 – 156	0.9752	1500	11.6
Mean			0.9923		8.0



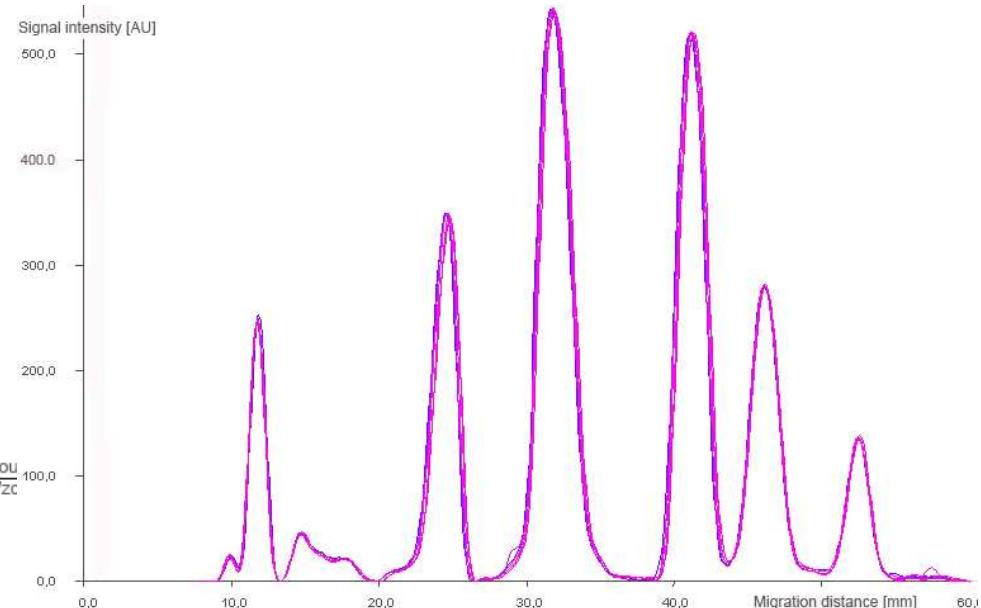
# Performance data obtained with the TLC-MS interface

⇒ before: check of performance data by HPTLC-Vis

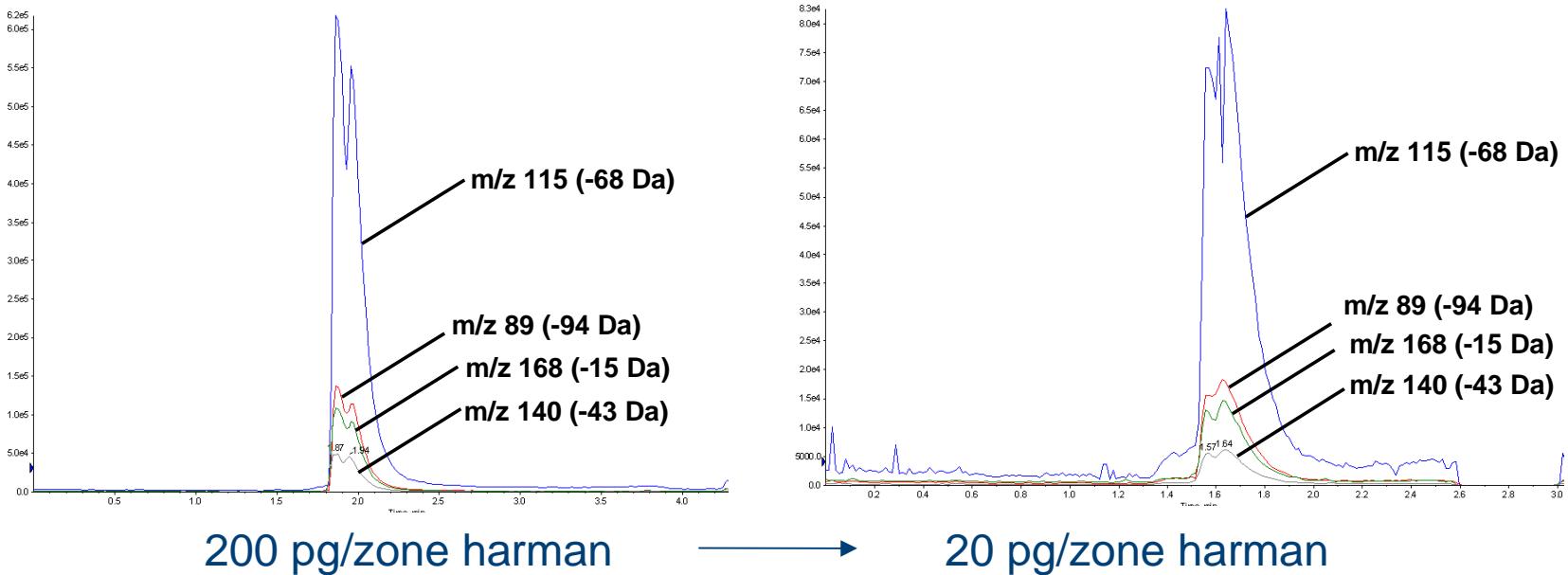
Calibration for Solvent Blue 35 (%RSD = 1.3%)



Overlay of 5 analog curves (%RSD ≤ 1.3%)

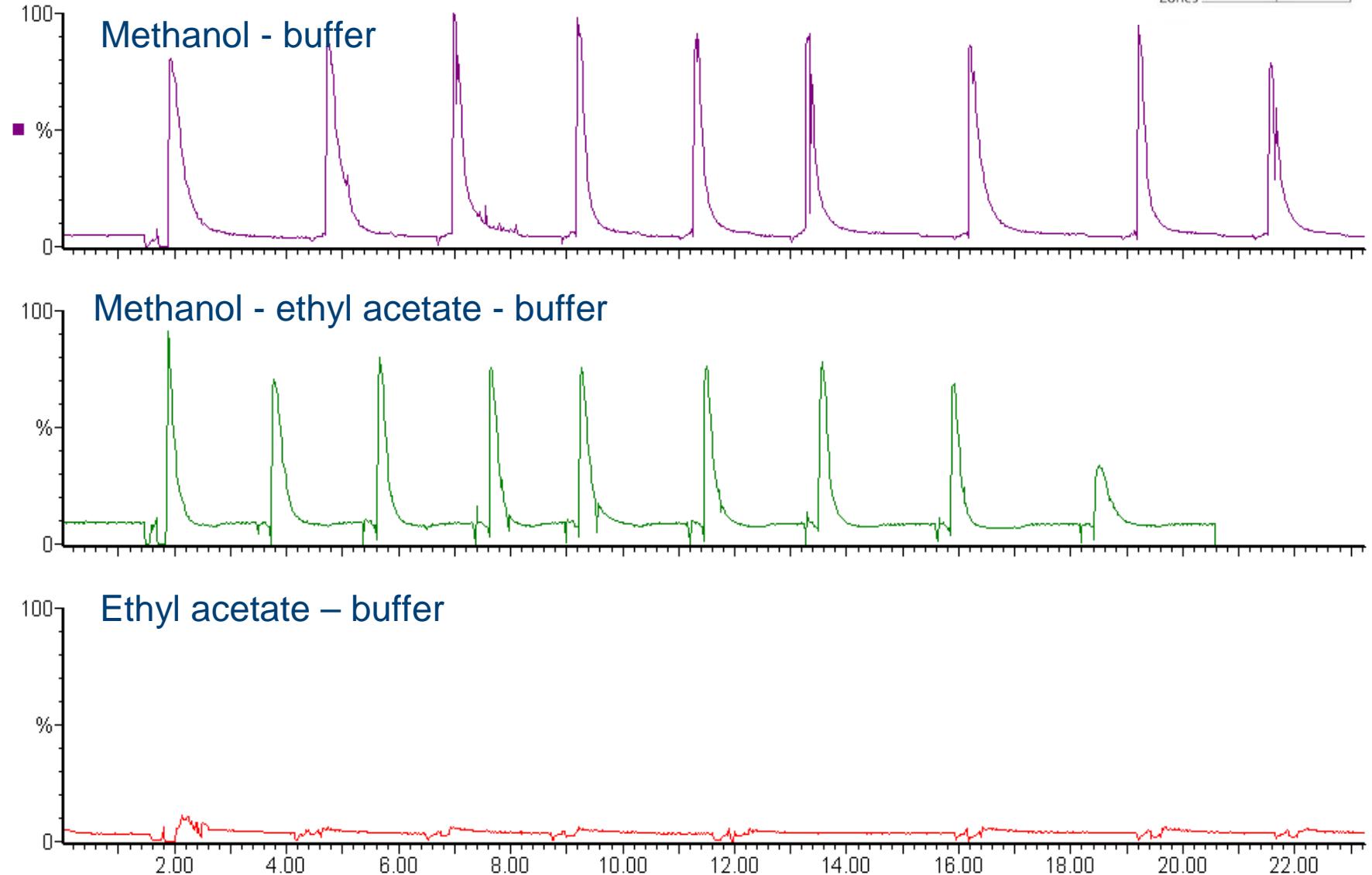
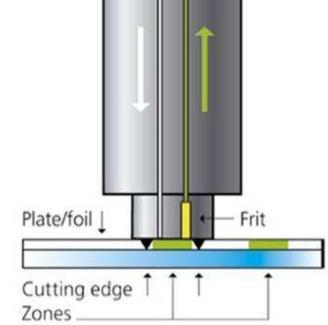


# Detectability by HPTLC-ESI-MS/MS

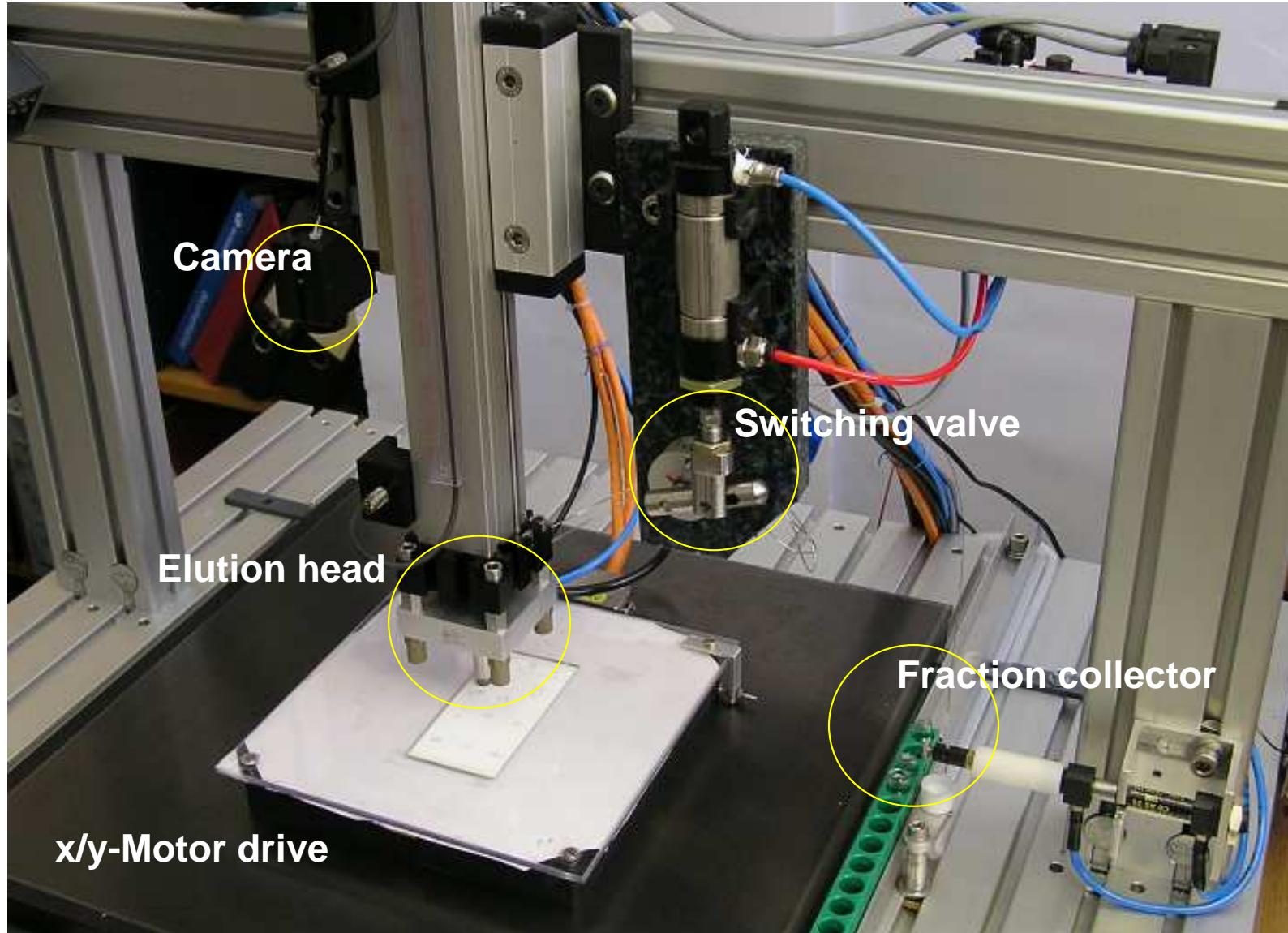


- LOQ better than 20 pg/zone harman (S/N 20)
- Detectability comparable to HPLC/MS

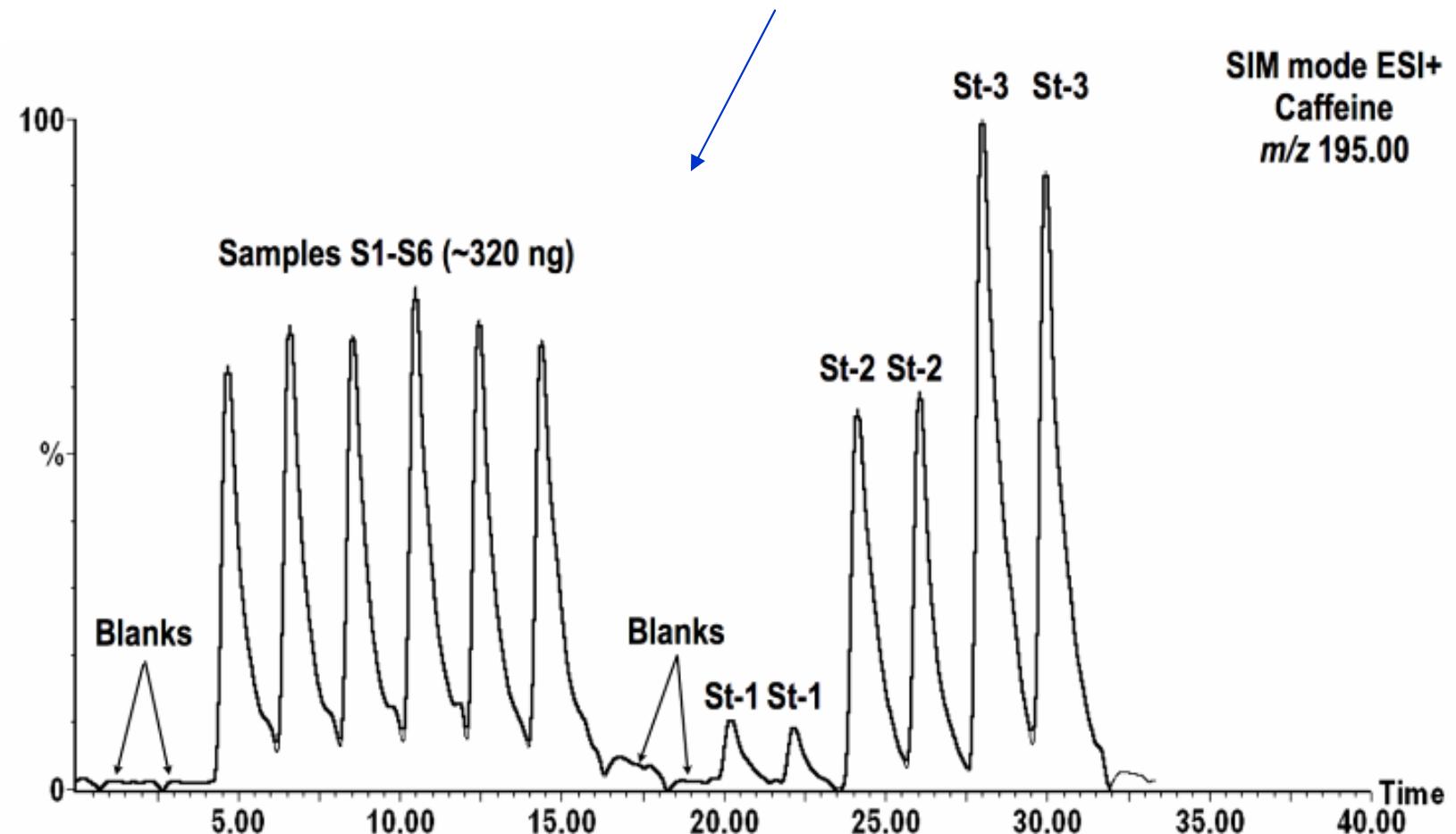
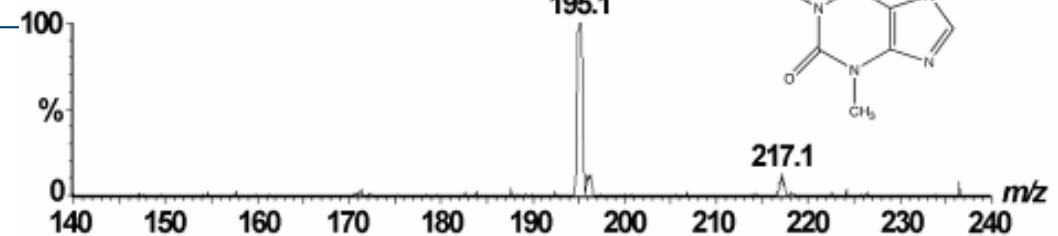
# Elution profiles with different solvents



# Hands-free interface called 'R3D3'

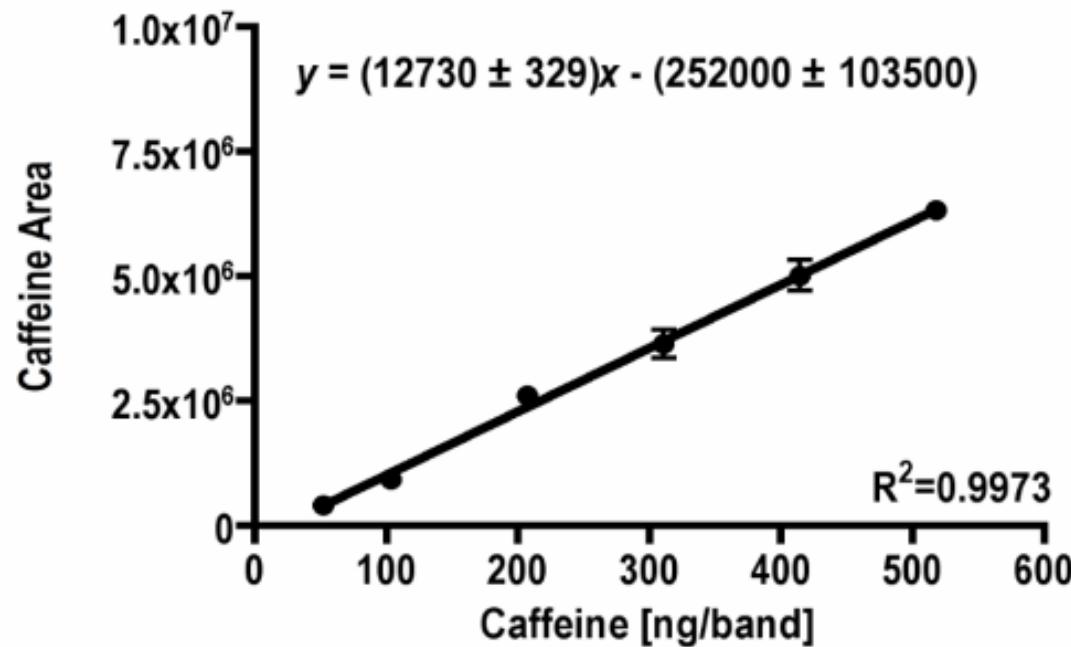


## Elution profiles



## Data of validation without IS

- Repeatability (%RSD, n = 6) in matrix: 5.6 %
- Linearity R<sup>2</sup>: 0.9973



H. Luftmann, M. Aranda, G. Morlock, Rapid Commun Mass Spectrom 21 (2007) 3772-3776

# Analysis of samples containing caffeine

Sample	Pharmaceutical mean $\pm$ SD (mg/tablet)	Energy drink mean $\pm$ SD (mg/100 mL)
HPTLC/ESI-MS RSD (%), n = 6	<b>102.09 <math>\pm</math> 5.76</b> (5.6)	<b>32.91 <math>\pm</math> 1.60</b> (4.9)
HPTLC/UV RSD (%), n = 5	<b>101.98 <math>\pm</math> 2.30</b> (2.3)	<b>33.71 <math>\pm</math> 0.96</b> (2.8)
Label	<b>100</b>	<b>32</b>

→ Comparable findings to validated HPTLC/UV methods (F-test, t-test)

# Comparison of automated interfaces

Parameter	Precision %RSD	Linear Response $r^2$
-----------	-------------------	--------------------------

Quantification **without** internal standard

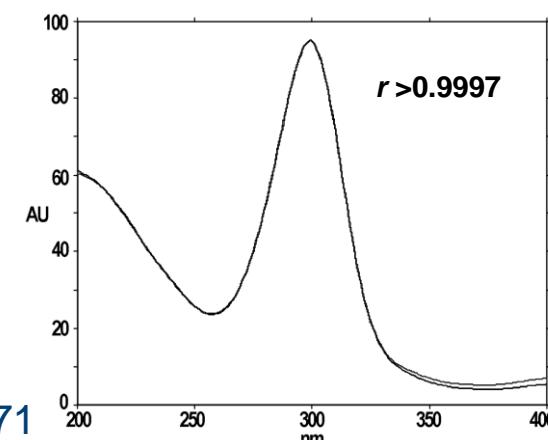
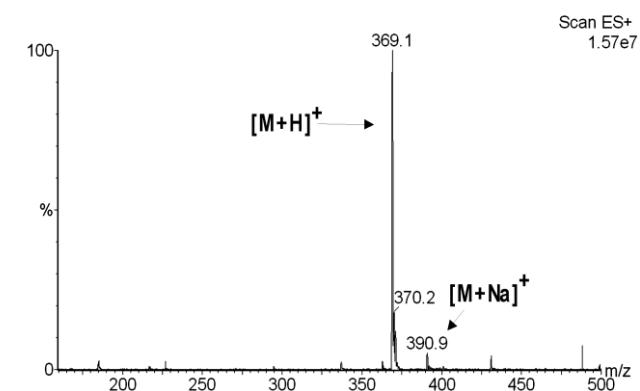
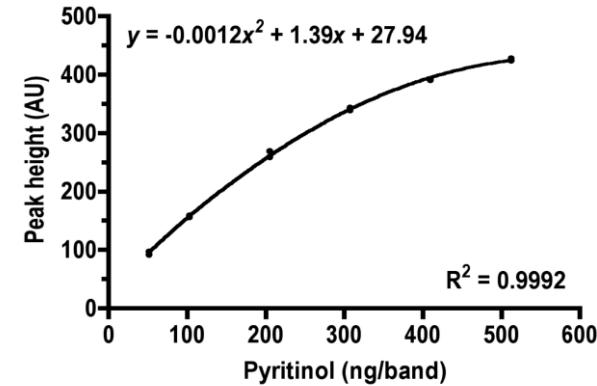
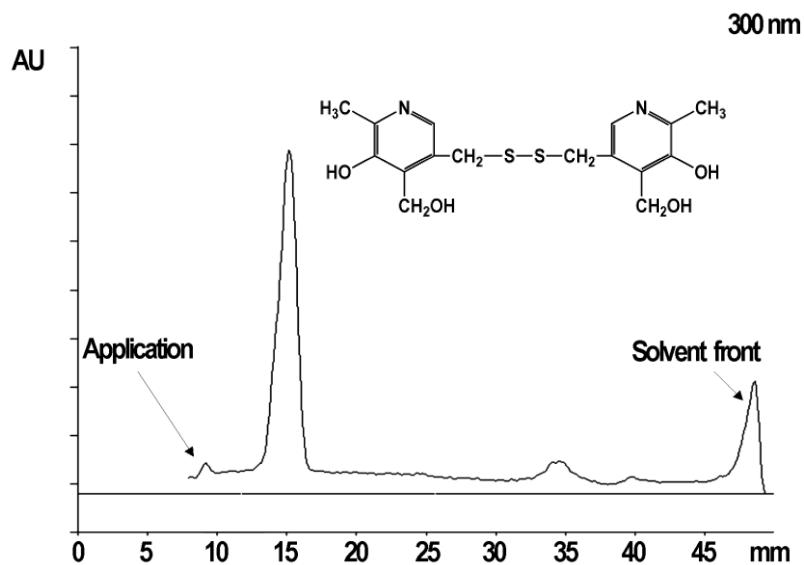
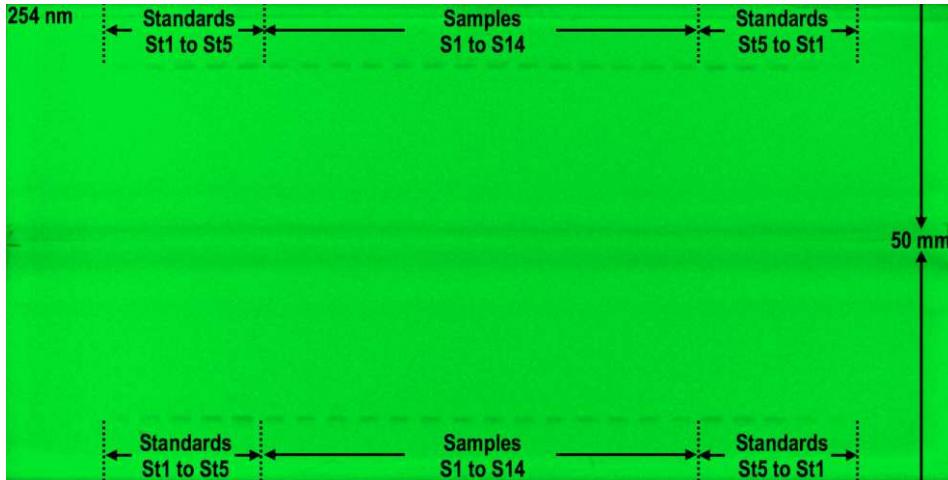


Elution head (autom.)	$\leq 5.6 \%$	0.9973
DESI	$\leq 16.8 \%$	0.95 - 0.98
MALDI	10 %	-
LA-ICP	17 – 41 %	$\geq 0.90$

Quantification **with** internal standard

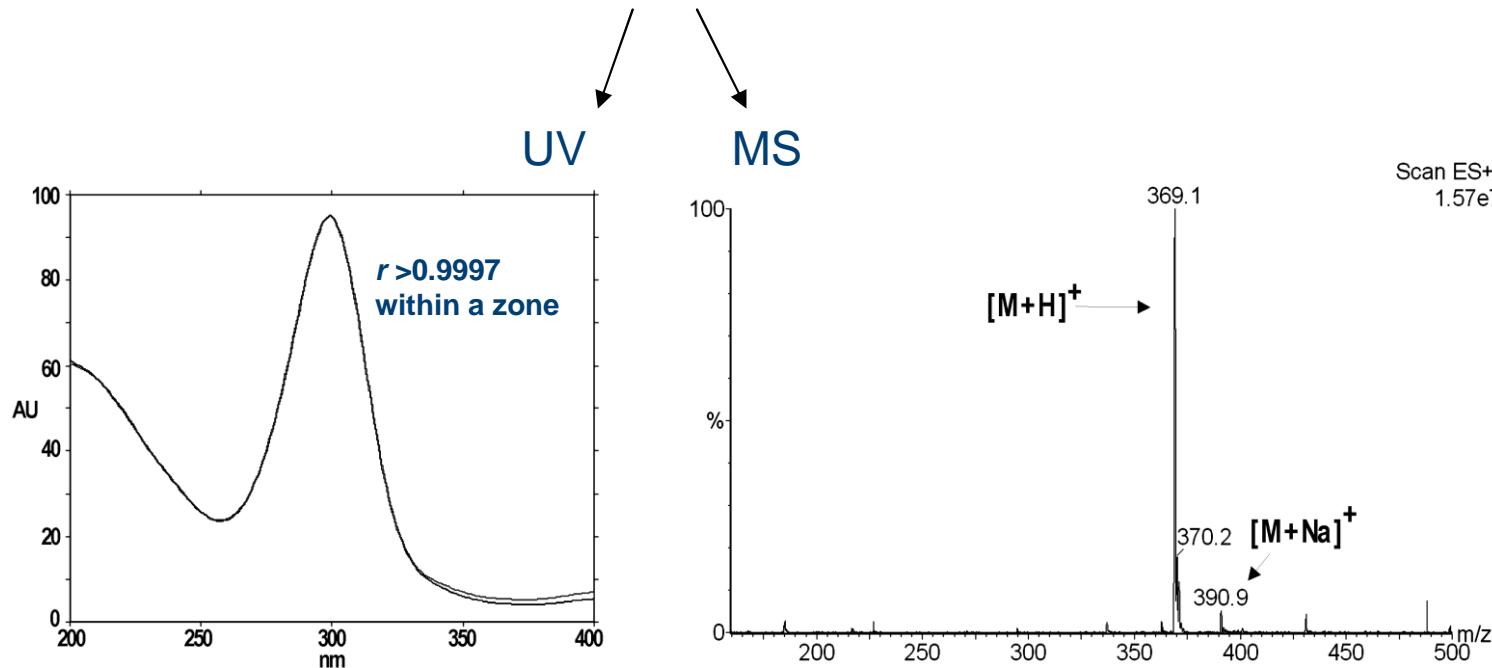
Micro-junction ESI	$\leq 4.4 \%$	0.9999
SALDI/APCI	7 %	0.9991
MALDI	$\leq 8.9 \%$	0.9969
LA-ICP	3 – 40 %	$\geq 0.98$

# Pyridinol in tablets

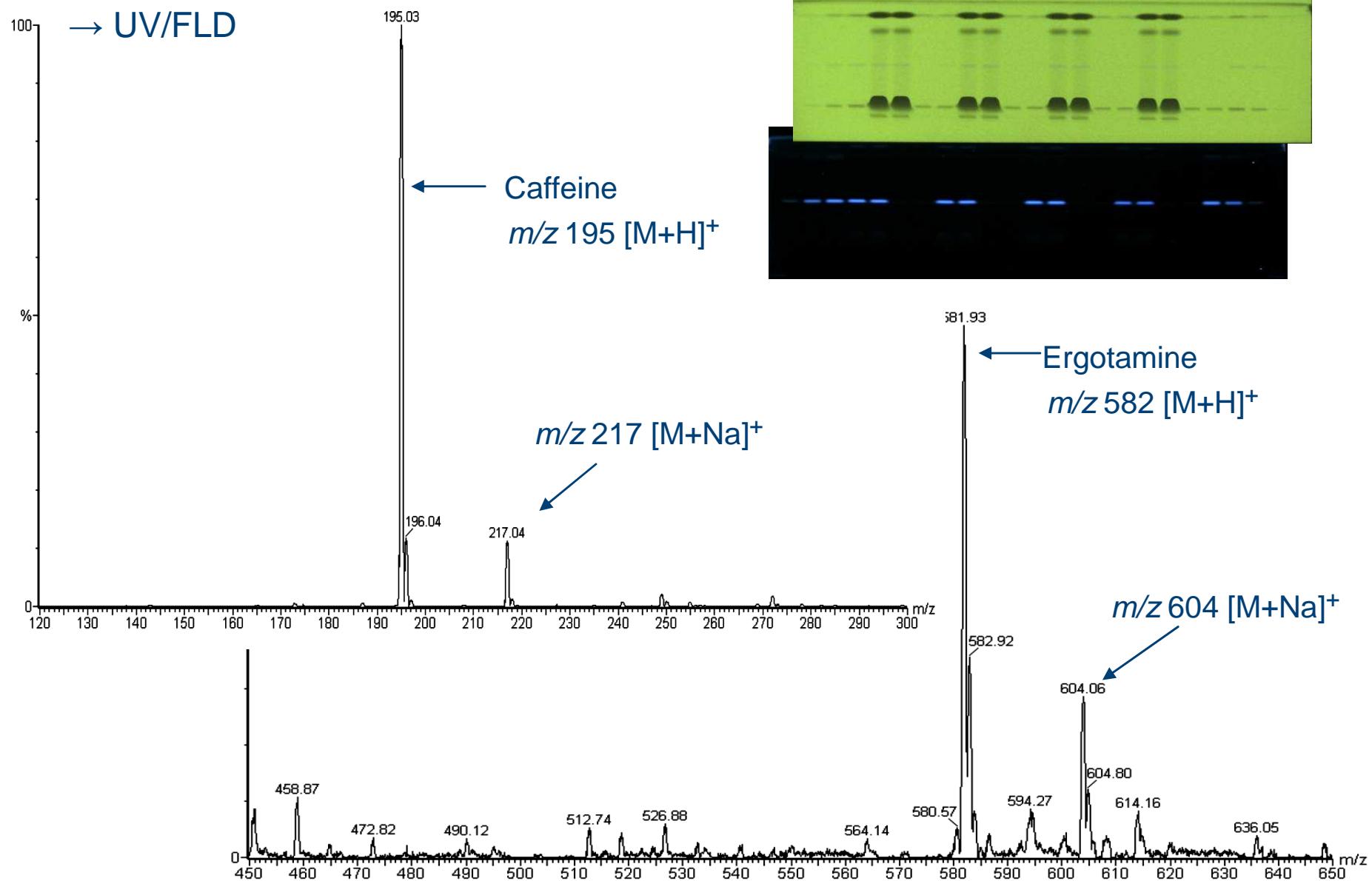


## ...no need for a higher separation power...

- Repeatability (%RSD, n = 6) in matrix: 0.4 %
- Intermediate precision (%RSD, n = 3) in matrix: 2.95 %
- Recoveries of spiked samples (three levels): 98.5 - 101.9 % ( $\pm$  3.6 - 4.7%)
- LOD/LOQ: 0.6/2.0 µg/mL (6/20 ng/band)
- Up to 17 times less mobile phase consumption
- Up to 8 times faster
- Selectivity proven by spectral purity

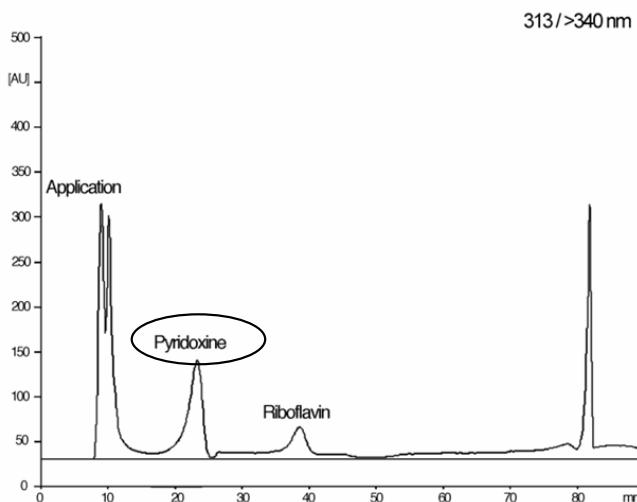
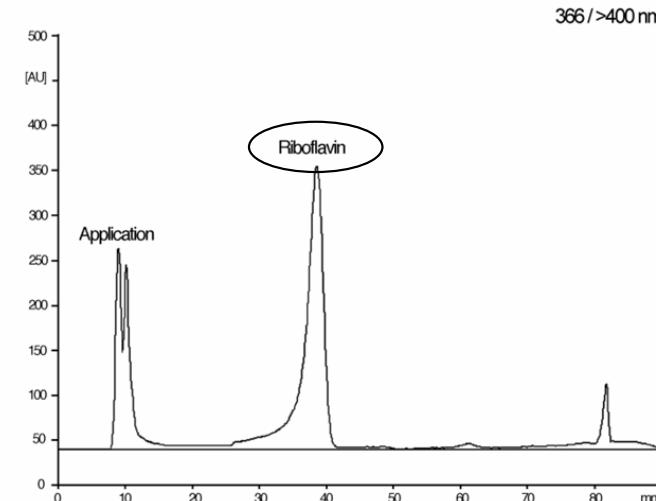
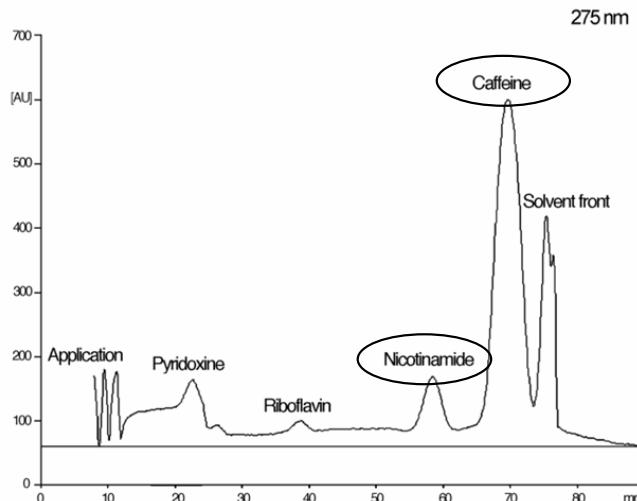


# Caffeine, ergotamine and metamizol in tablets

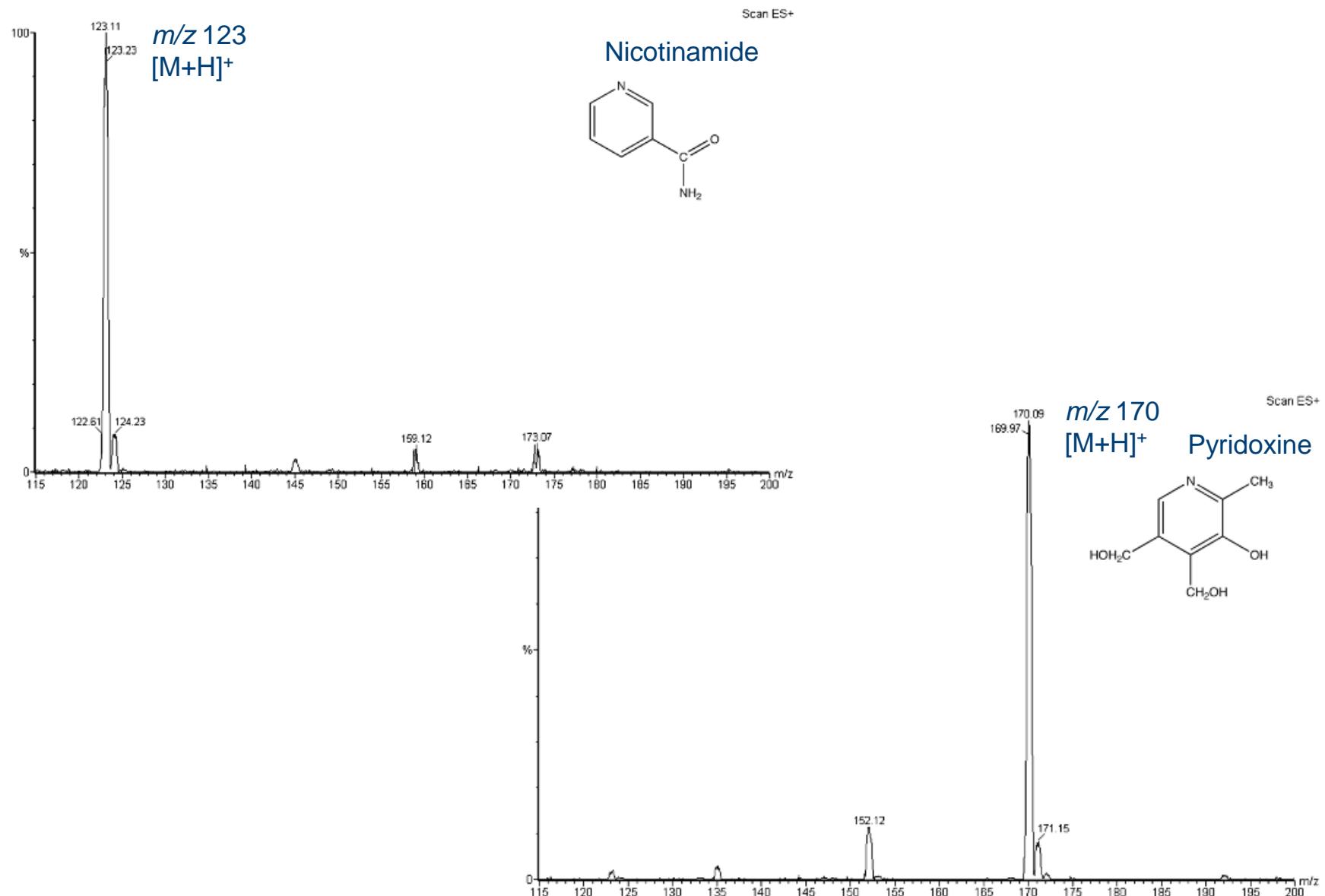


# Active ingredients in energy drinks

Simultaneous determination by MWL scan (UV/FLD) → Derivatization → Vis

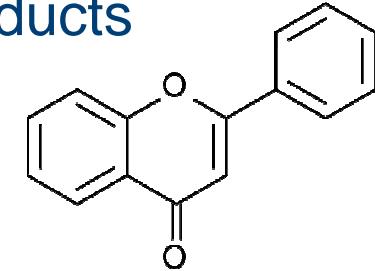


# Confirmation by MS

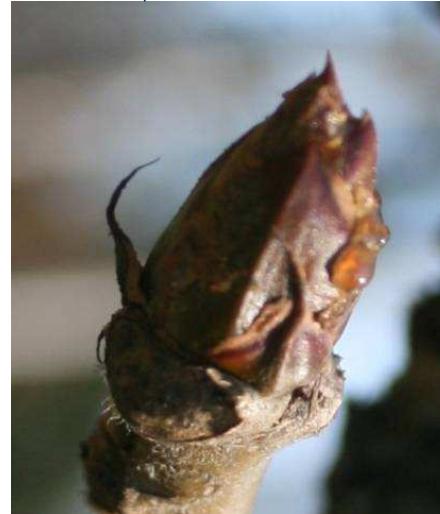


# Are there different types of German propolis?

- Use as food supplement and for cosmetic products
- Flavonoid/phenolic acid profil



Resinated buds, e.g. of  
poplar and horse chestnut

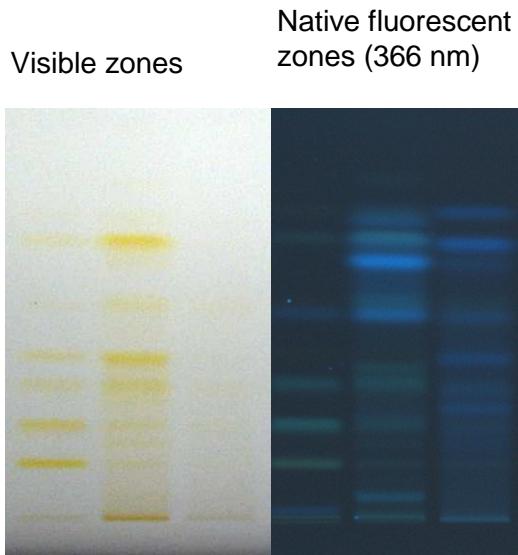


German propolis

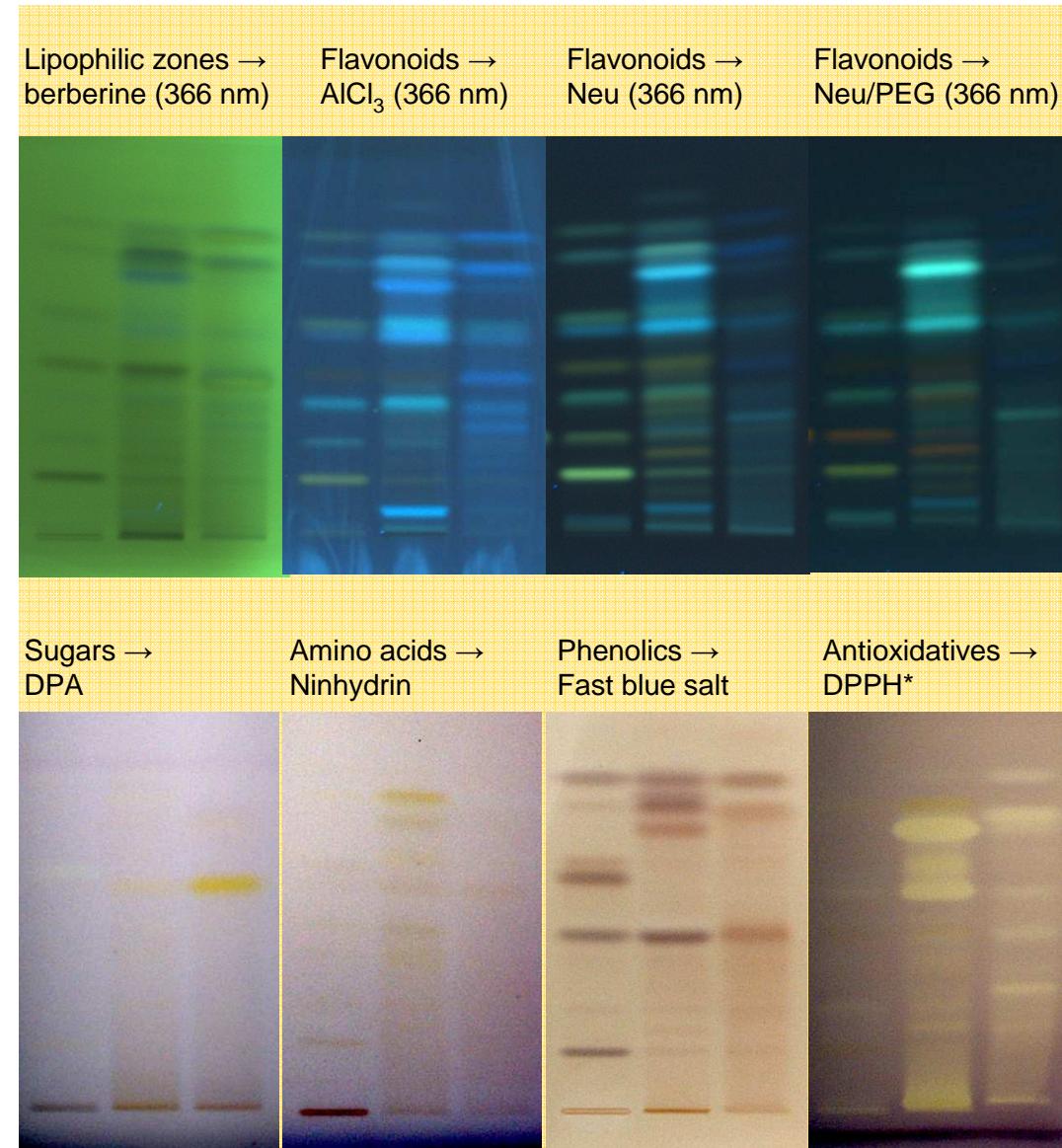


# Information obtained from one plate

## Fast characterization of samples by HPTLC

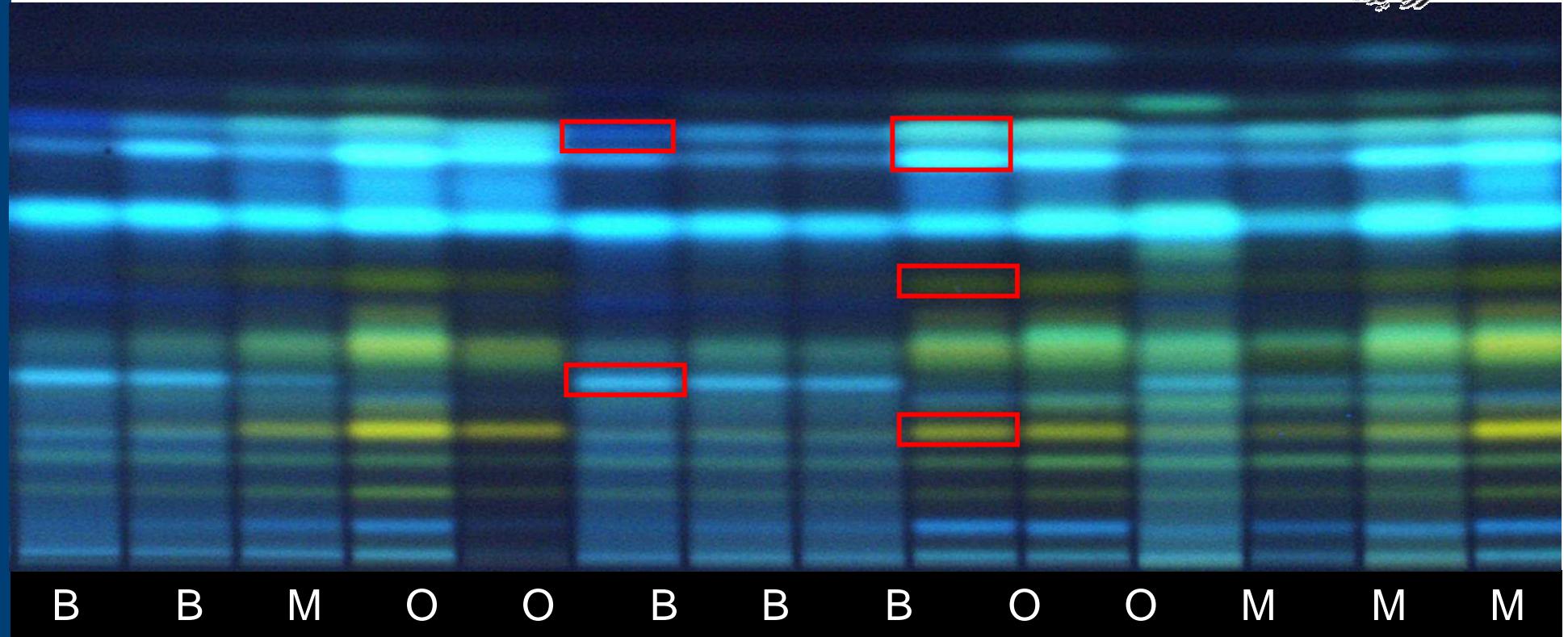


## Selective derivatizations



# Flavonoid/phenolic acid profil

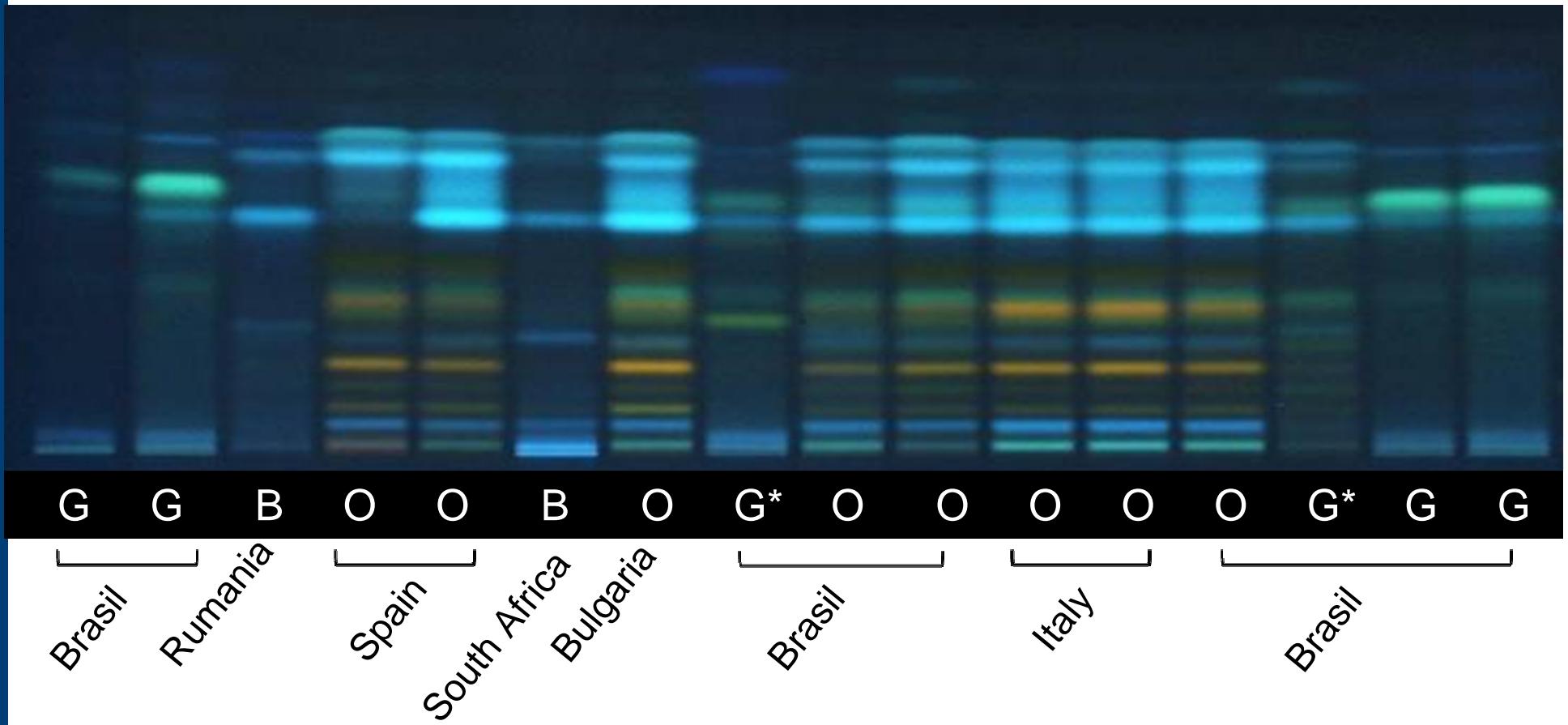
→ German propolis



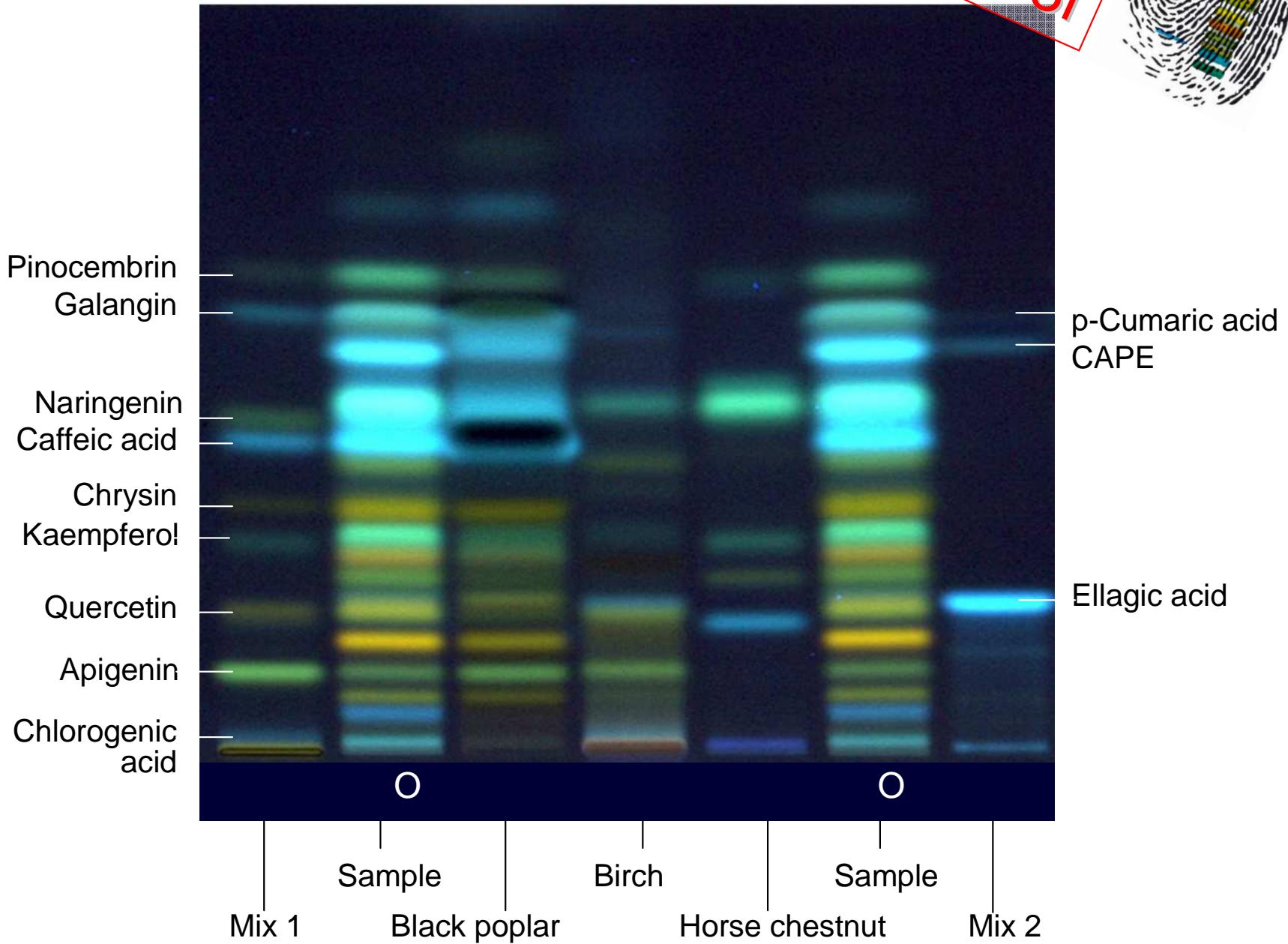
- Screening of 100 samples showed characteristic marker compounds
- 2 types of propolis

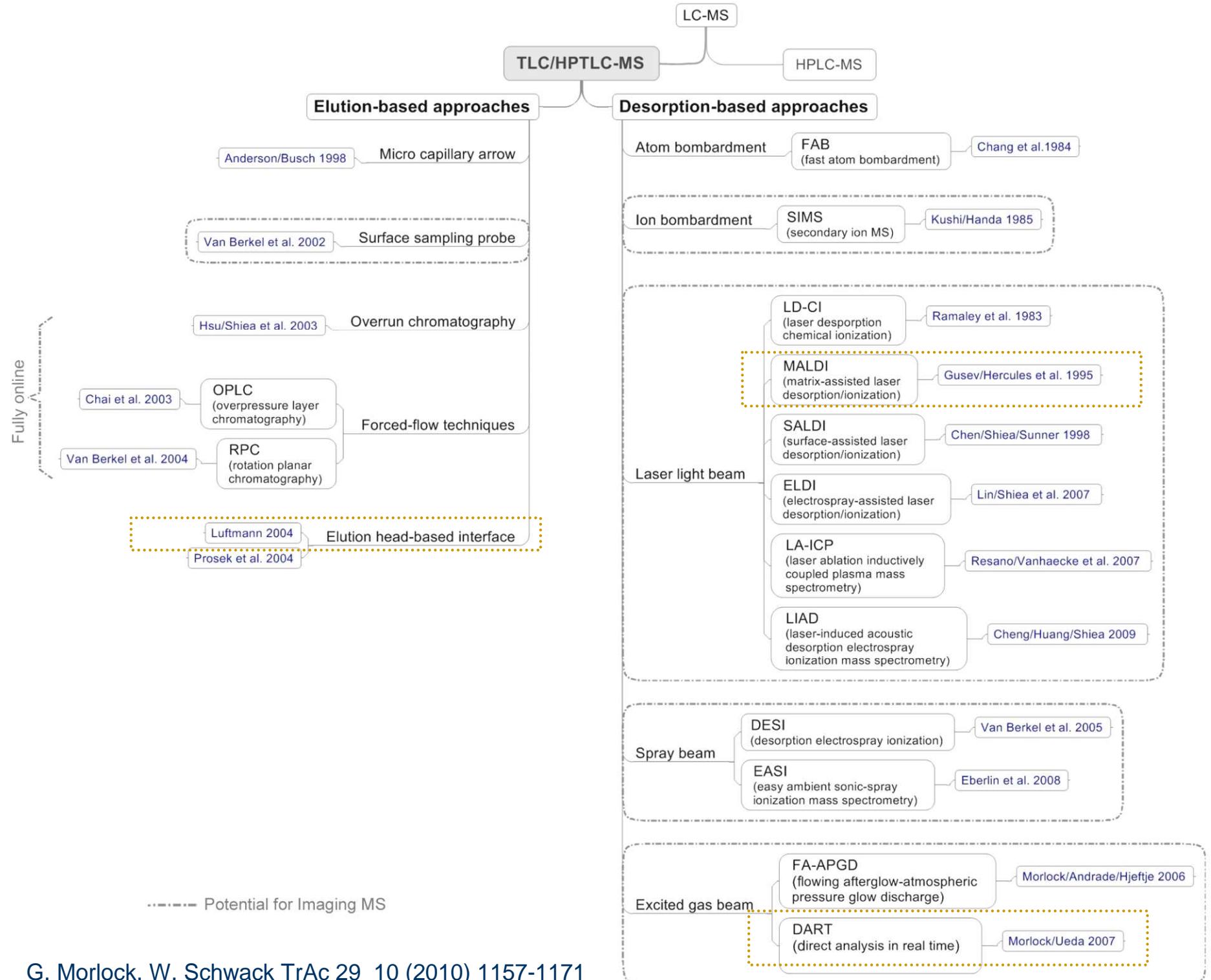
# Flavonoid/phenolic acid profil

→ Foreign propolis: additional green type



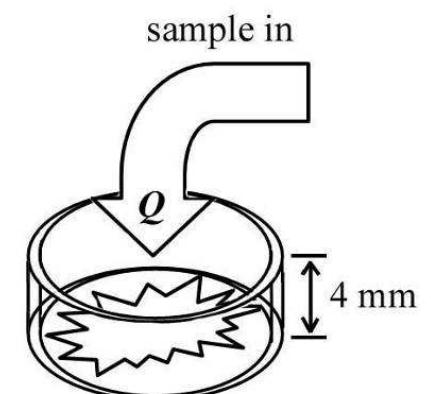
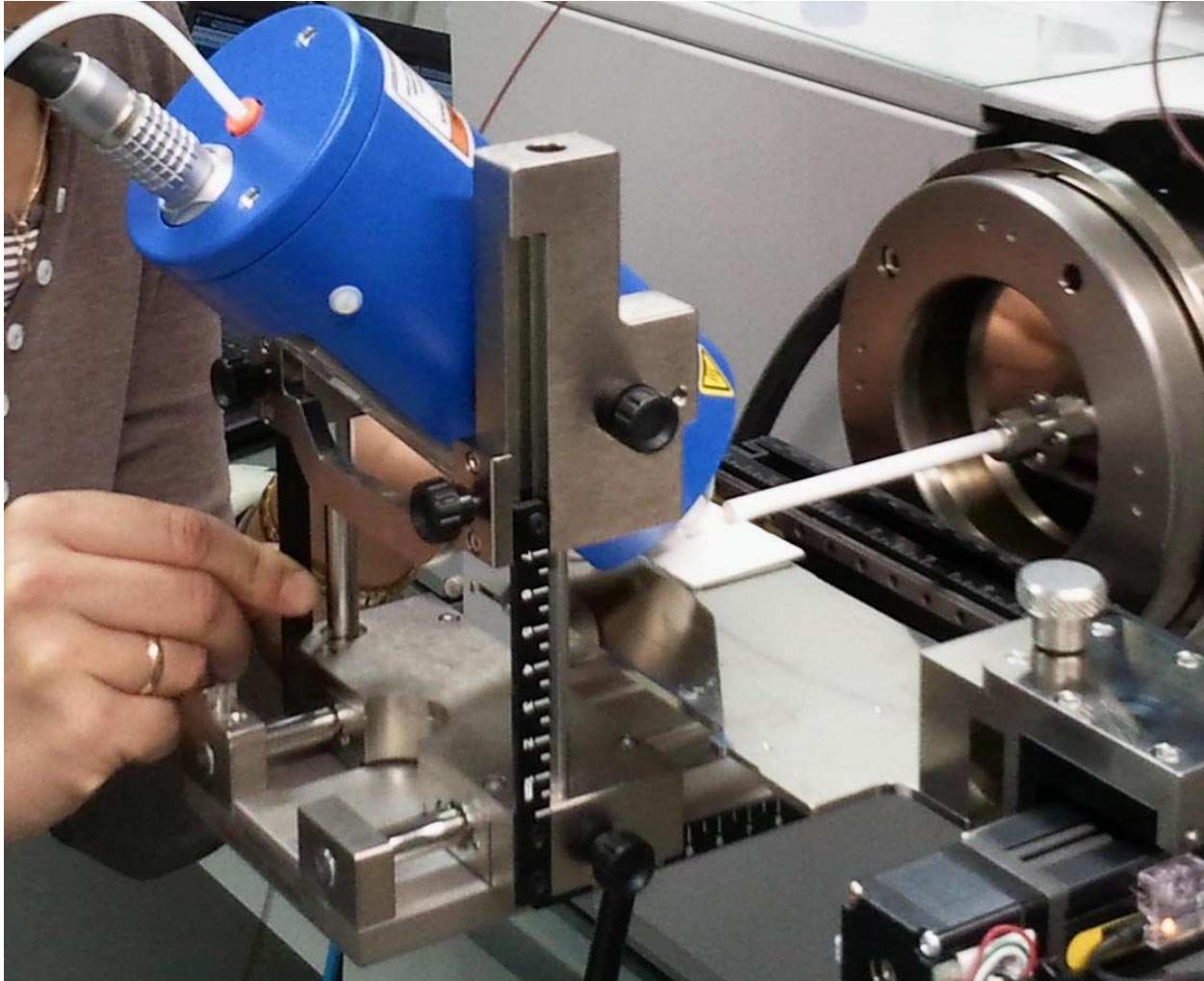
# Plant origin in type O?





# Desorption-based HPTLC-MS → HPTLC-DART-MS

Direct Analysis in Real Time Mass Spectrometry



E. Chernetsova, G. Morlock, *Rapid Commun Mass Spectrom*, in print

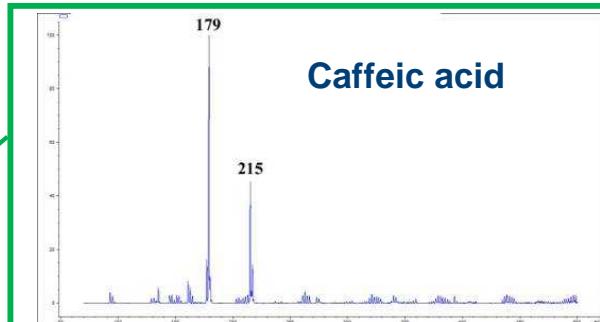
# Confirmation of marker compounds by MS

Lecture 9b

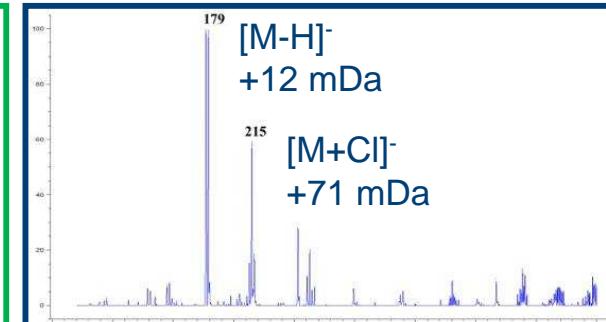
ESI-MS full scan spectra



Spectra of standard



Spectra of sample



EIC of DART-MS



$m/z$  179 ( $[M-H]^-$  of caffeoic acid)



G. Morlock, W. Schwack, *TrAC* 2910 (2010) 1157-1171  
E. Chernetsova et al. *Mass Spectrom Rev* 2011, in print

# Confirmation of marker compounds by MS

Lecture 9c

## TLC-MS VERSUS TLC-LC-MS FINGERPRINTS OF HERBAL EXTRACTS. PART II. PHENOLIC ACIDS AND FLAVONOIDS

Mieczysław Sajewicz,<sup>1</sup> Dorota Staszek,<sup>1</sup> Maja Natić,<sup>1,2</sup> Łukasz Wojtal,<sup>1</sup> Monika Waksmundzka-Hajnos,<sup>3</sup> and Teresa Kowalska<sup>1</sup>

<sup>1</sup>Institute of Chemistry, University of Silesia, Katowice, Poland

<sup>2</sup>Faculty of Chemistry, University of Belgrade, Belgrade, Serbia

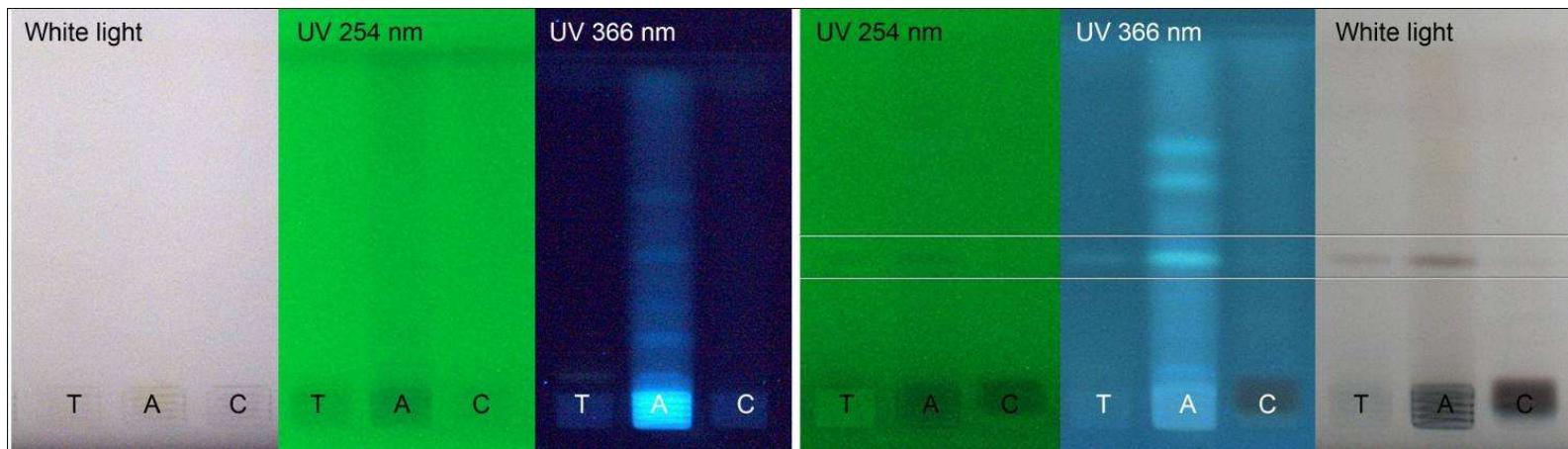
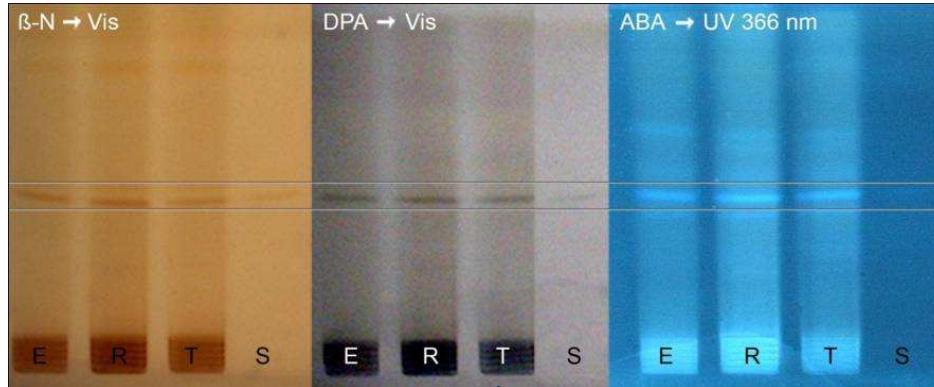
<sup>3</sup>Department of Inorganic Chemistry, Medical University of Lublin, Lublin, Poland

- In the previous paper from this series, we proposed mass spectrometric fingerprinting of a complex and volatile botanical sample upon an example of the essential oil derived from *Salvia lavandulifolia*. In that paper, we compared two variants of fractionation of such a mixture. A simpler one-dimensional variant consisted of the low-temperature thin-layer chromatographic fractionation coupled with mass spectrometric fingerprinting of each separated fraction (1D LT TLC-MS). A more sophisticated variant was the two-dimensional liquid chromatographic system

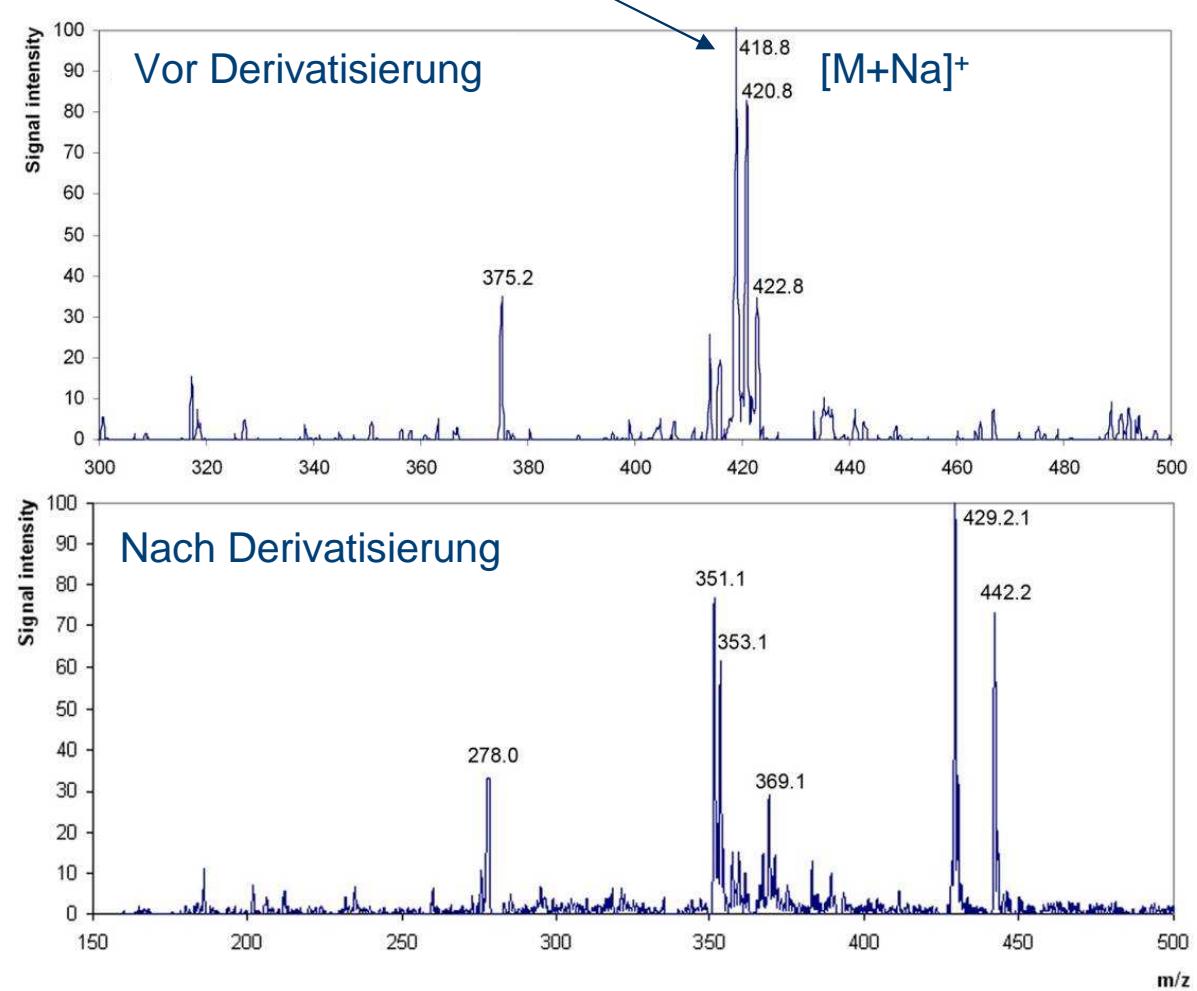
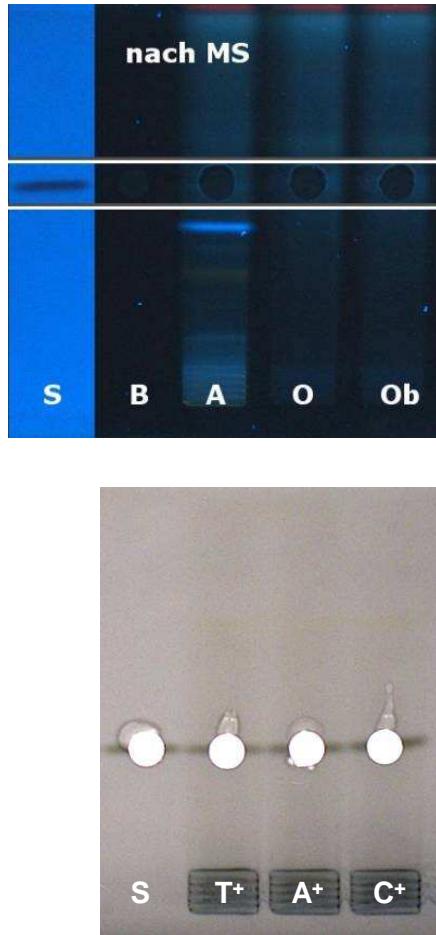
# More information about unknown samples

Sucralose in sewage and waste water

Variety of derivatization options



# Confirmation by HPTLC-MS

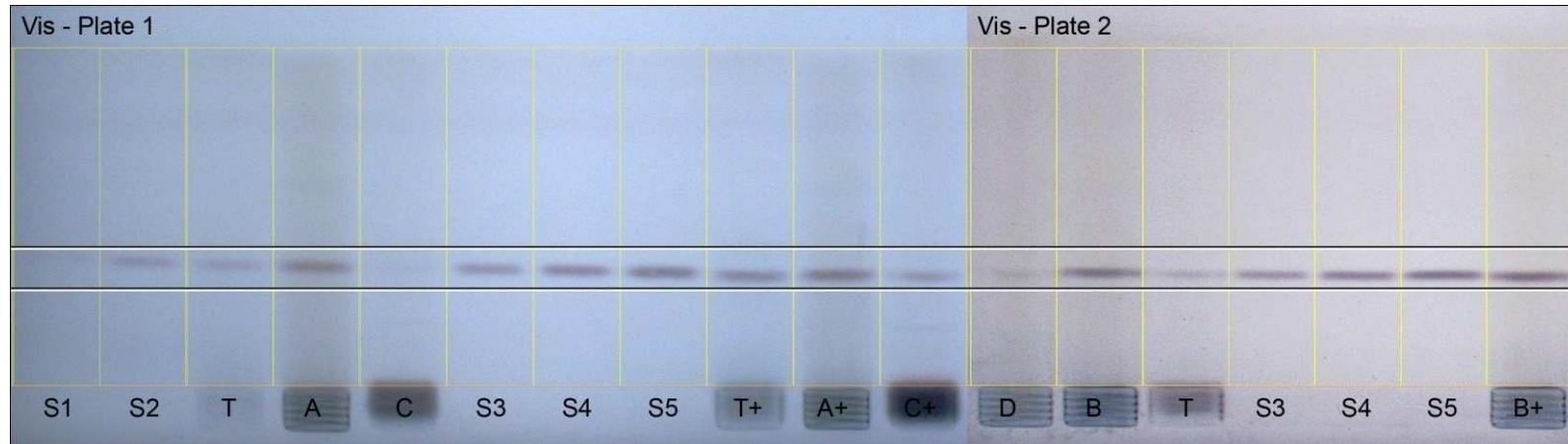


# Analysis of sewage effluent and river water



HPLC-TOF or -MS/MS with isotopically labeled standard

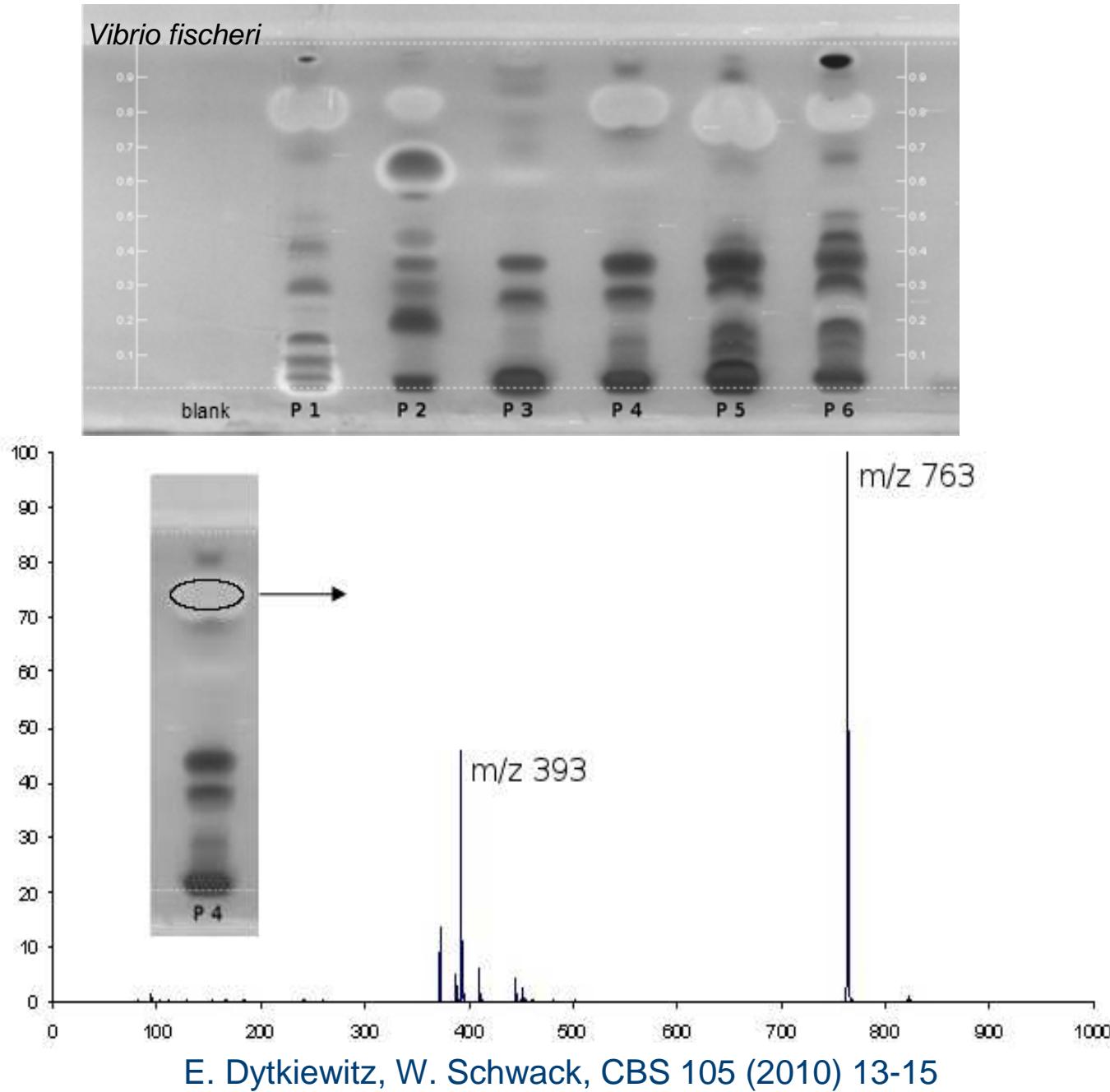
...or HPTLC-Vis?



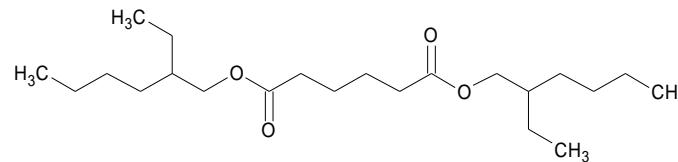
Mean value (ng/L)	Sample A	Sample B	Sample C	Sample D
HPLC-TOF or -MS/MS (n = 6 laboratories)	5869	7302	186	200
HPTLC-Vis (n = 2)	5863	7034	247	218

# Additives in food packaging foils

Poster 8f

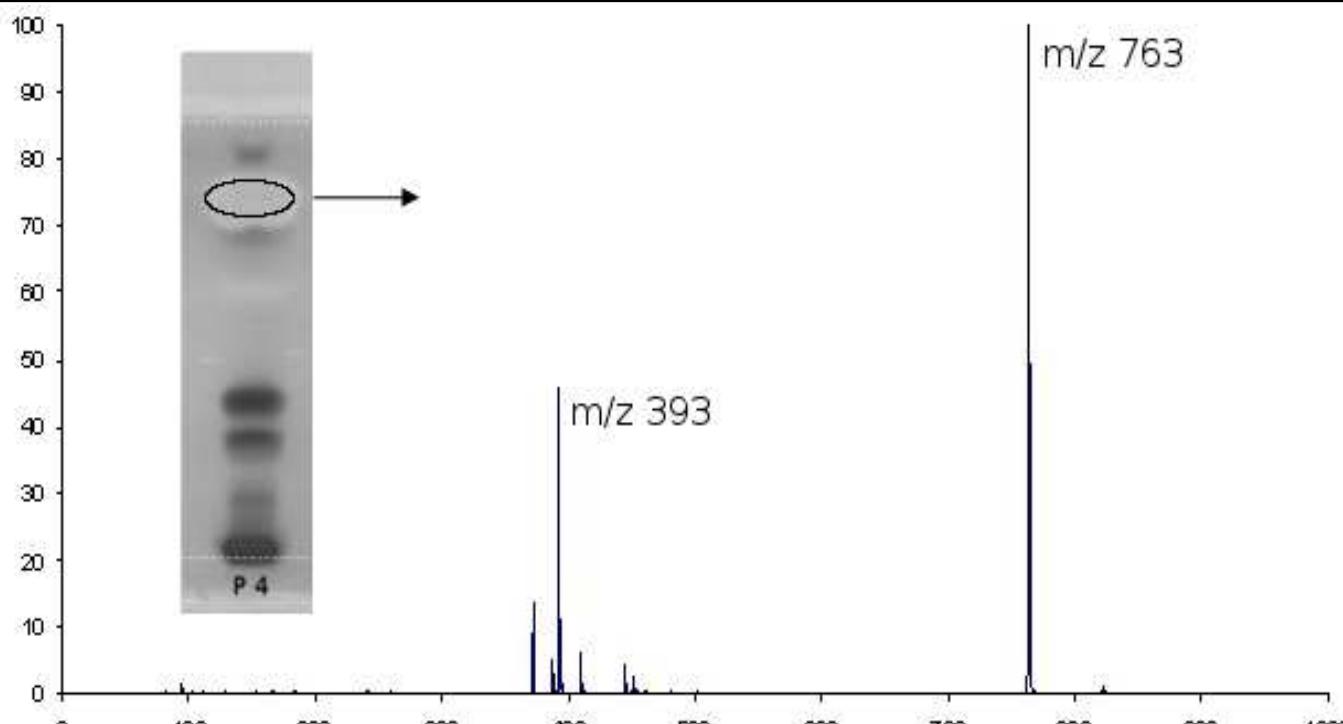


# Detection of additives in polymer packaging foils

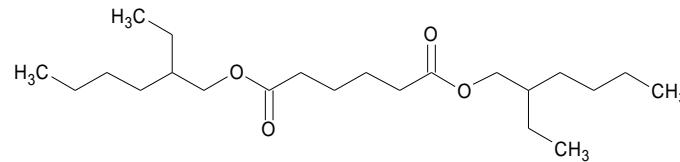


Bis-2-ethylhexyladipate

MS signals of	Mass determined	Mass theoretical	$\Delta$ (ppm)	Sum formula	Assignment
HPTLC zone	393,2985	393,2981	-1,0691	C <sub>22</sub> H <sub>42</sub> O <sub>4</sub> Na	[M+Na] <sup>+</sup>
	763,6077	763,6064	-1,7164	C <sub>44</sub> H <sub>84</sub> O <sub>8</sub> Na	[2M+Na] <sup>+</sup>

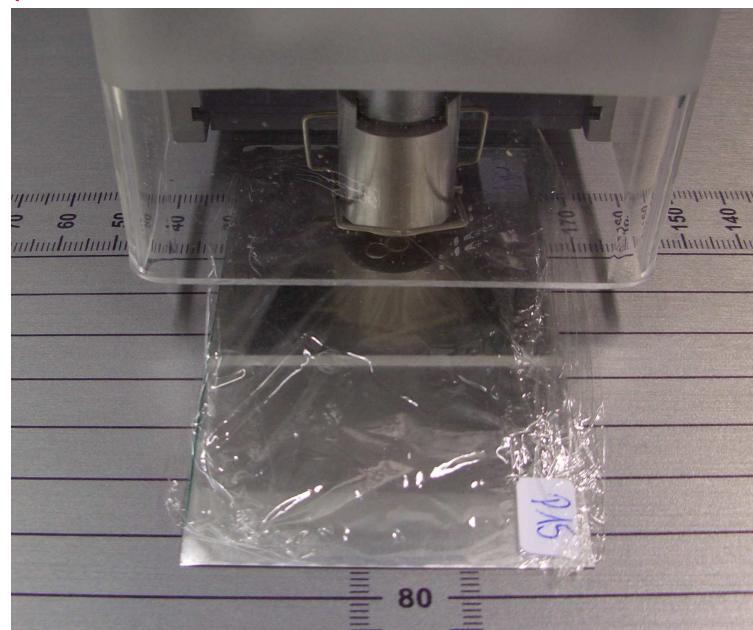


# Detection of additives in polymer packaging foils

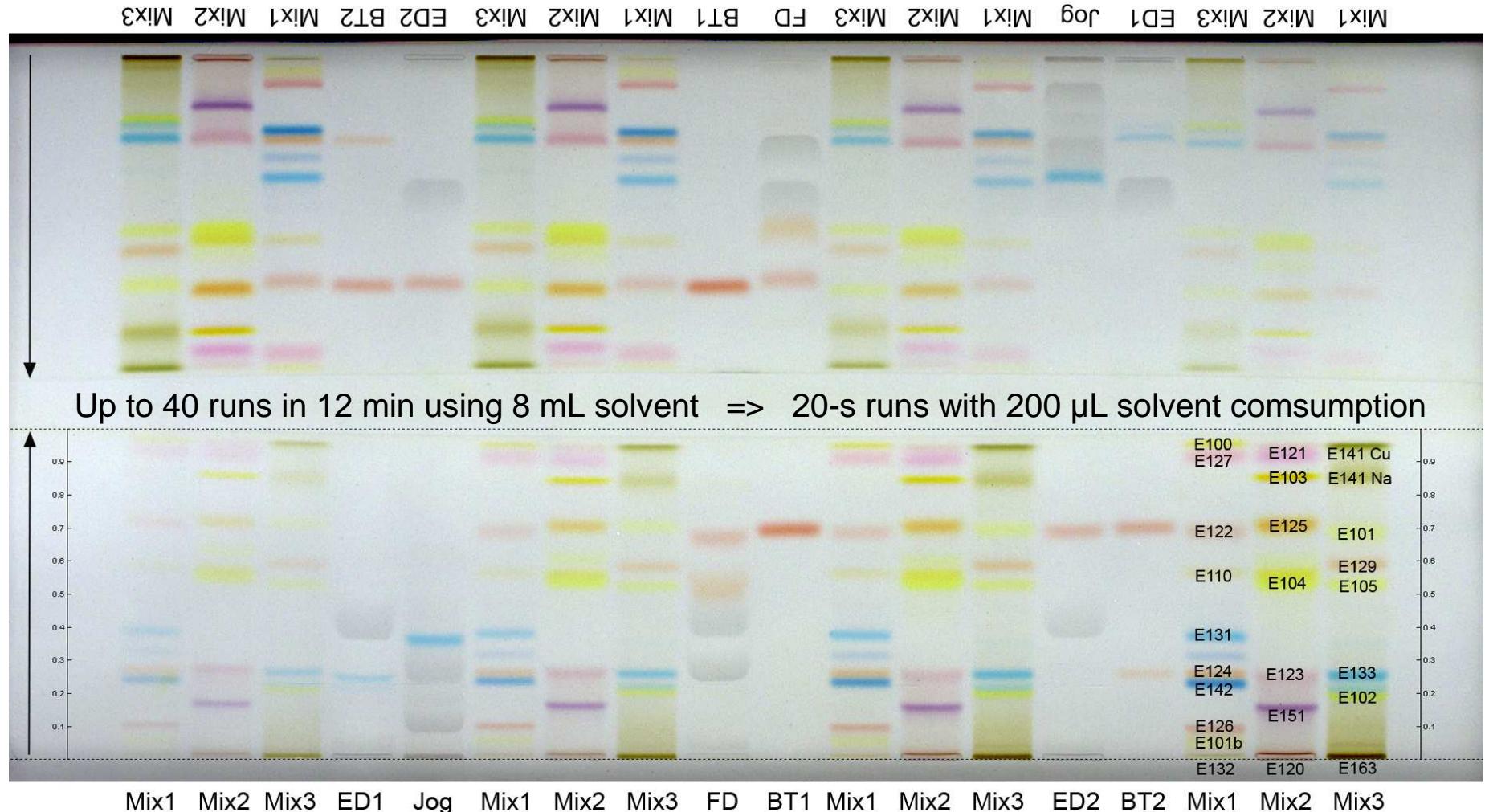


Bis-2-ethylhexyladipate

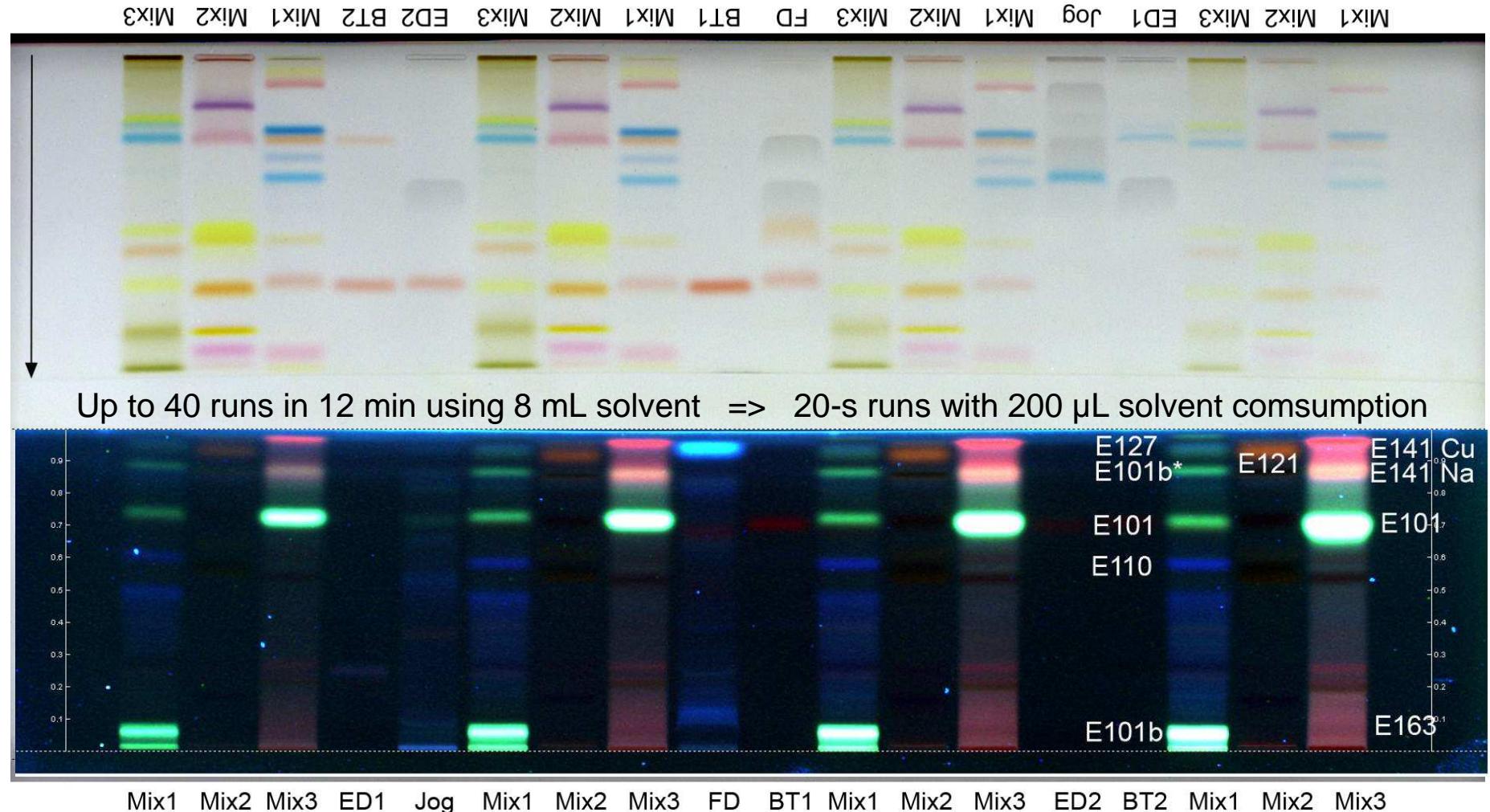
MS signal of	Mass determined	Mass theoretical	$\Delta$ (ppm)	Sum formula	Assignment
<b>Plastic foil</b>	<b>371,3174</b>	<b>371,3161</b>	<b>-3,4071</b>	<b>C<sub>22</sub>H<sub>43</sub>O<sub>4</sub></b>	<b>[M+H]<sup>+</sup></b>
<b>HPTLC zone</b>	393,2985	393,2981	-1,0691	C <sub>22</sub> H <sub>42</sub> O <sub>4</sub> Na	[M+Na] <sup>+</sup>
	763,6077	763,6064	-1,7164	C <sub>44</sub> H <sub>84</sub> O <sub>8</sub> Na	[2M+Na] <sup>+</sup>



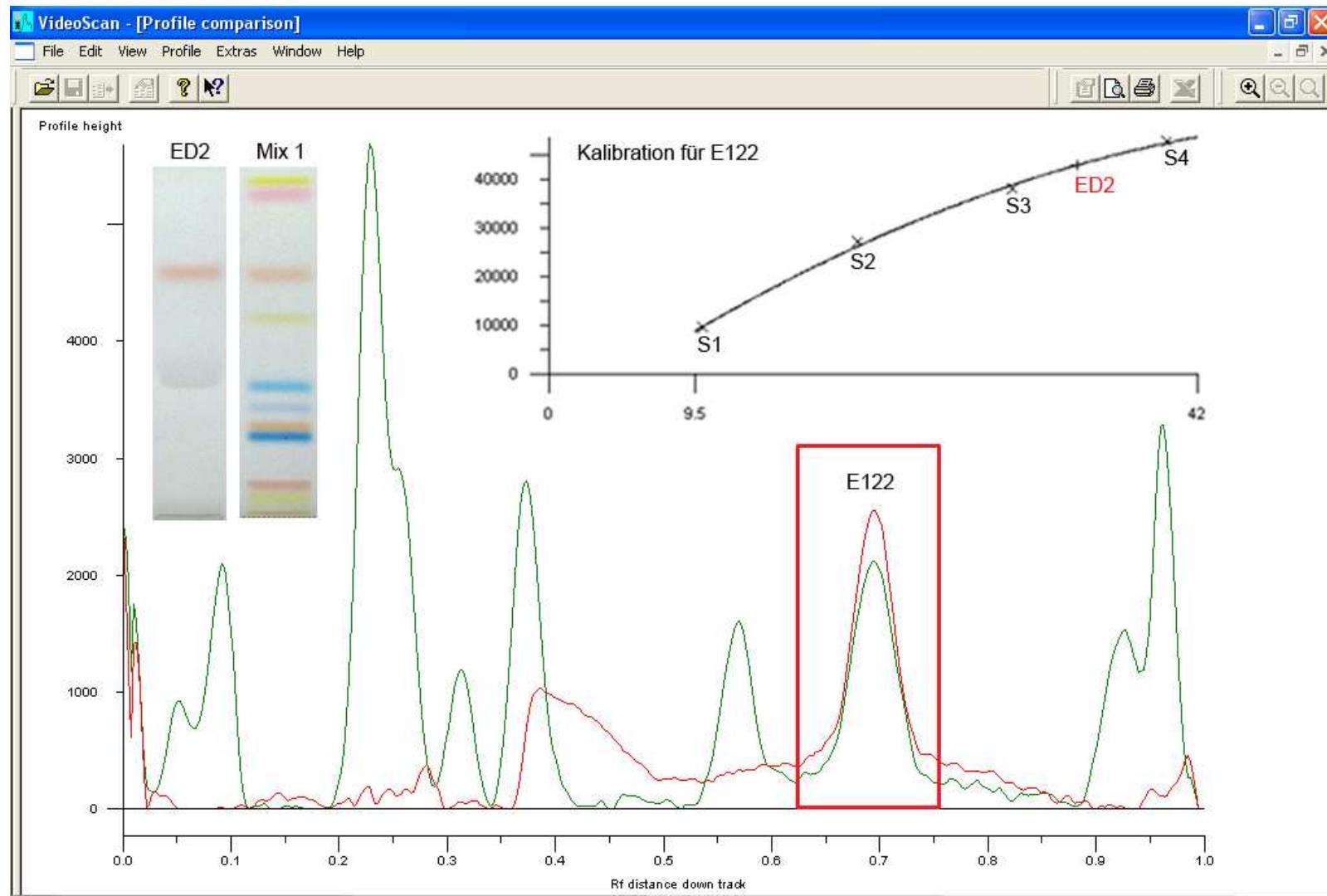
# Dye analysis



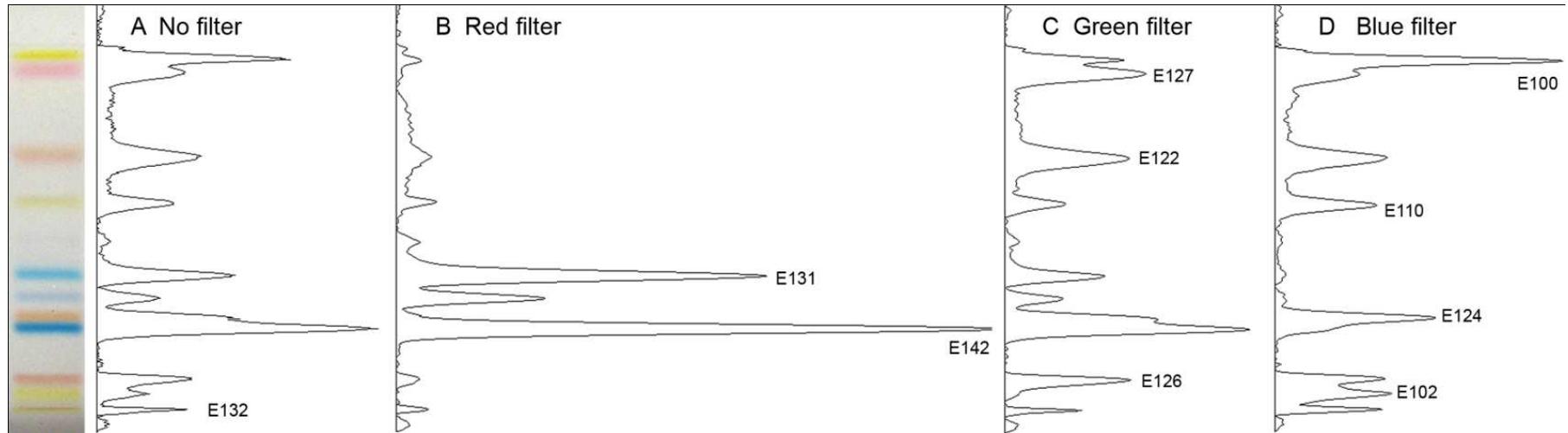
# Dye analysis



# Digital quantification

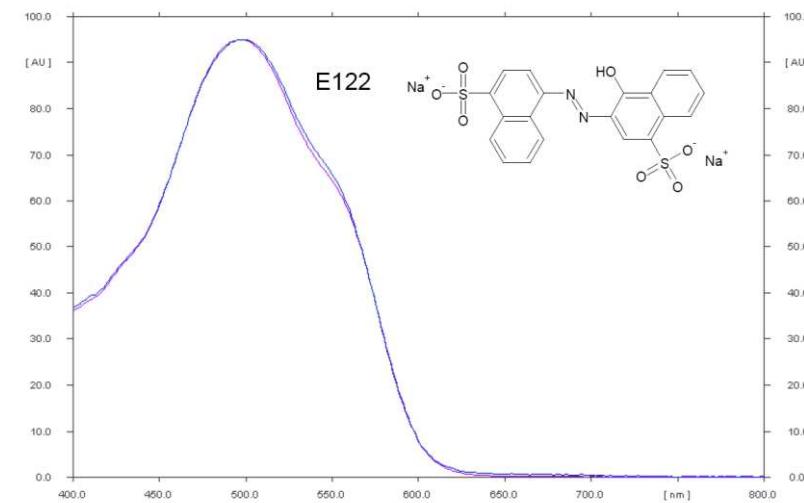
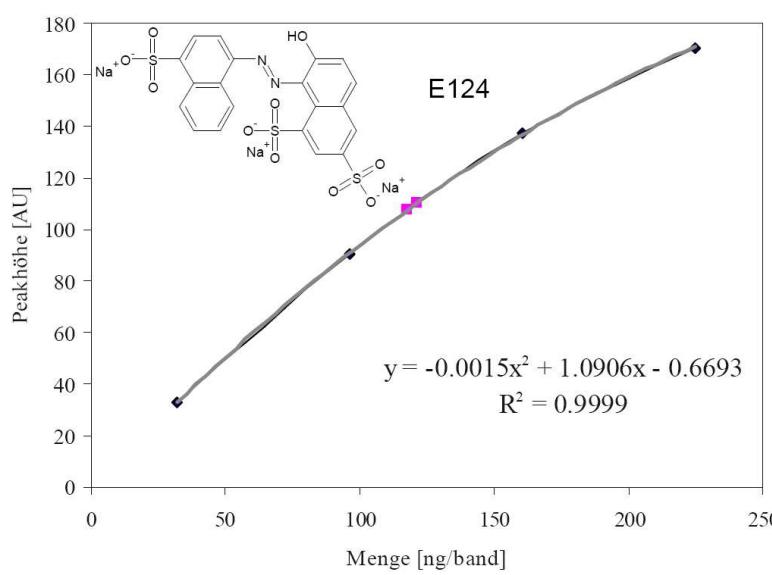
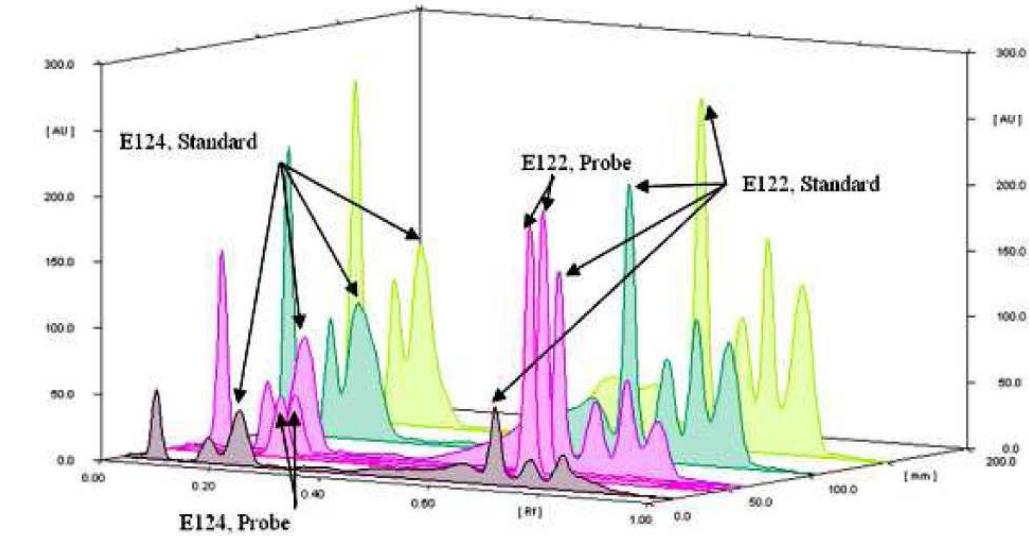
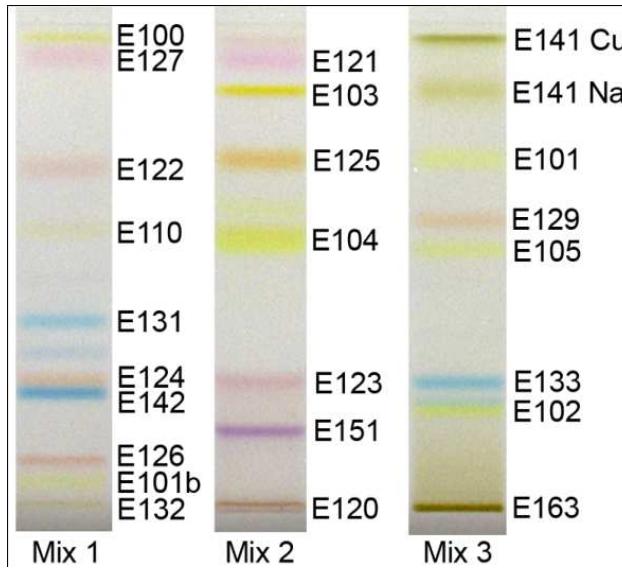


# Digital filters

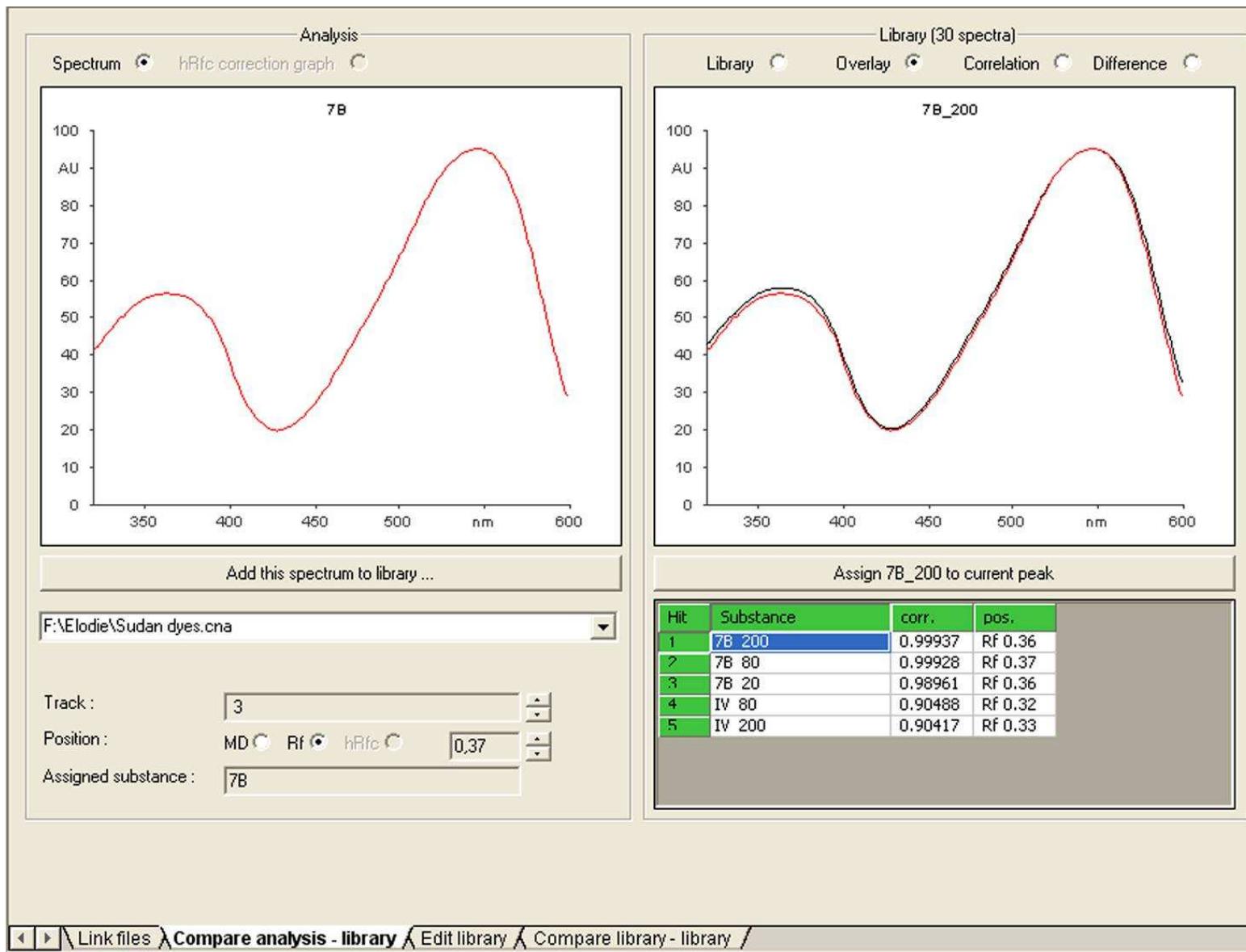


G. Morlock, W. Schwack, Die Aktuelle Wochenschau der GDCh,  
Woche 26 (2009), [www.aktuelle-wochenschau.de](http://www.aktuelle-wochenschau.de)

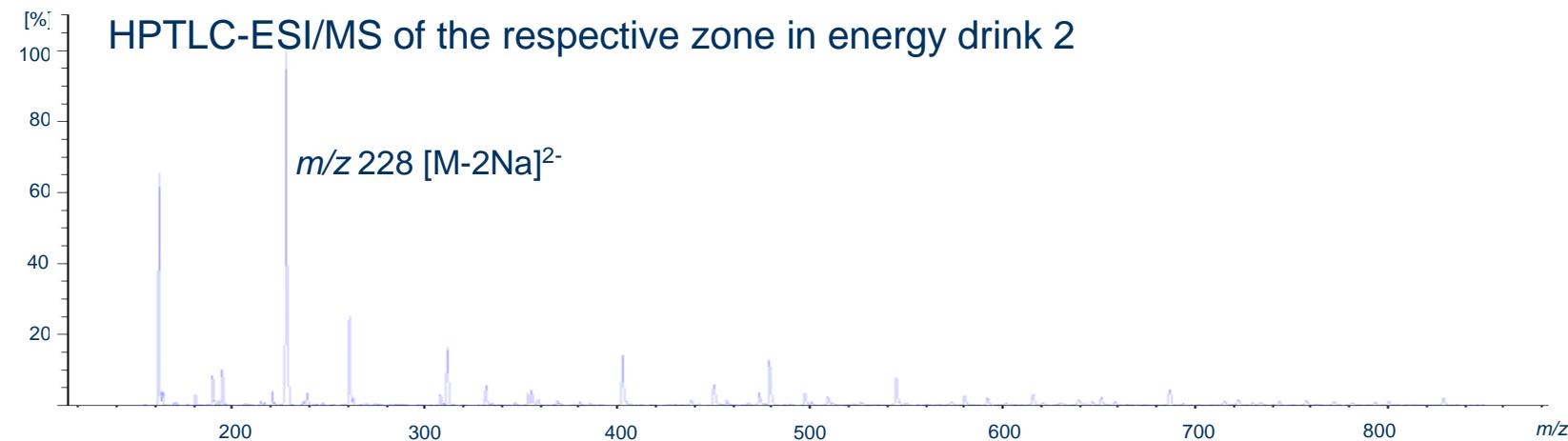
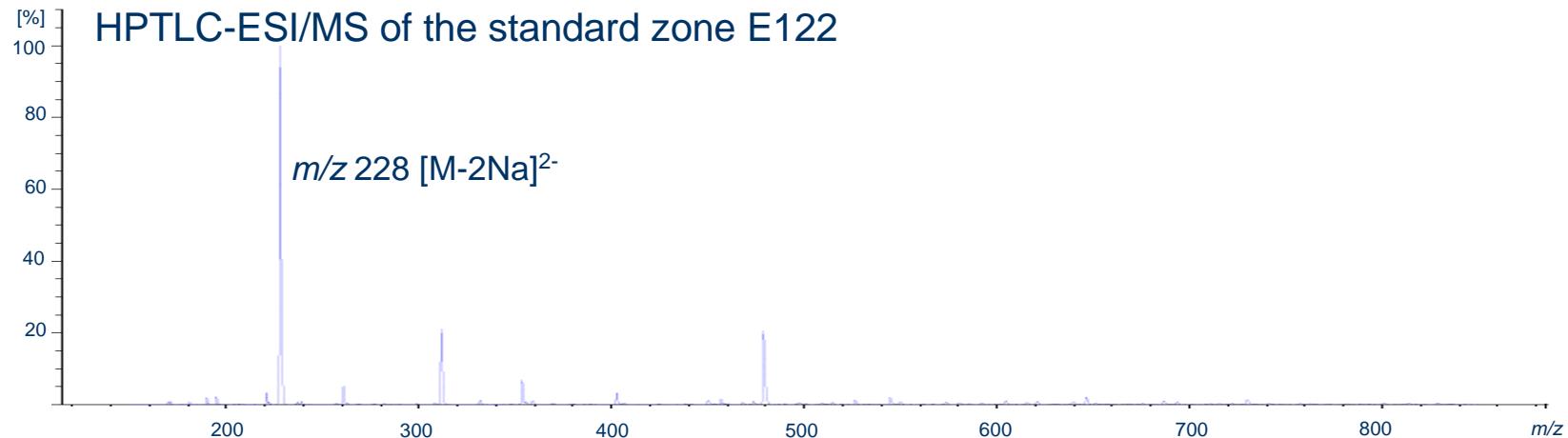
# Dye analysis



# Search in spectra library



# Confirmation by mass spectra



G. Morlock, C. Oellig, J AOAC Int 92 (2009) 547-554

# Rare examples for HPTLC



# Rare examples for HPTLC

Information obtained from a single plate				Identity	
Sample	Dyes found	Concentration calculated	%RSD (n=2)	Spectra correlation (400–800 nm) of standard and sample	Mass signal(s) (full scan, m/z 100–900)
Bakery ink formulation	122	66.4 g/L	0.0	≥ 0.99996	228 [M-2Na] <sup>2-</sup>
	124	13.3 g/L	2.1	≥ 0.99957	279 [M-2Na] <sup>2-</sup> 178 [M-3Na] <sup>3-</sup>
Energy drink 1	133	9.1 mg/L	0.1	≥ 0.99964	373 [M-2Na] <sup>2-</sup>
Energy drink 2	122	76.2 mg/L	3.6	≥ 0.99958	228 [M-2Na] <sup>2-</sup>

# Cost comparison

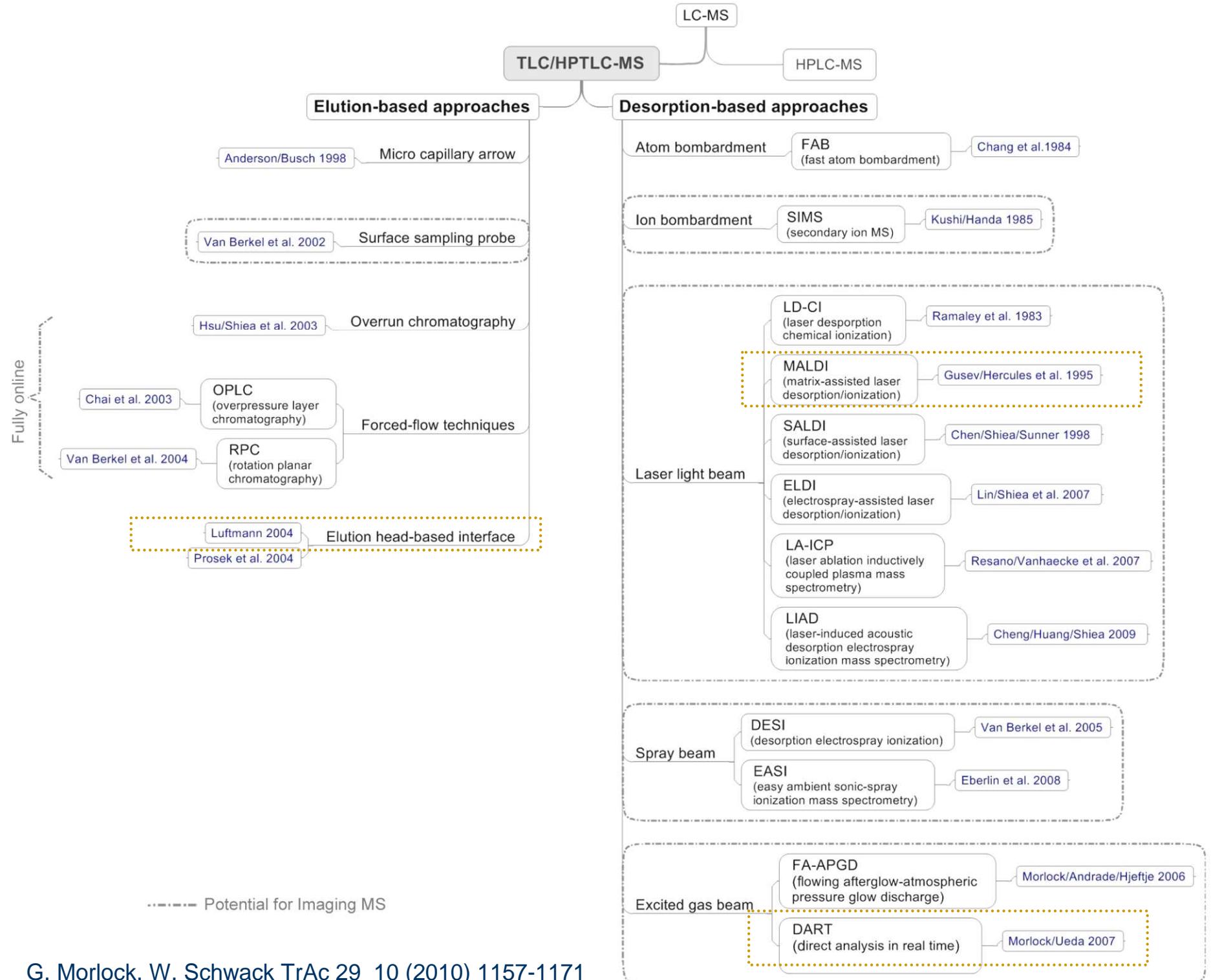
Operating costs/run (€)	HPLC <sup>1</sup>	HPTLC <sup>2</sup>
Mobile phase	0,58	0,003
Stationary phase	0,64	0,11
Disposal	0,04	0,0001
<b>Sum</b>	<b>1,26</b>	<b>0,11</b>

=> 11 x lower

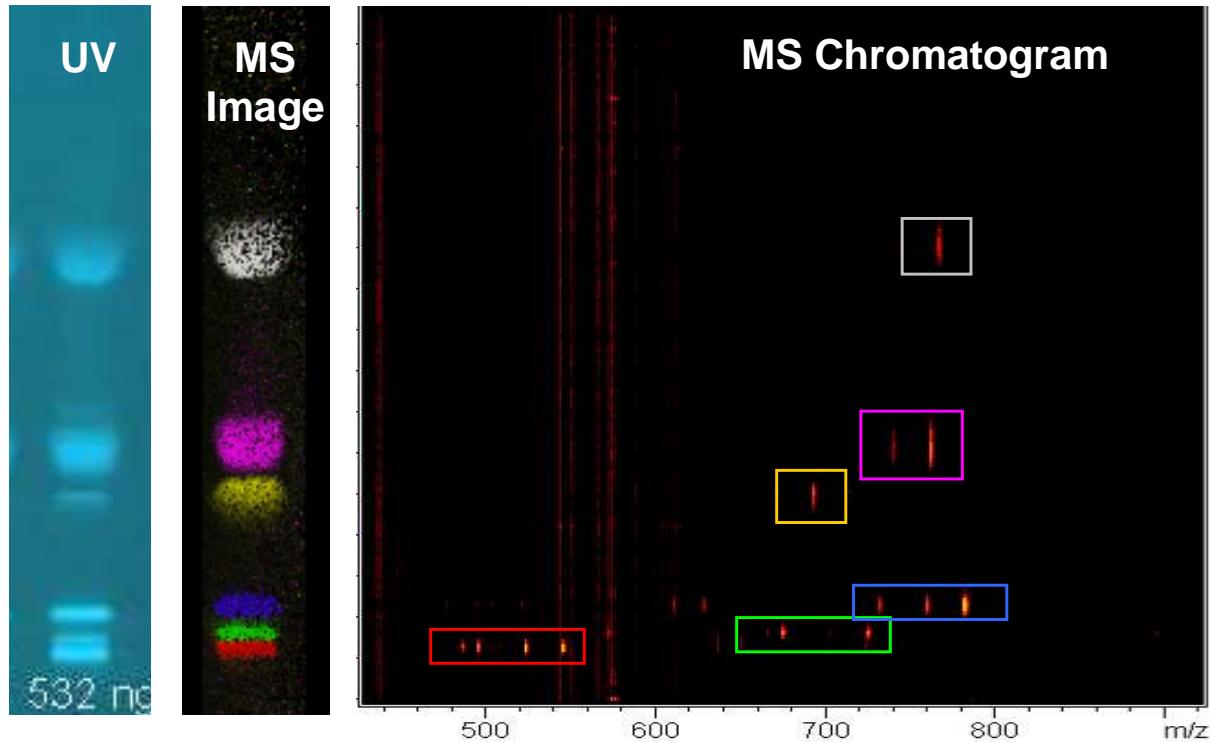
Time/run (min)	HPLC	HPTLC
Application/Injection		0,50
Run time	43	0,20
Detection		0,10
<b>Sum</b>	<b>43</b>	<b>0,80</b>

=> 54 x faster

thereof labor time/40 runs	none	5 min
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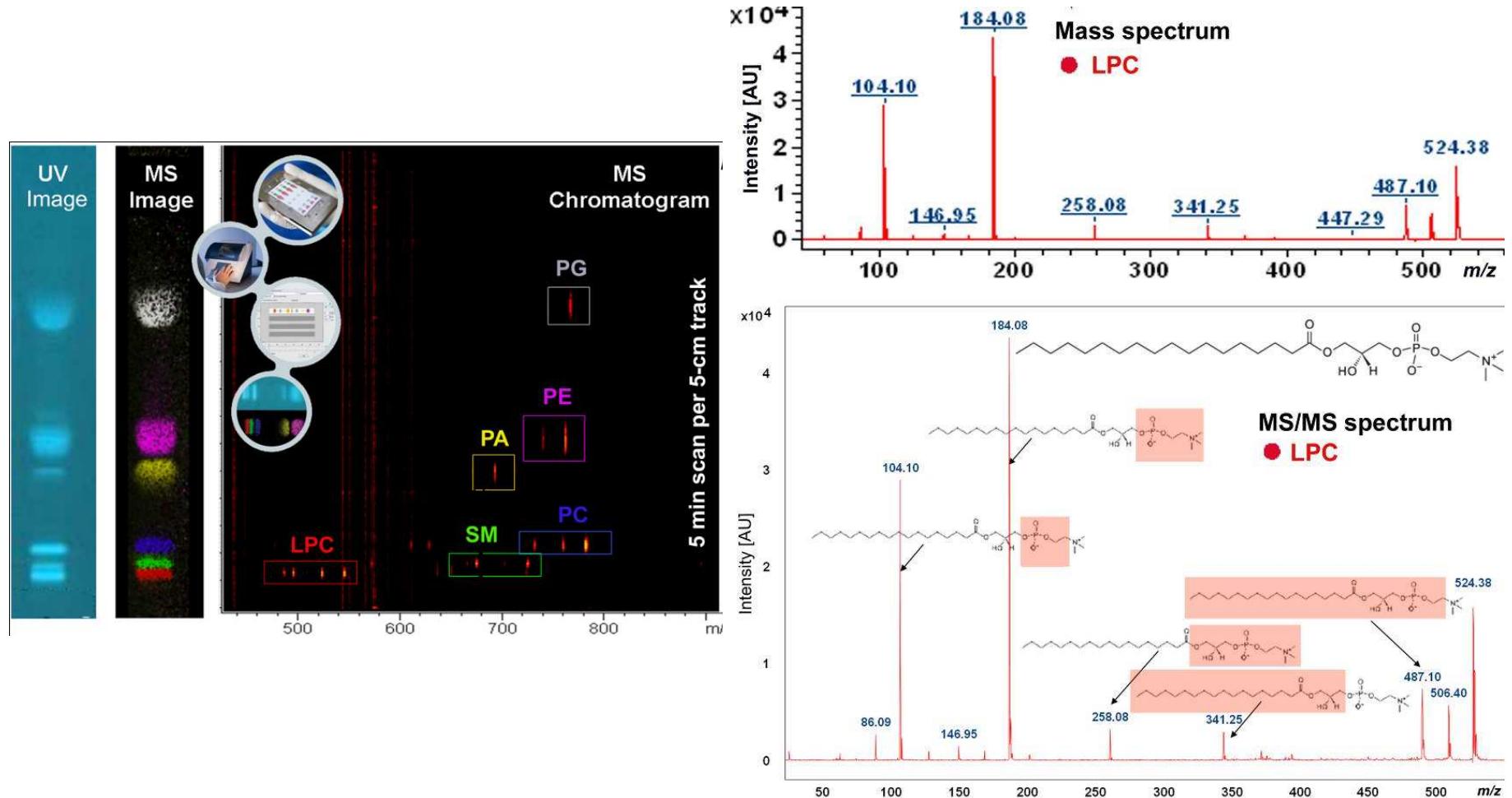


# HPTLC-FLD-MALDI-TOF MS

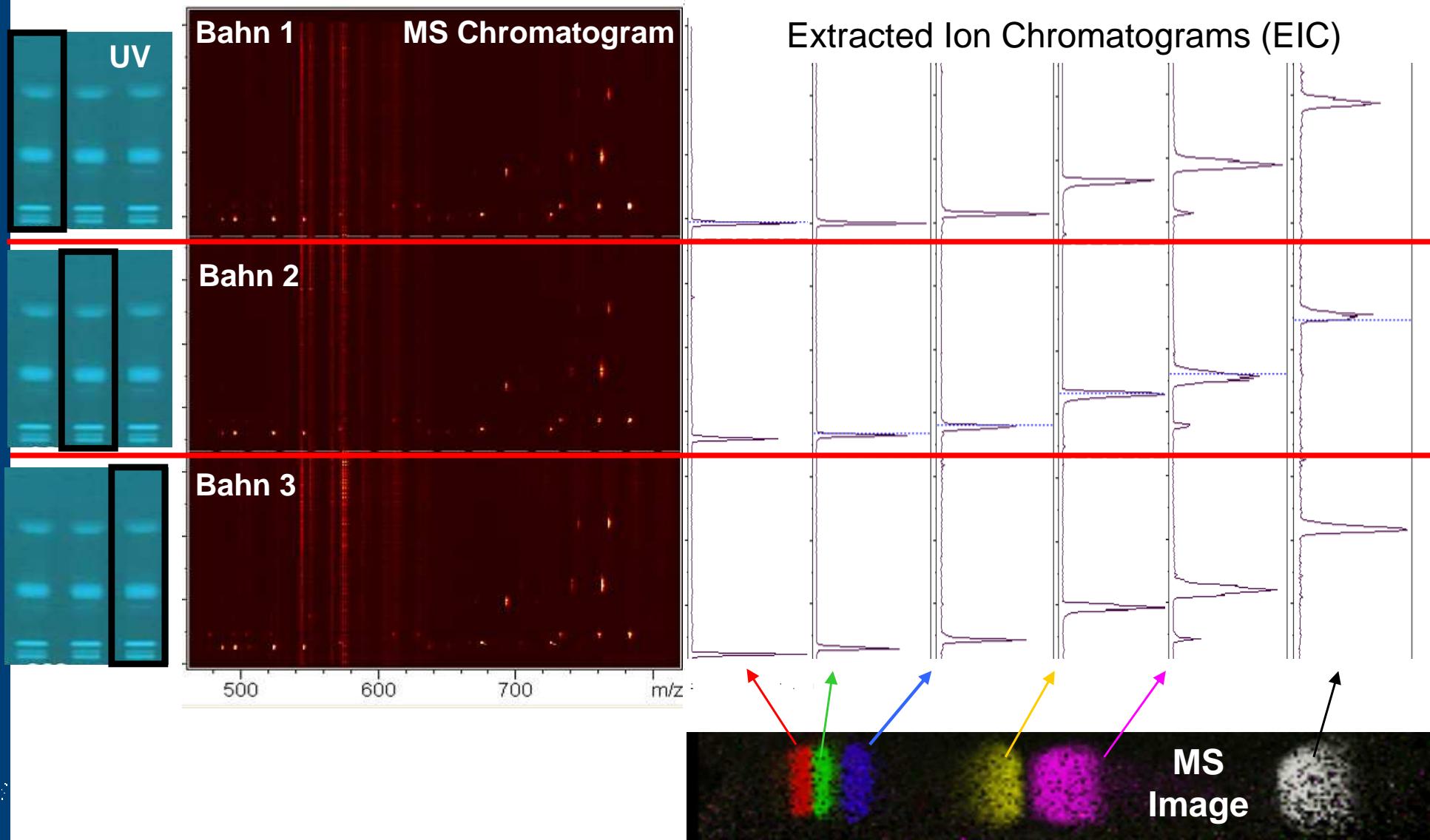


M. Schuerenberg *et al.*, IMSC 2009, Bremen, Poster PMM 386

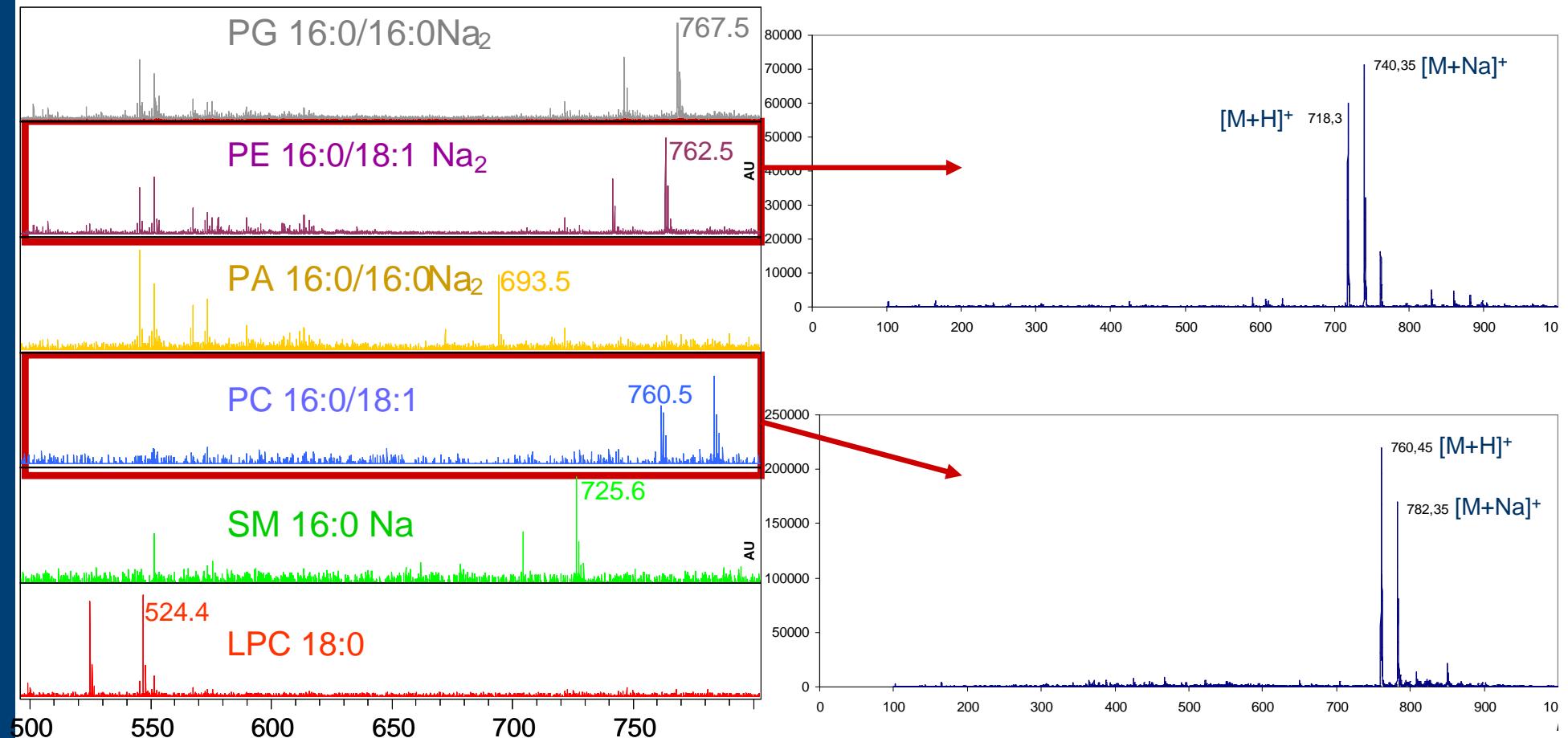
# HPTLC-FLD-MALDI-TOF MS



# Quantification?



# Comparison of mass spectra



# Comparison of the approaches

DART/APGD → dry desorption technique ↔ DESI



→ no plate preparation etc. ↔ SALDI, MALDI

→ ambient conditions, no high voltage ↔ micro junction

→ simple spectra ↔ MALDI, SIMS

→ quantitativ *with* internal standard → scan function

MALDI

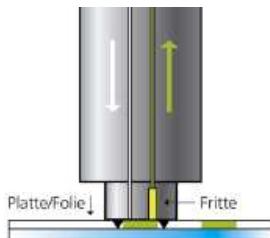


✓ strict protocol for plate preparation

✓ complex spectra

✓ quantitativ *with* internal standard → scan function

Elution-head  
based  
Interface



✓ *universally* connectable to any LC-MS system given

✓ plug & play interface (without adjustments or modifications)

✓ whole plate (no cut)

✓ all carriers on mostly all layers ↔ micro junction

✓ whole zone incl. depth profile → high detectabilities

✓ quantitativ *without* internal standard ↔ desorption techniques

✓ targeted recording → cost-effective, but *no* scan function

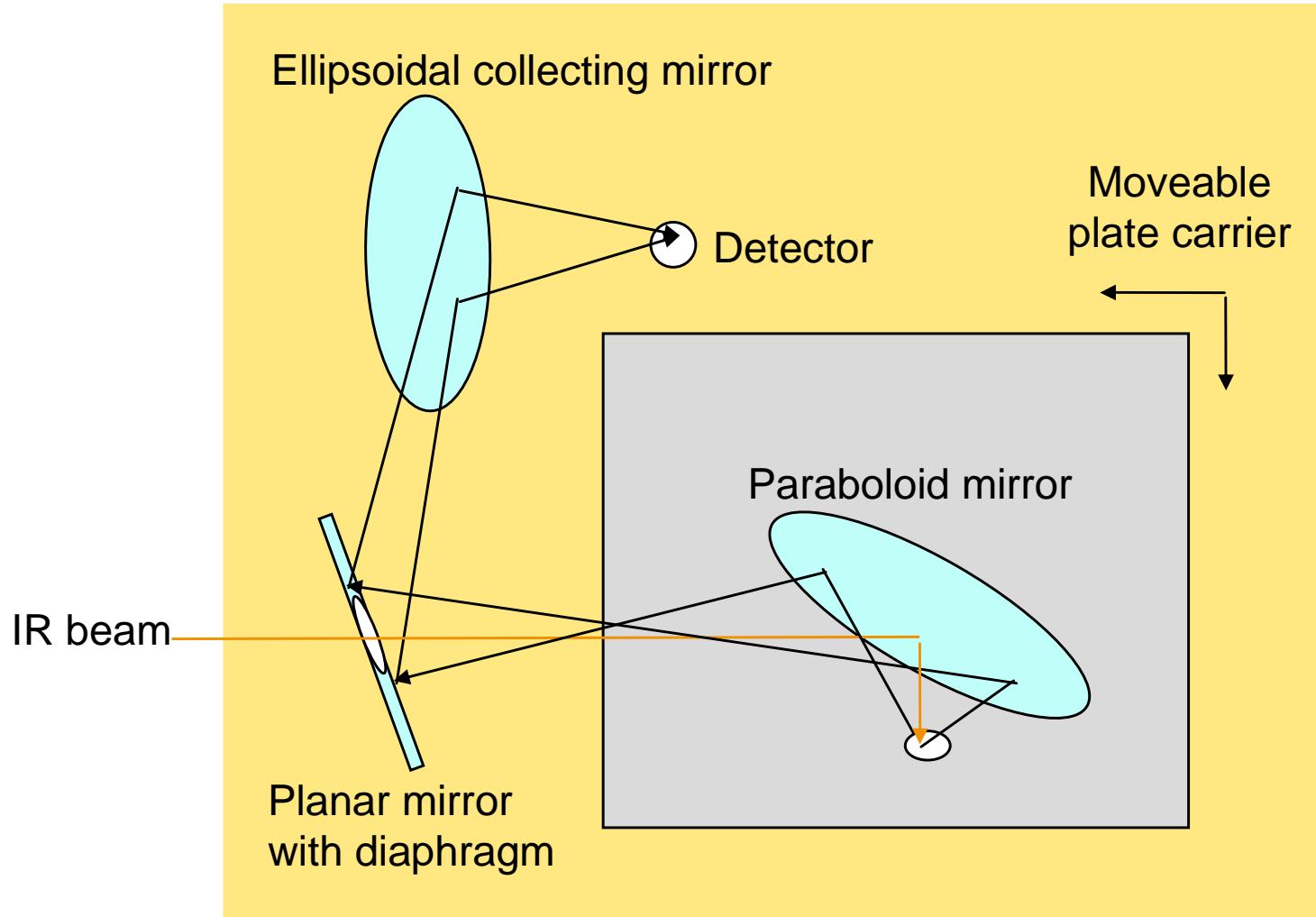
# Content

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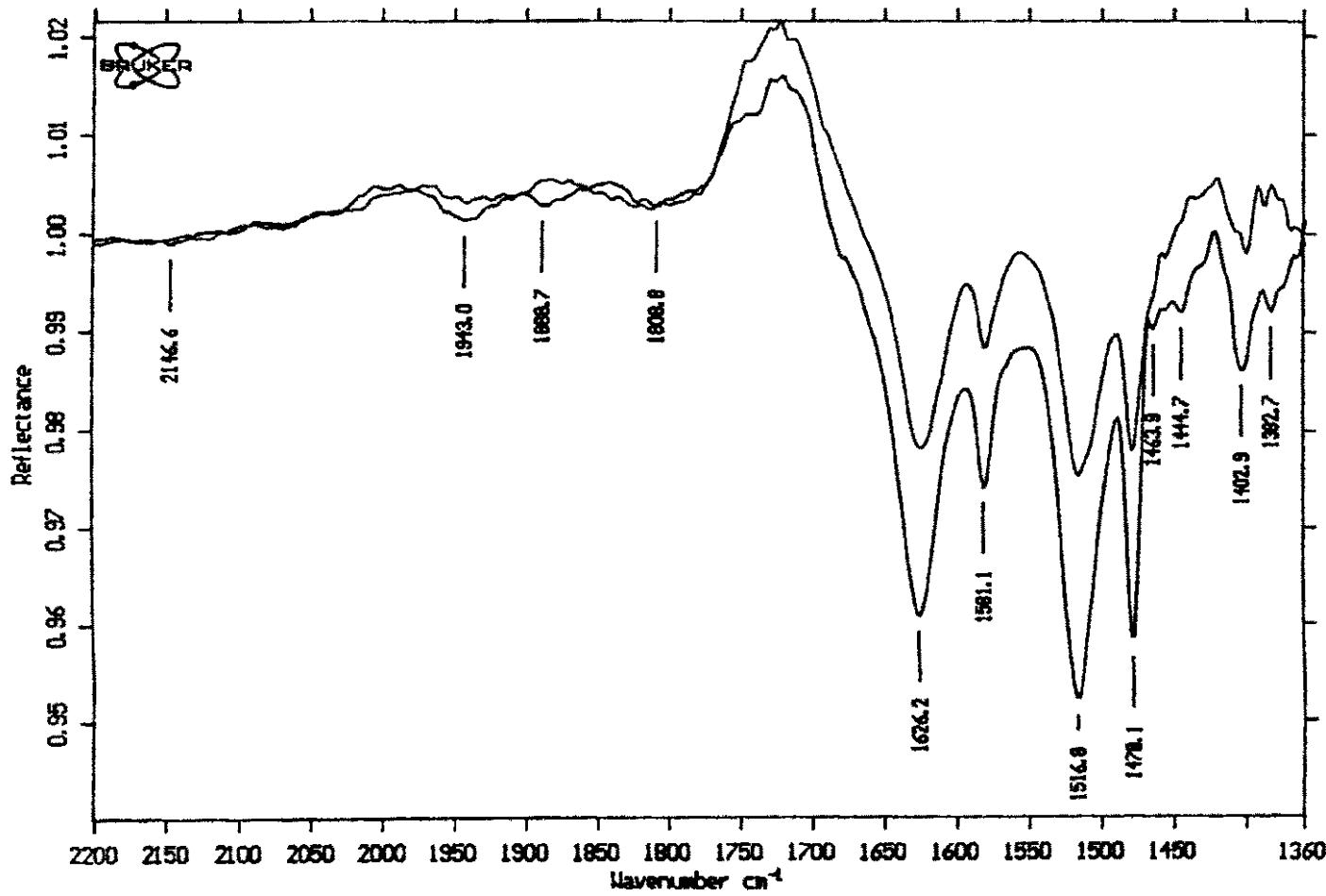
## Hyphenations with

1. UV/VIS/FLD/derivatizations
2. MS
3. FTIR
4. NMR
5. Bioassays

# HPTLC-DRIFT

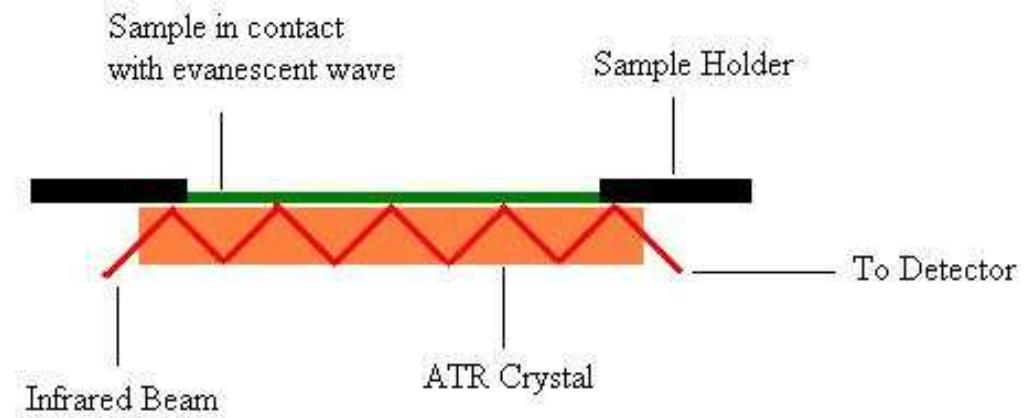
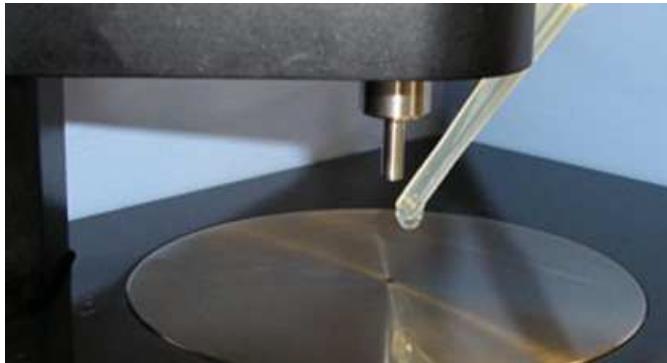


# FTIR spectrum of neburon in drinking water



## HPTLC-ATR FTIR

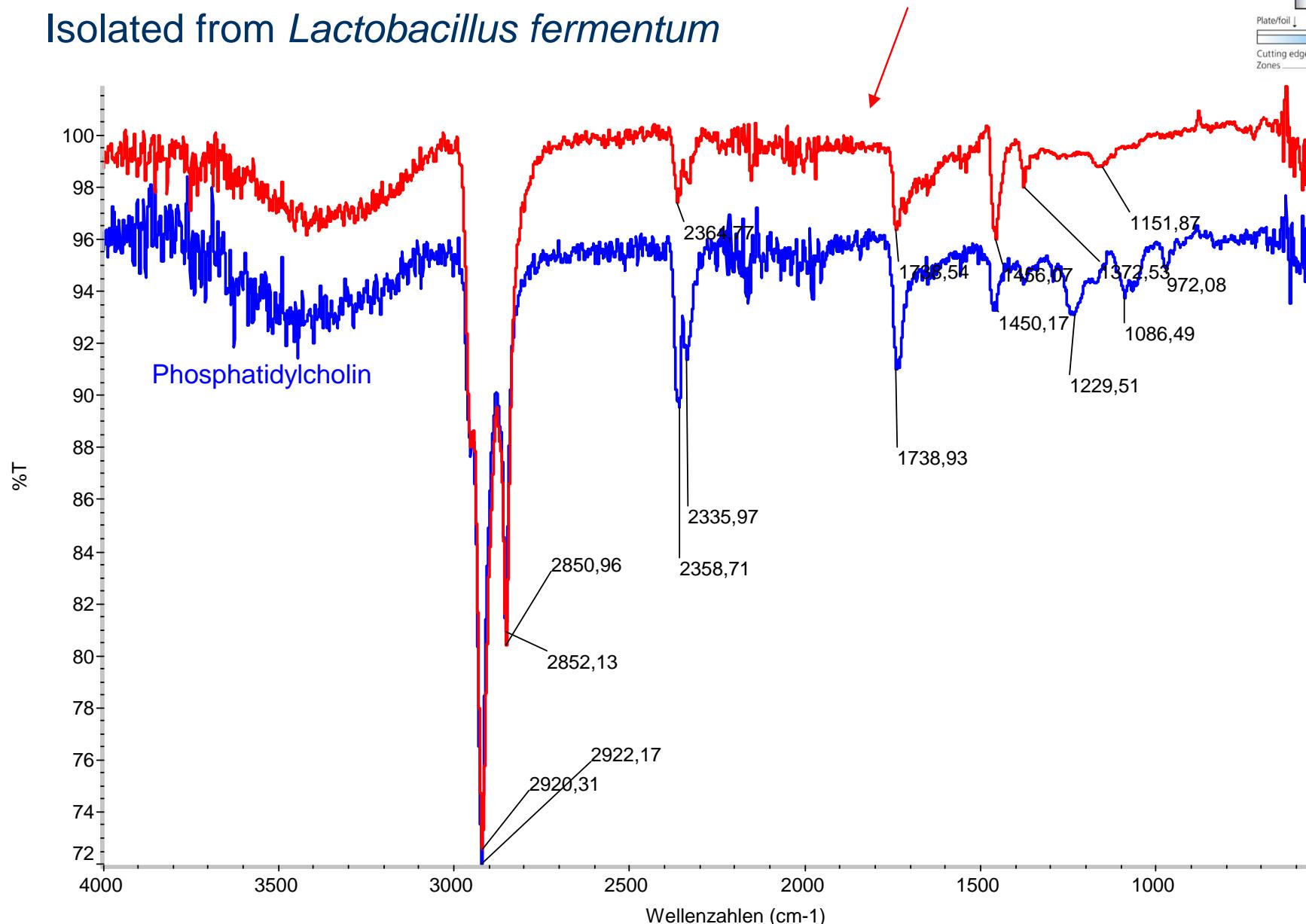
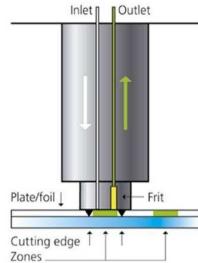
- attenuated total reflection infrared (ATR IR) spectroscopy
- samples examined directly (solid or liquid) without any preparation



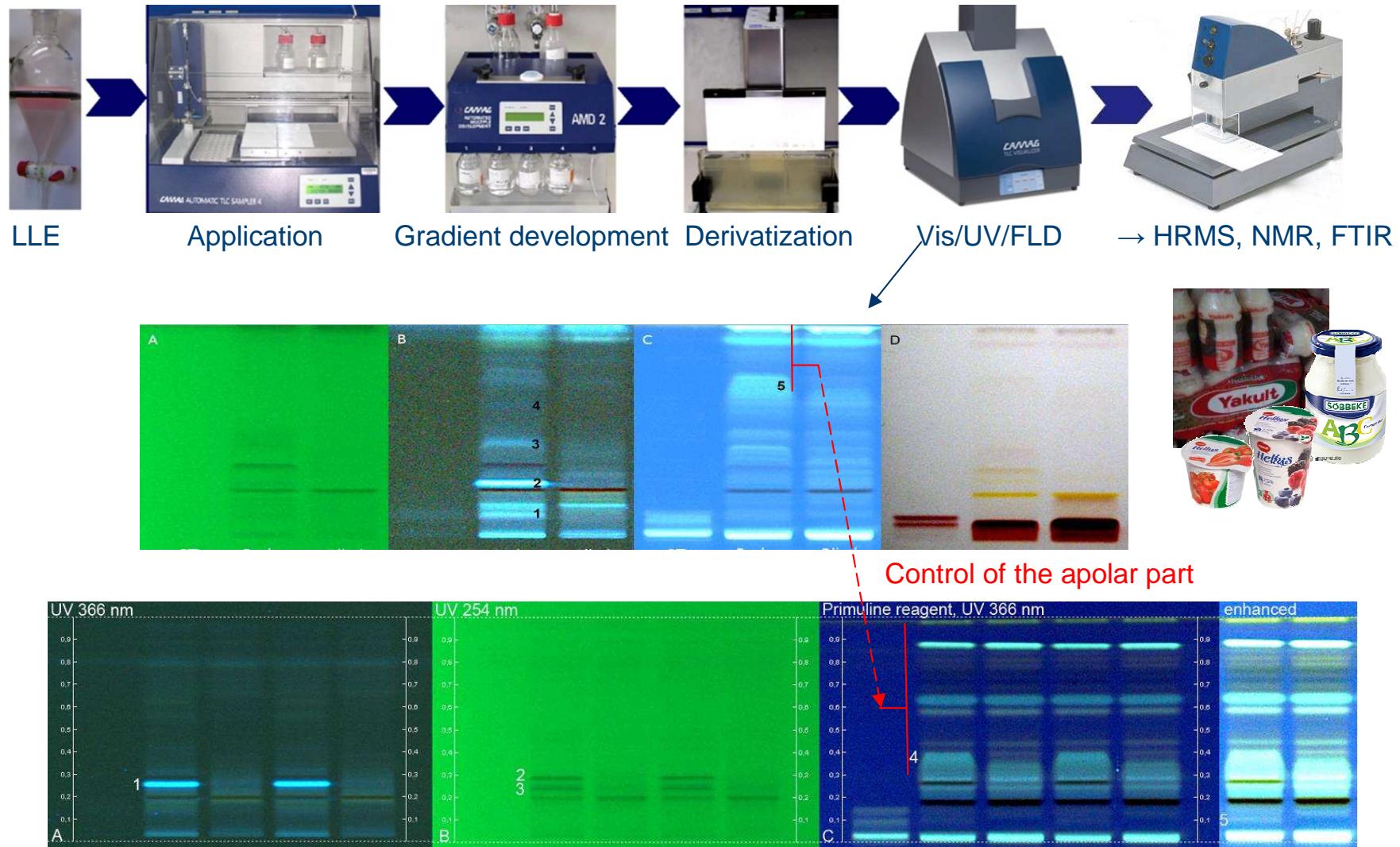
- zone eluted via TLC-MS Interface in 100 µL
- drop is directly applied and solvent rapidly evaporates
  - ☺ fast protocol
  - ☺ µg-amount per zone

# HPTLC-ATR FTIR of an anti-inflammatory compound

Isolated from *Lactobacillus fermentum*

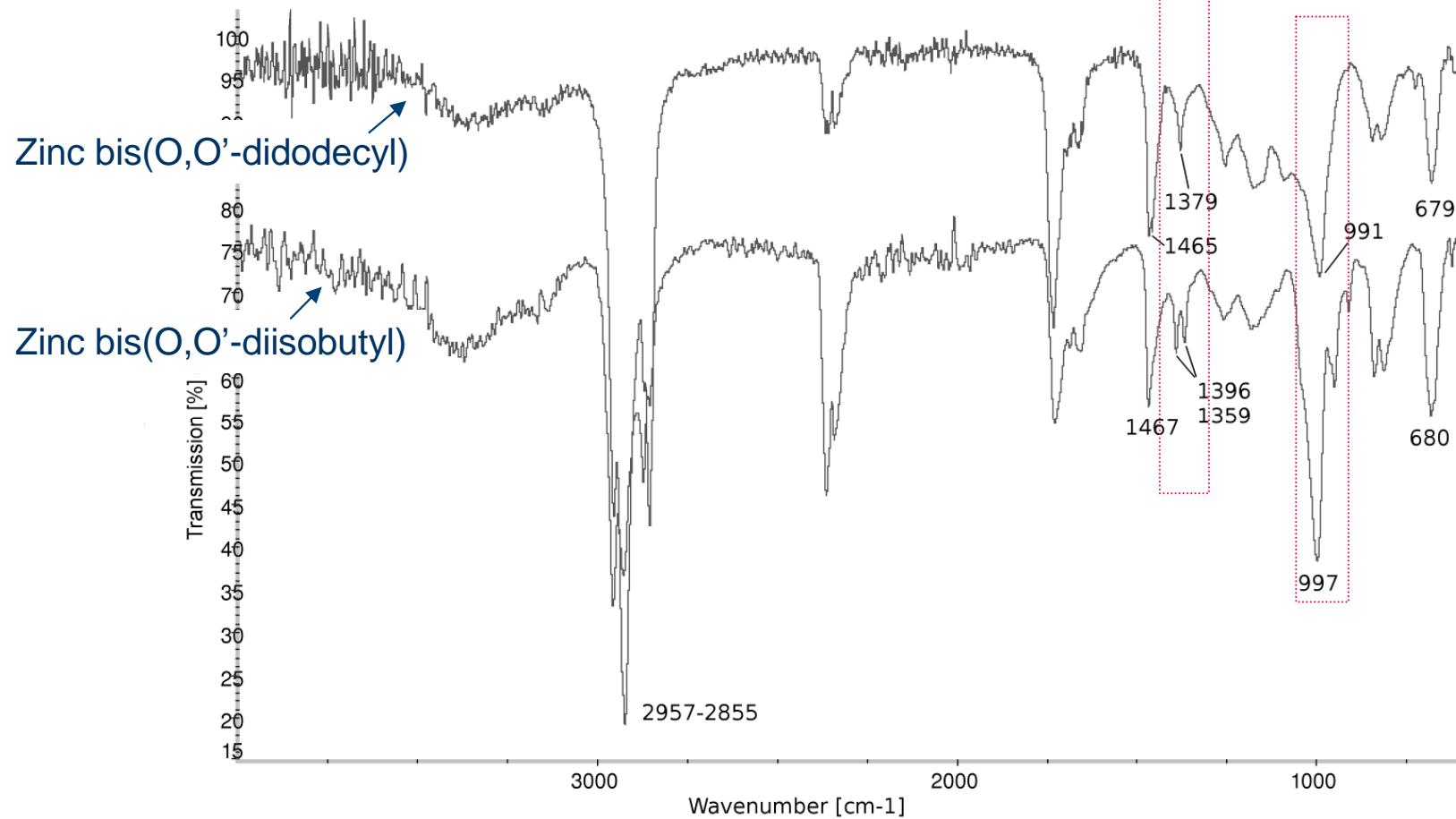
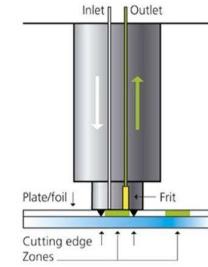


# Active compounds of *Lactobacillus fermentum*



# HPTLC/ATR-IR spectra

Dithiophosphate additives in mineral oil

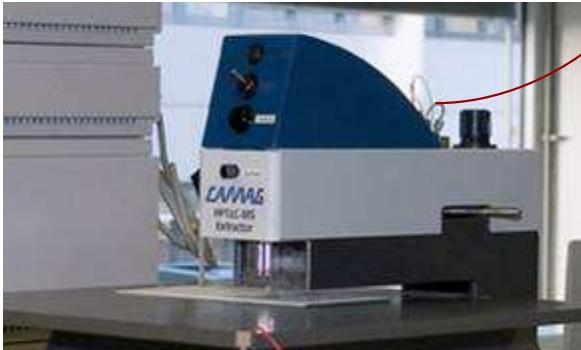


E. Dytkiewitz, G. Morlock, J AOAC Int 91 (2008) 1237-1244

# Hyphenation with NMR

- direct and online hyphenation
- limitations and possibilities
- coming soon

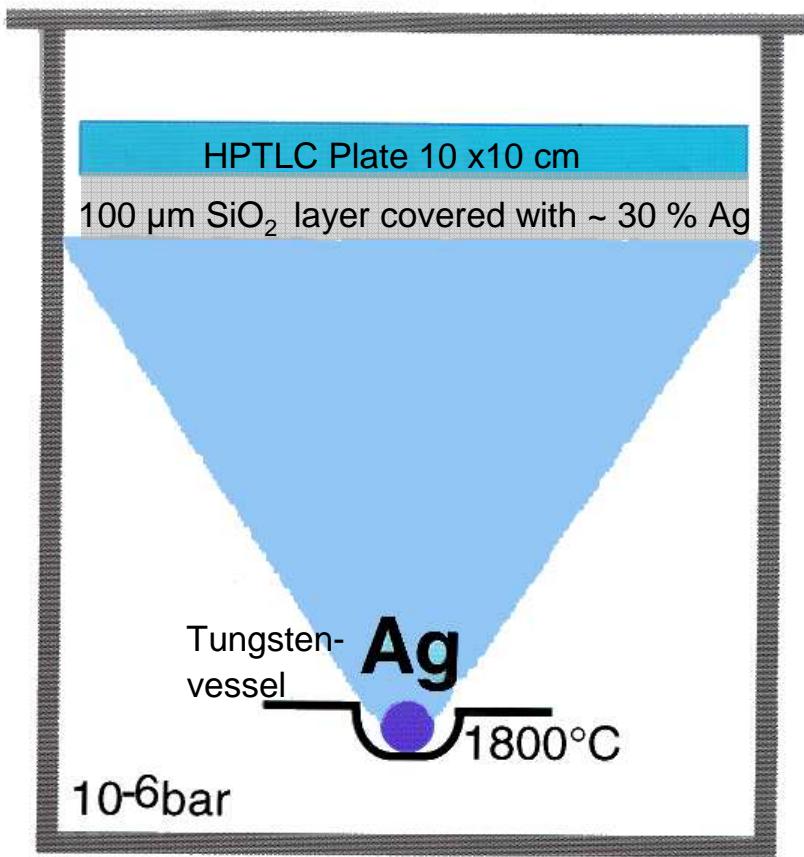
to NMR



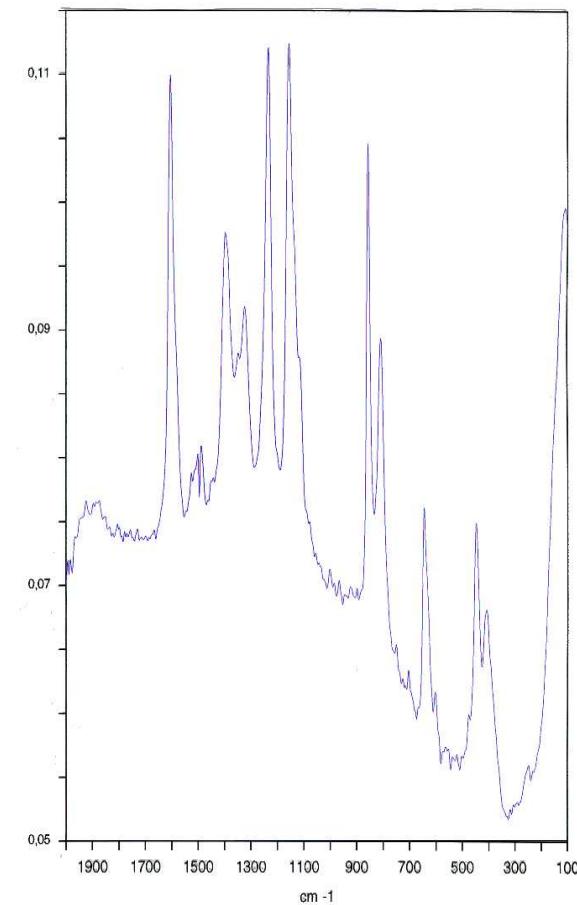
# Raman: FT-SERS

→ based on the work of Dr. Klaus Burger†; Bayer Laboratories, Germany

Vacuum transfer



10 ng/zone p-nitrophenol



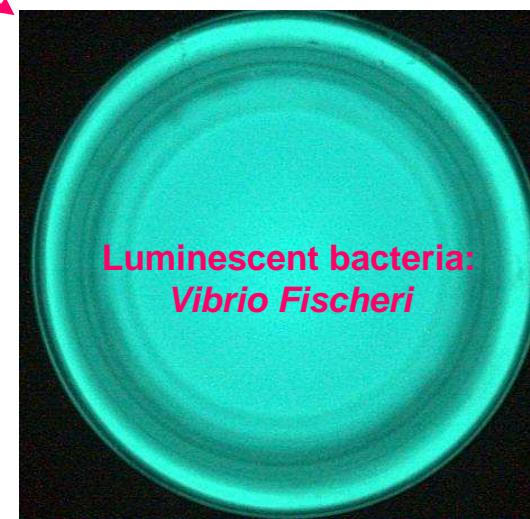
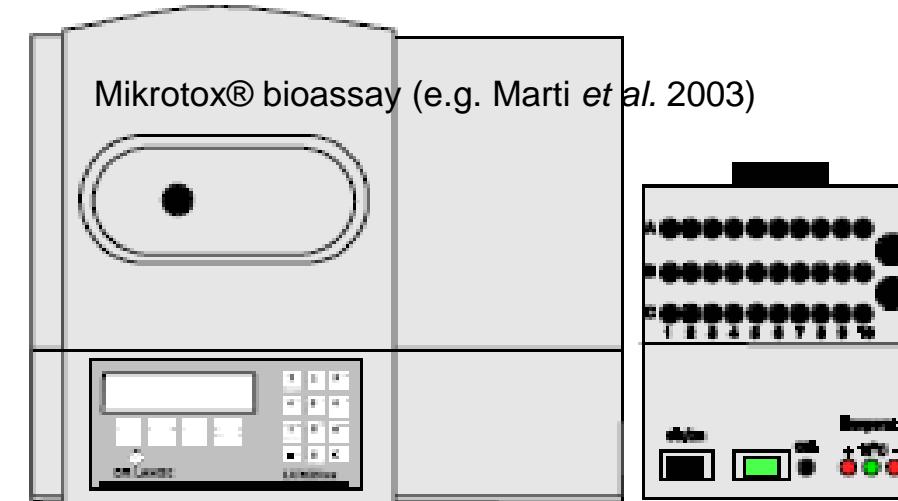
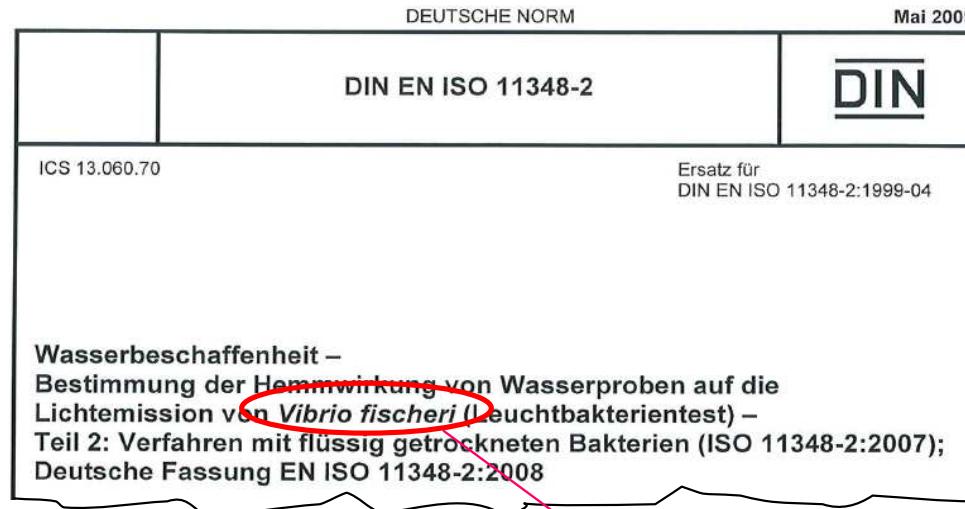
# Content

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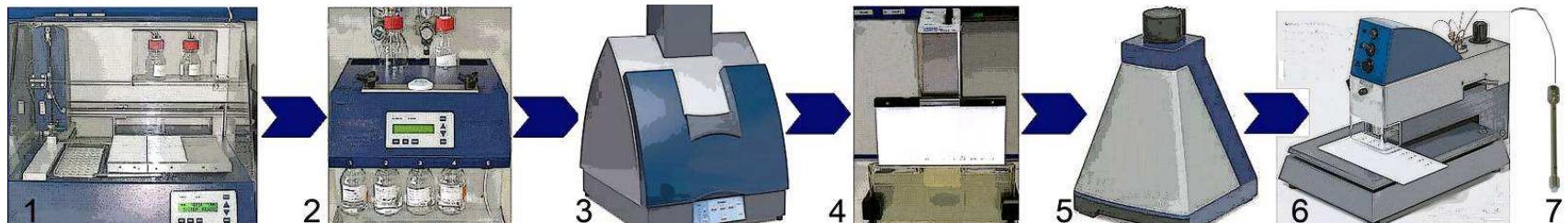
## Hyphenations with

1. UV/VIS/FLD/derivatizations
2. MS
3. FTIR
4. NMR
5. Bioassays

# Effect-directed analysis → sum parameter!

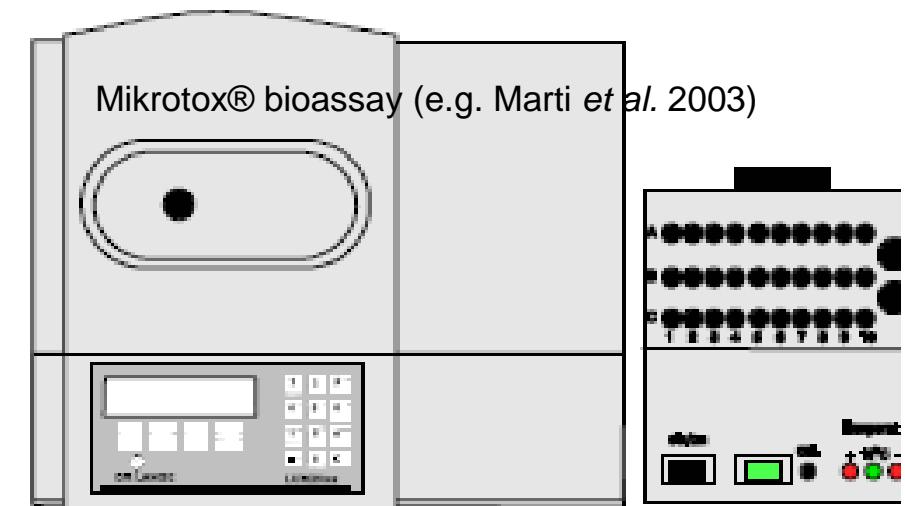


# Chromatography-bioassay versus cuvette test



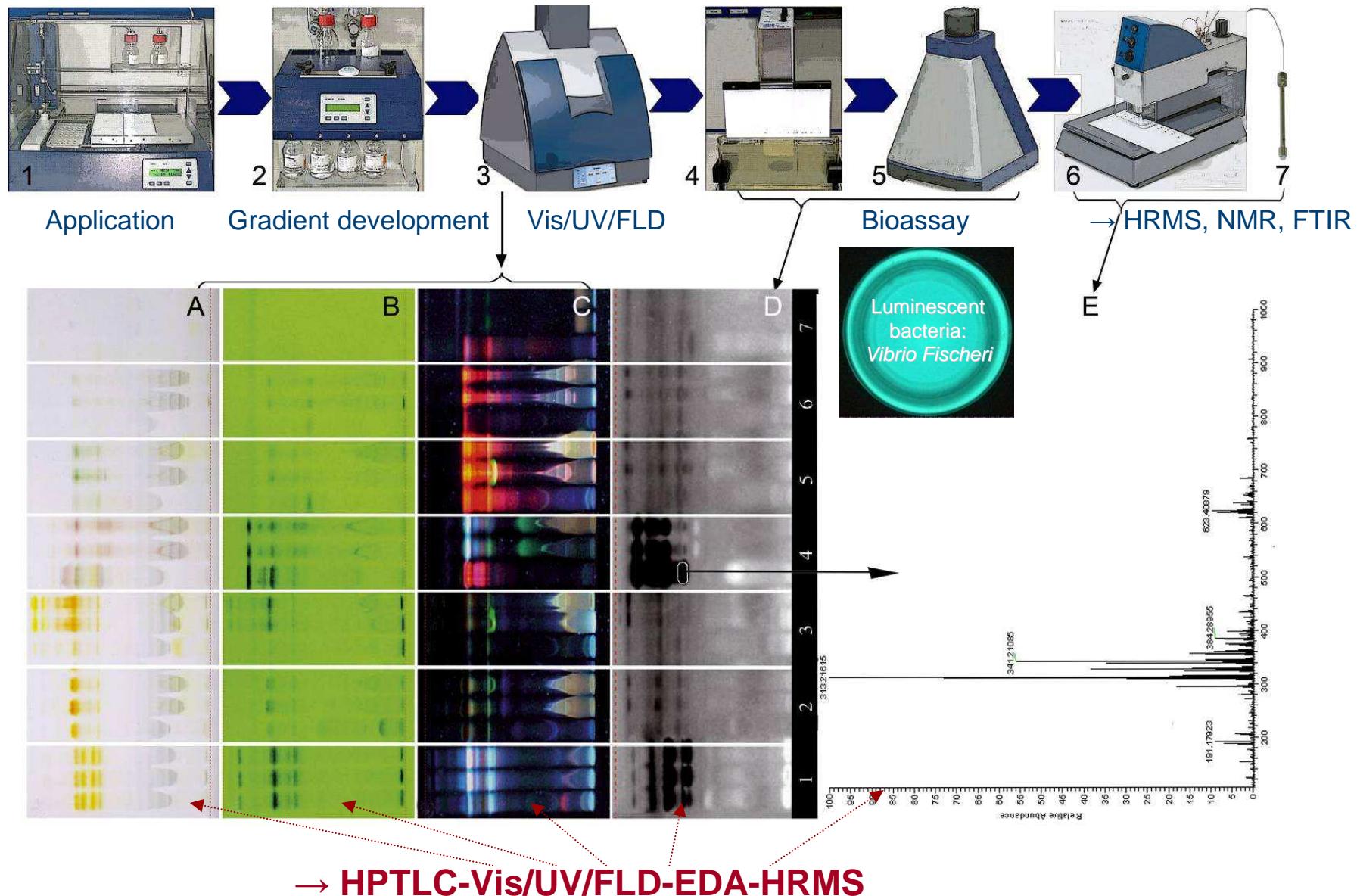
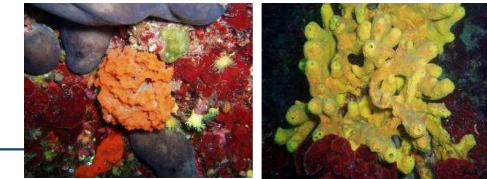
- Sample as original as possible
- Assignment of single compounds
- Bio-pattern is visible
- Polarity is clear

Some days versus some months



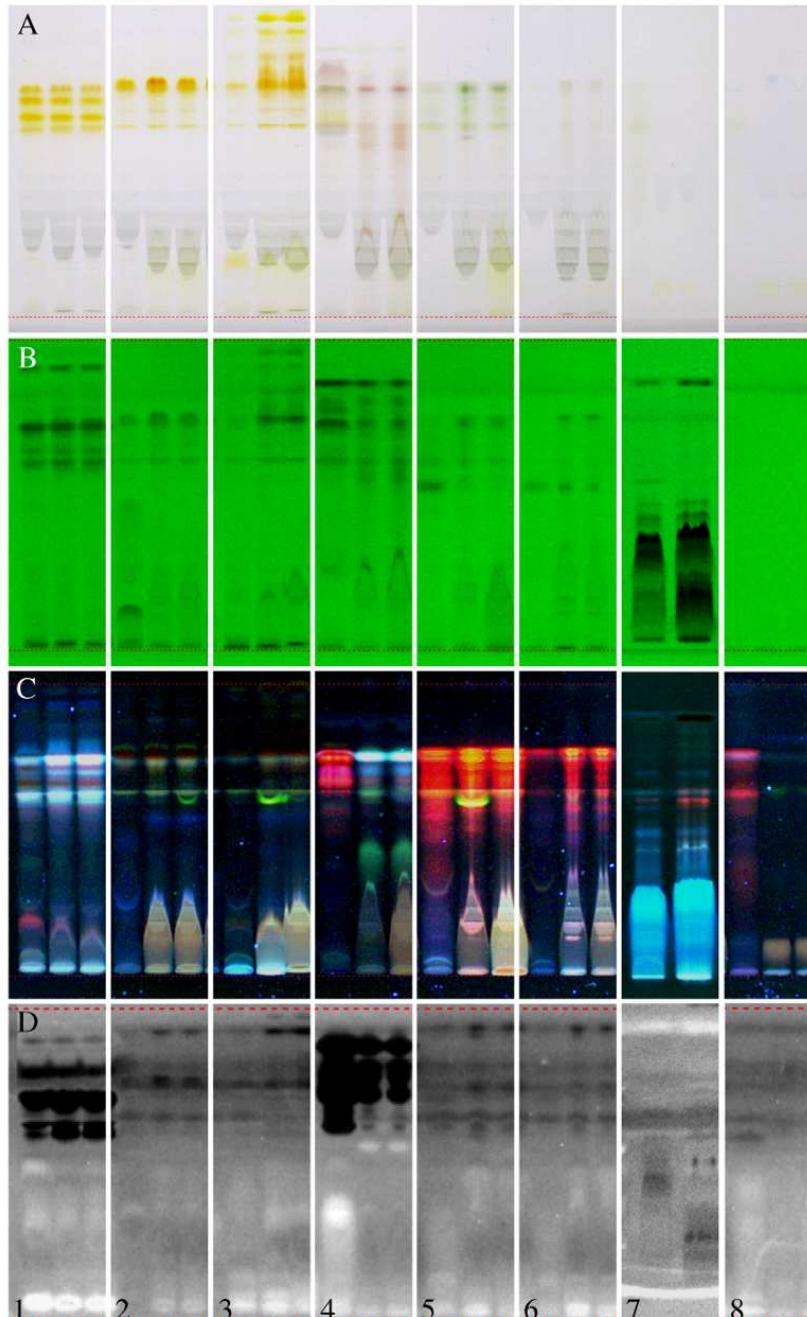
- Schlesinger et al., *Crambe crambe* marine sponge extract targets pancreatic and prostate cancer stem-like characteristics, in submission

# Bioactive products in marine sponges



# Primmorphe as bioreactor

Poster 3b



**1 *Acanthella acuta*, very toxic**

**2 *Axinella polypoides*, primmorphs more toxic than *in situ* sponge**

**3 *Suberites domuncula*, primmorphs more toxic than *in situ* sponge**

**4 *Dysidea avara*, bioactive metabolites besides Avarol and Avarone, loss of toxic metabolite, synthesis of enhancing metabolite**

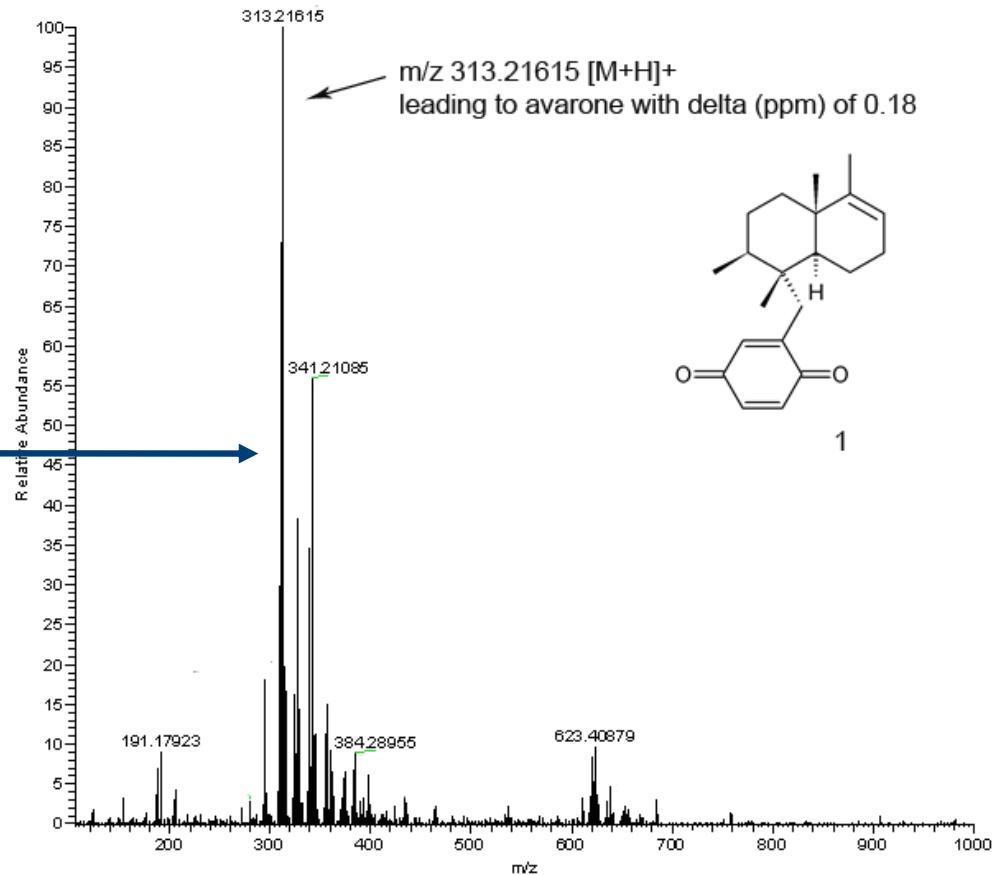
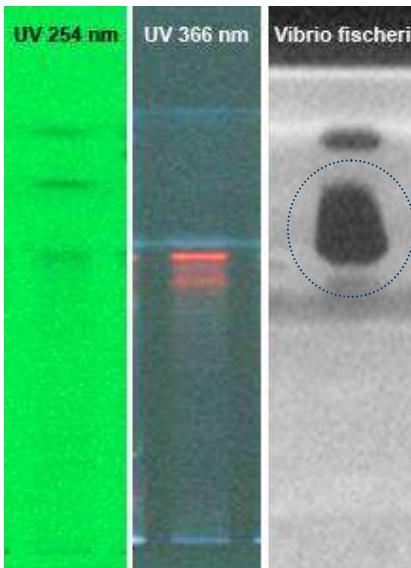
**5 *Petrosia ficiformis* with symbiotic cyanobacteria,**

**6 *Petrosia ficiformis* without symbiotic cyanobacteria, different pattern but equal toxicity**

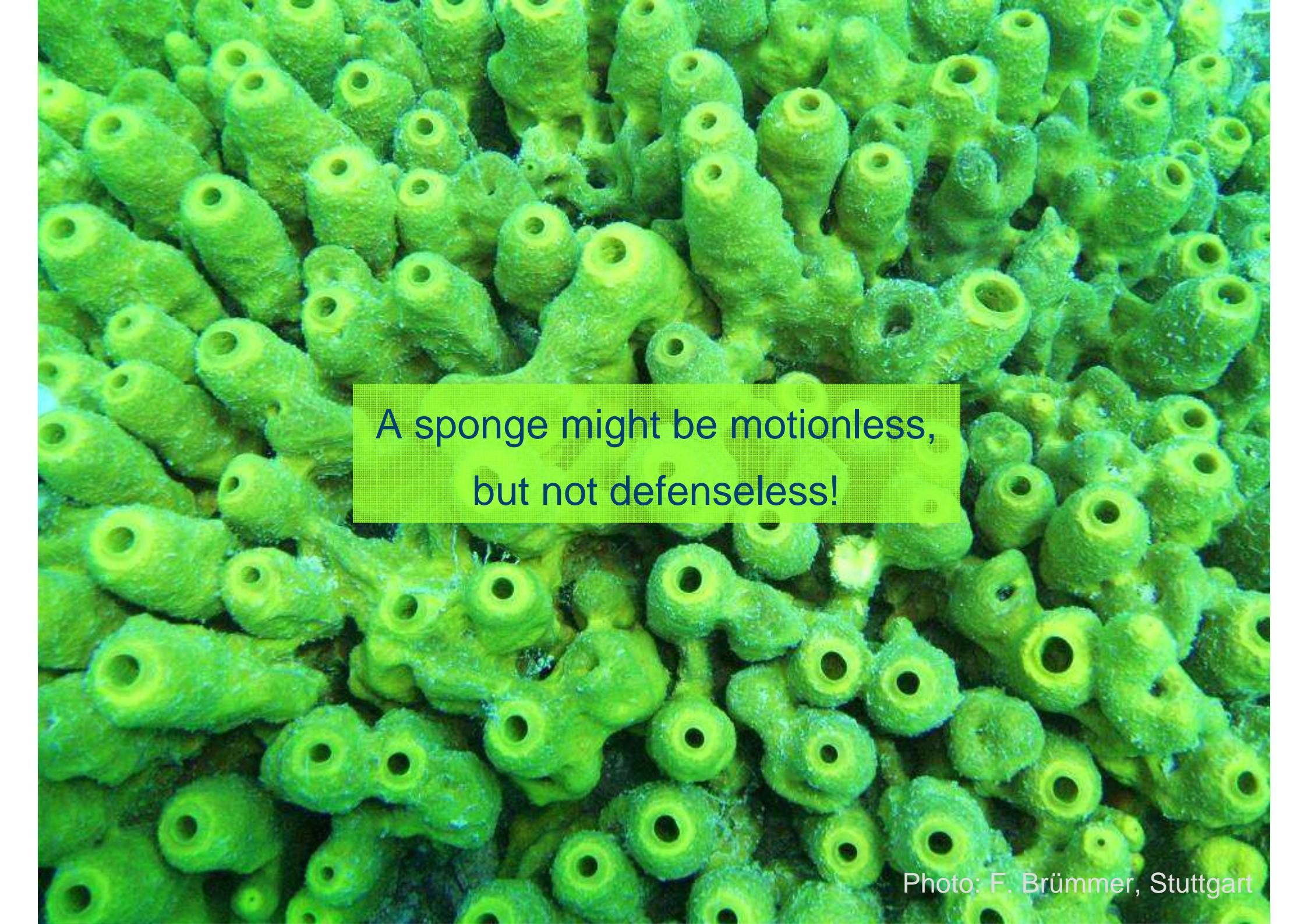
**7 *Axinella damicornis*, synthesis of a new toxic metabolite in primmorphs**

**8 *Ephydatia fluviatilis*, first documentation of bioactivity, loss in primmorphs**

# HPTLC-Bioactivity-HRMS



G. Morlock, W. Schwack, *LCGC Eur* 21 (2008) 366-371

A close-up photograph of a dense cluster of green sponges. The sponges have a textured, porous surface with numerous small, circular pores or oscula. They are packed closely together, creating a complex, organic pattern.

A sponge might be motionless,  
but not defenseless!

Photo: F. Brümmer, Stuttgart

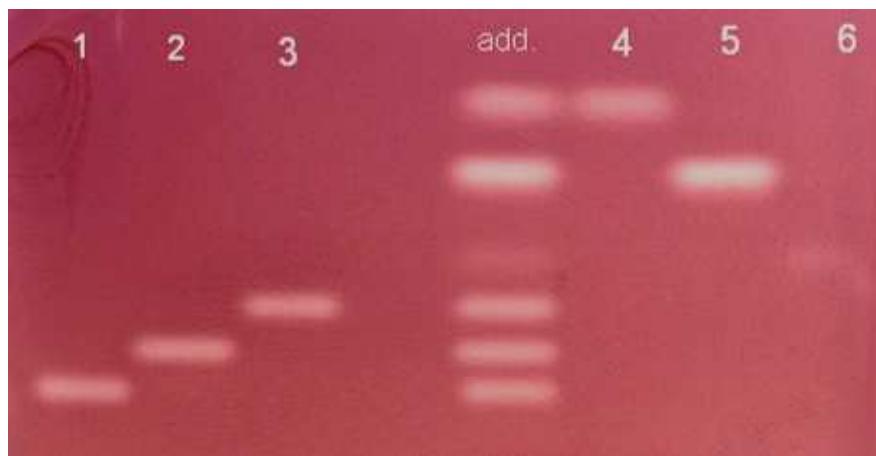
# Screening of natural products by HPTLC-Bioactivity-HRMS

- ✓ Combination of different methods (SPE, GPC, prep. HPLC) for fractionation can be skipped → HPTLC is highly matrix-tolerant
- ✓ Isolation and purification of substances, always followed by bioactivity testing, can be skipped
- ✓ 30 sponge extracts separated in parallel under identical chromatographic and environmental conditions
- ✓ Directly extracted/desorbed from the HPTLC plate and transfer into the MS within seconds or half a minute
- ✓ Highly targeted coupling with HRMS → after evaluation just from zones of interest → very cost-effective
- ✓ Detectability of the extraction interface comparable to HPLC → the whole zone inclusive its depth profile is extracted
- ✓ Bioassays not interfered by solvents → evaporated after chromatography → no inactivation

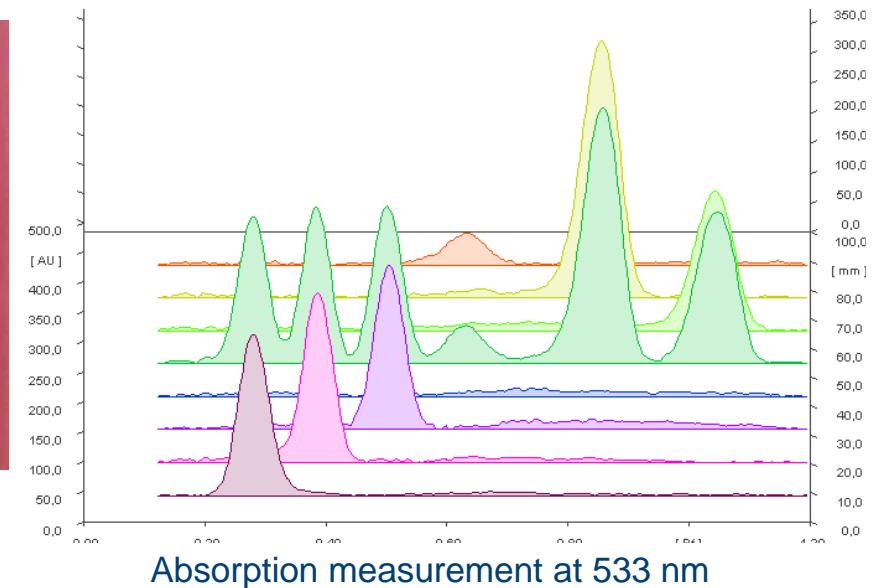
# HPTLC-VIS/UV/FLD-EDA

Cholinesterase inhibiting pesticides by esterases

- detectability down to 2 pg/zone
- using an esterase and substrate (1-naphthylacetate/fast blue salt B) solution
- white zones on a pink background



1. Paraoxon-methyl, 2. malaoxon, 3. paraoxon,  
4. ethiofencarb, 5. chlorfenvinfos, 6. dichlorvos



R. Akkad, W. Schwack, J Planar Chromatogr 21 (2008) 411-415

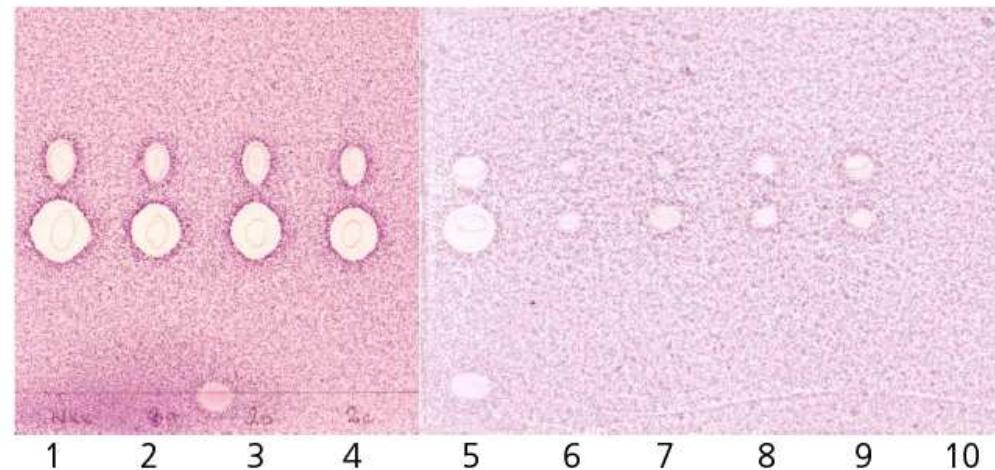
# Detection of antibiotics with *Bacillus subtilis*

## Determination of enrofloxacin and ciprofloxacin in milk by direct bioautography detection



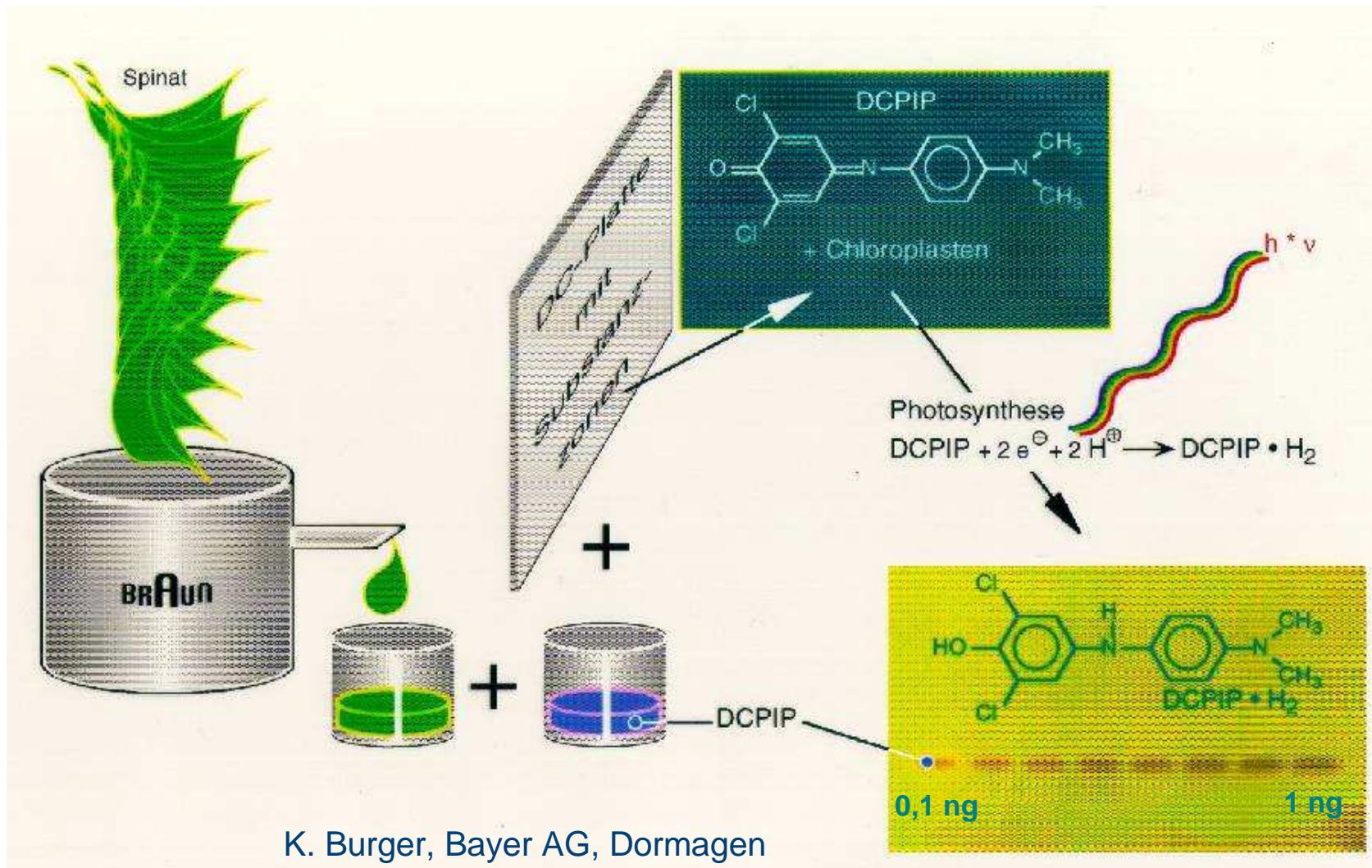
From left: M.Sc. Wioleta Bąk, Dr. hab. Irena Choma, M.Sc. Edyta Grzelak, Dr. Karol Pilorz and Dr. hab. Barbara Majer-Dziedzic.

Lecture 10a



# Detection with chloroplasts (spinach)

→ Photosynthesis inhibiting herbicides ( $\rightarrow 100 \text{ pg/zone}$ )



K. Burger, Bayer AG, Dormagen

# Content

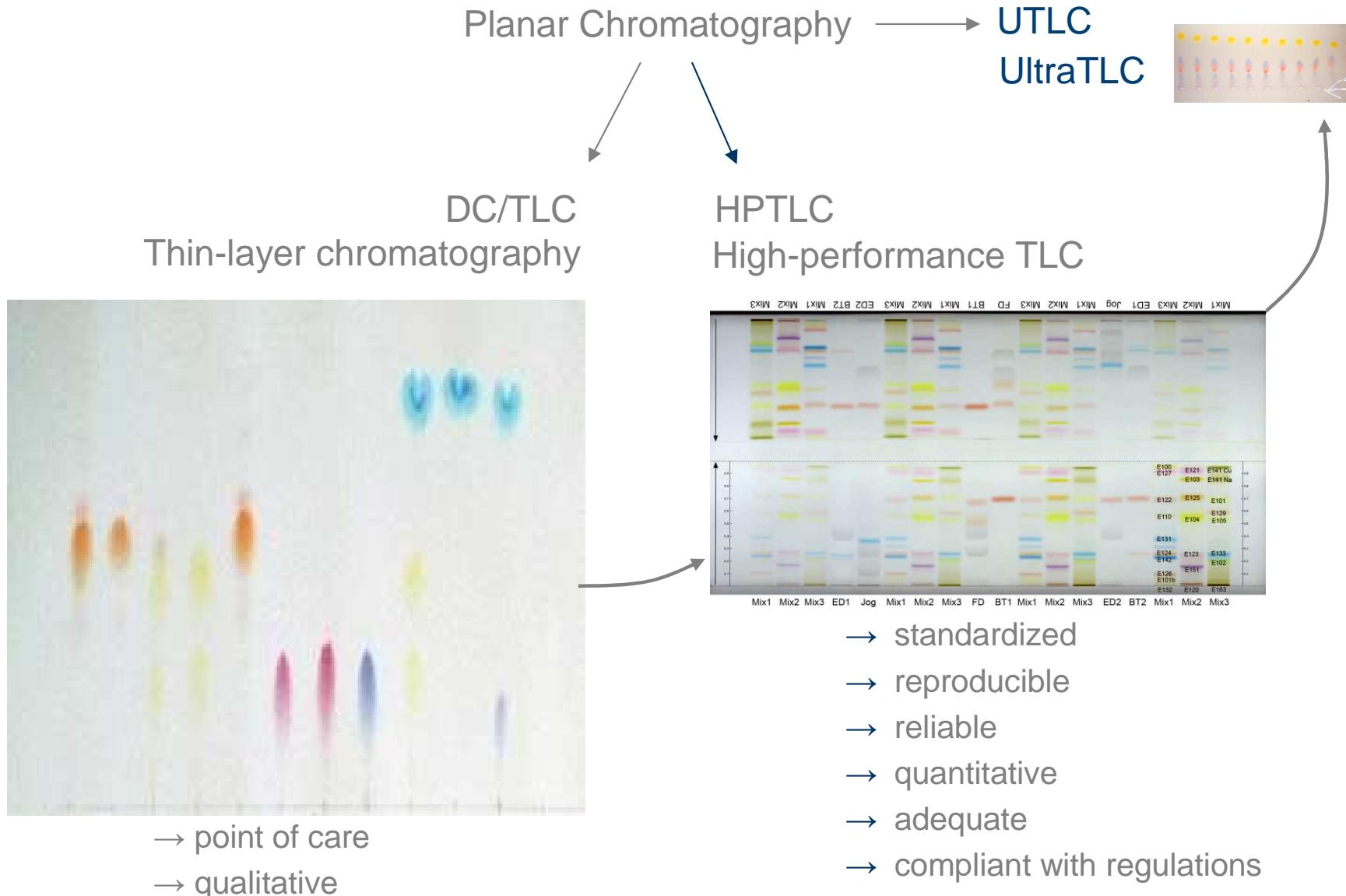
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## Hyphenations with

1. UV/VIS/FLD/derivatizations
2. MS
3. FTIR
4. NMR
5. Bioassays

*Session 10*

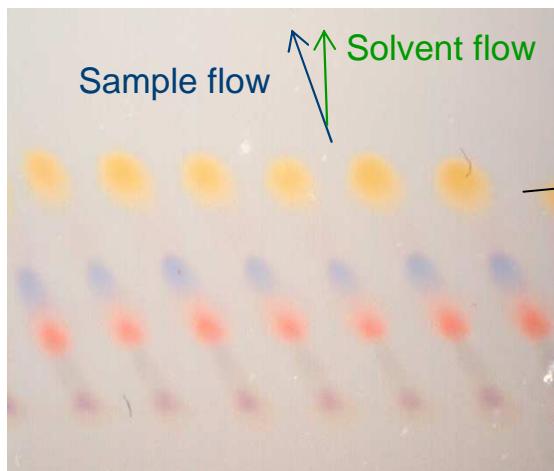
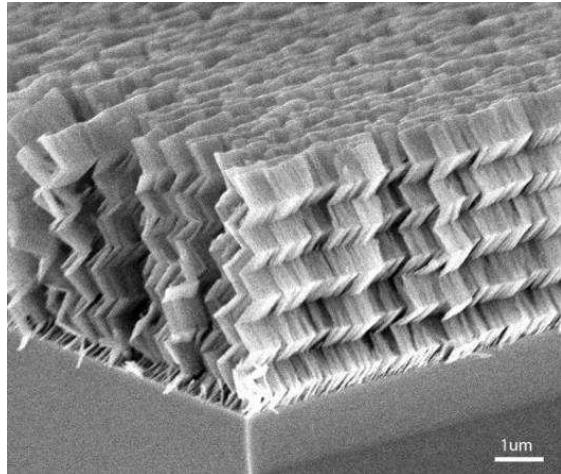
# High-Performance Thin-Layer Chromatography



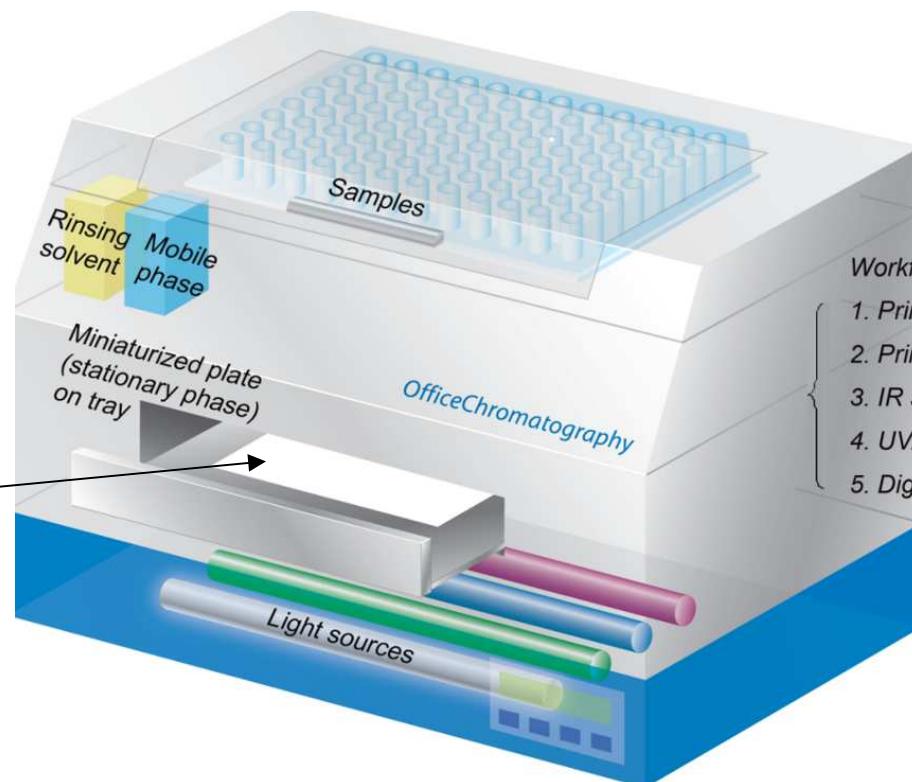
# Office Chromatography

Lecture 2b

Nanostructured UTLC plates

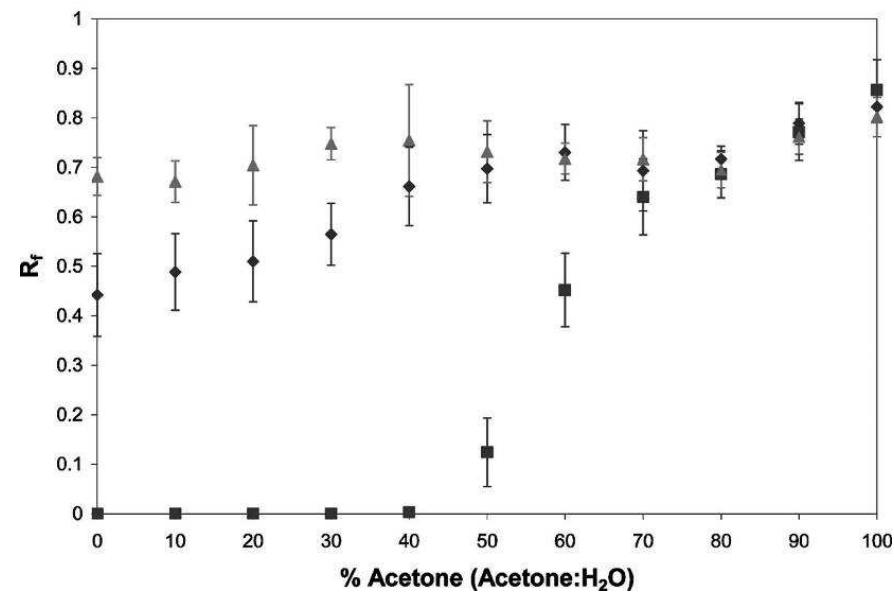
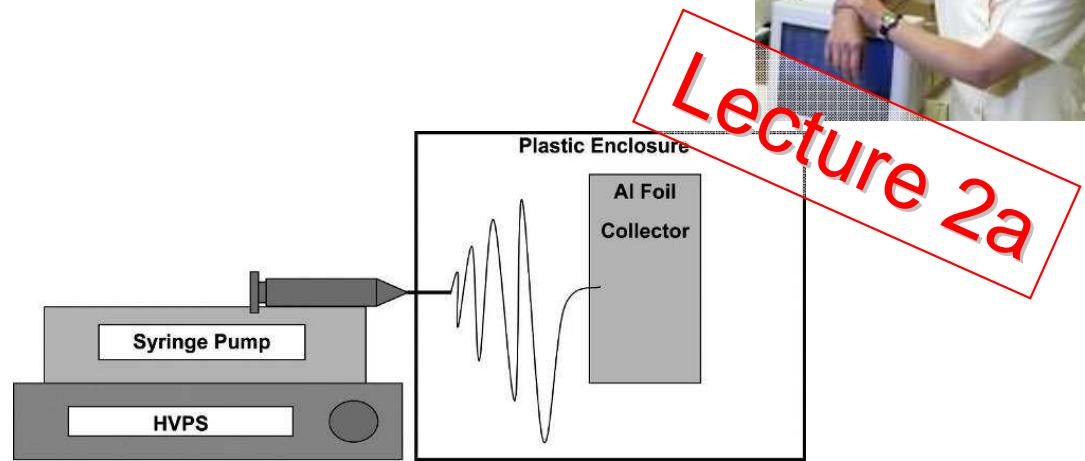
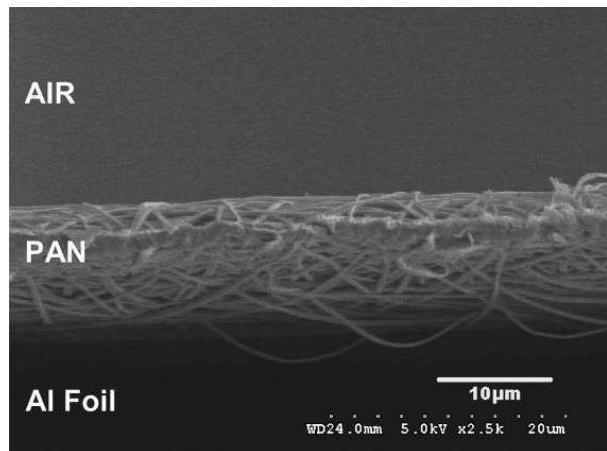
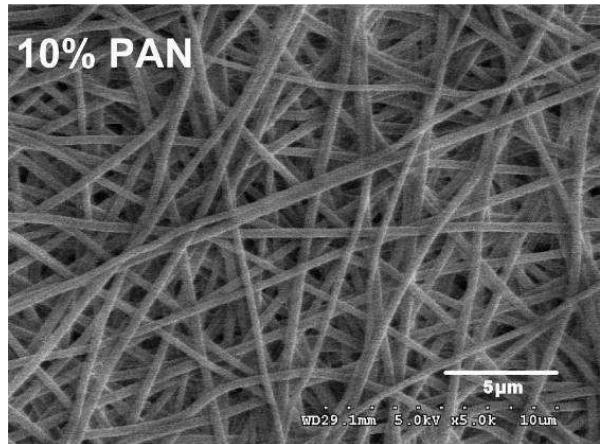


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G. Morlock, C. Oellig, L. Bezuidenhout, M. Brett & W. Schwack,  
Anal. Chem. 82 (2010) 2940-2946

# Electrospun UTLC



Jonathan E. Clark, Susan V. Olesik, Anal Chem 81 (2009) 4121–4129  
and J Chromatogr A 1217 (2010) 4655–4662



# Nanotube UTLC

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[www.MaterialsViews.com](http://www.MaterialsViews.com)

Lecture 2c

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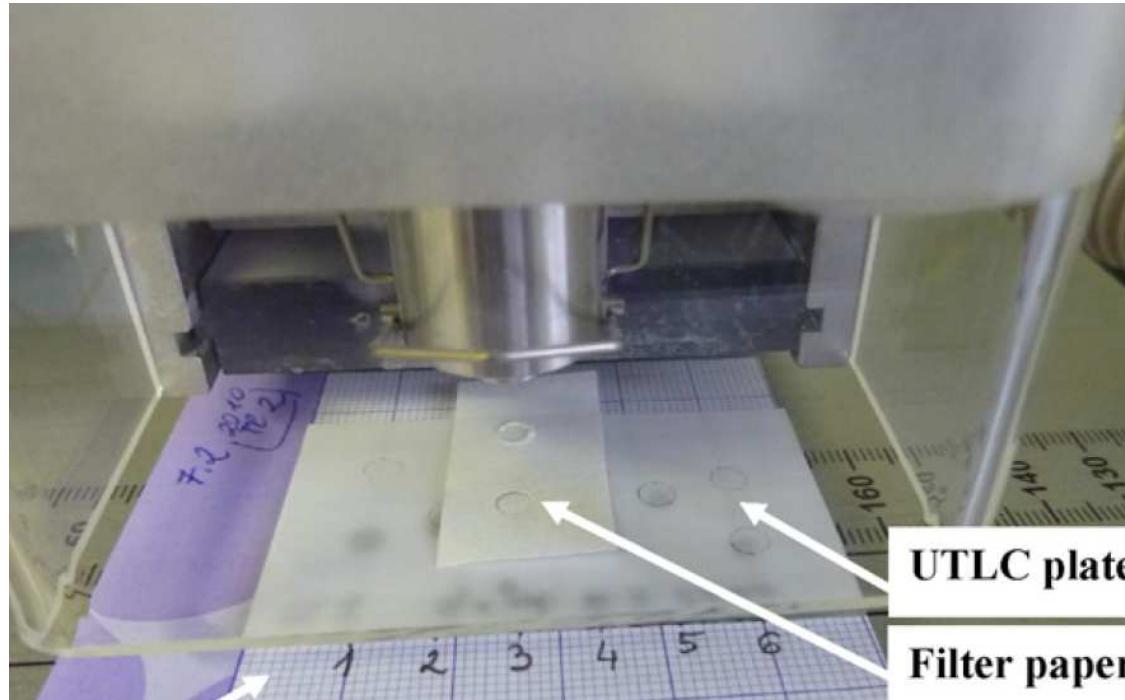
## Carbon-Nanotube-Templated Microfabrication of Porous Silicon-Carbon Materials with Application to Chemical Separations

*Jun Song, David S. Jensen, David N. Hutchison, Brendan Turner, Taylor Wood, Andrew Dadson, Michael A. Vail, Matthew R. Linford, Richard R. Vanfleet, and Robert C. Davis\**

Carbon-nanotube-templated microfabrication (CNT-M) of porous materials is demonstrated. Partial chemical infiltration of 3D carbon-nanotube structures with silicon results in a mechanically robust material, structured from the 10 nm scale to the 100 µm scale. The nanoscale dimensions are determined by the diameter and spacing of the resulting silicon/carbon nanotubes, while the microscale dimensions are controlled by the lithographic patterning

recombination rates, and mobilities are strongly influenced by nanoscale structuring.<sup>[4]</sup> Often, multiple physical properties are coupled and are jointly influenced by nanoscale structuring, as is the case for strained silicon: nanoscale strain control is used to produce higher mobilities than achievable in the bulk.<sup>[5]</sup> Coupling

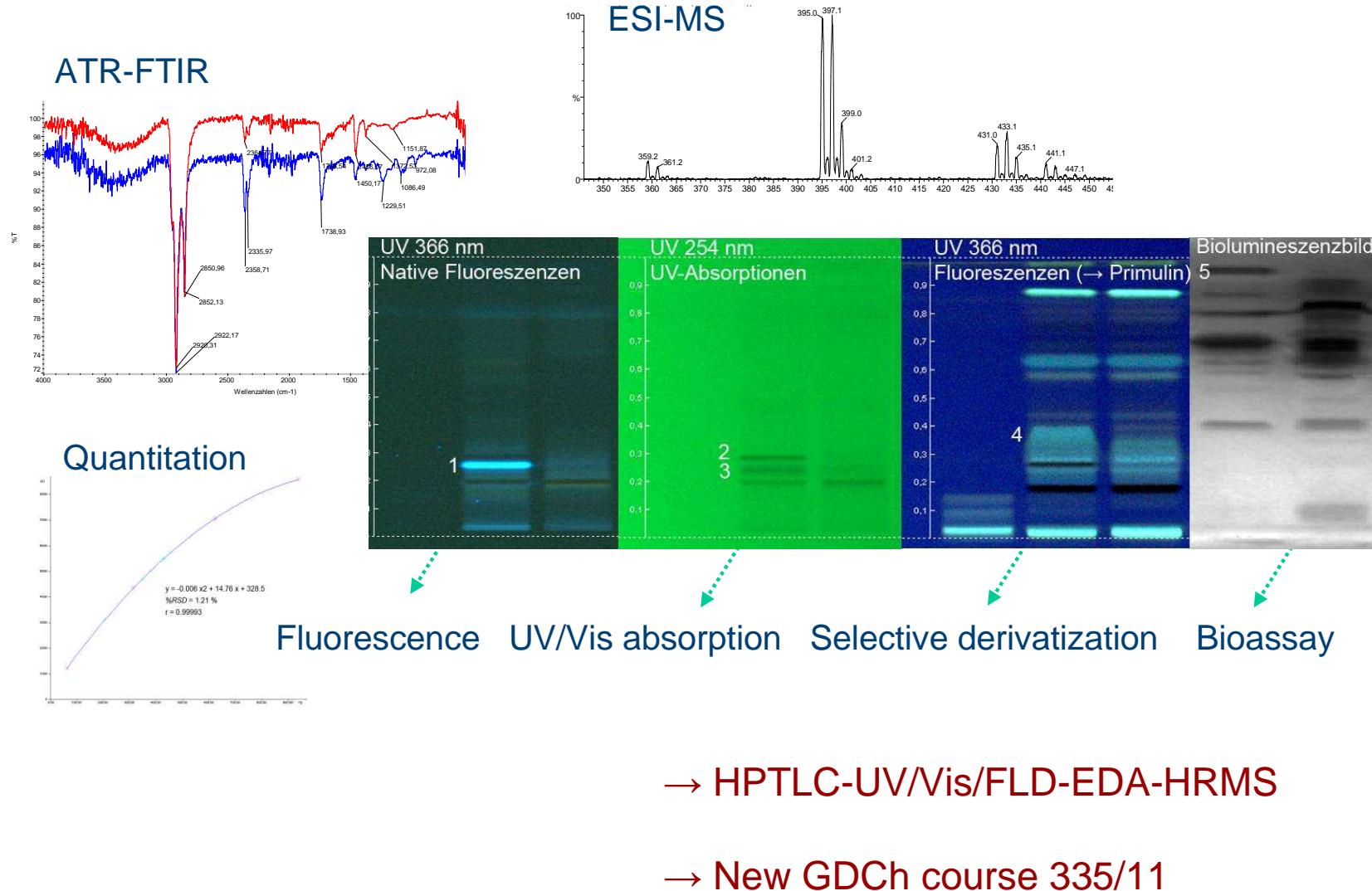
# UTLC-MS



Lecture 3a

I. Vovk et al. J Chromatogr A 1218 (2011) 3089–3094

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**Carbohydrates**

G. LODI\*, C. BIGHI, V. BRANDOLINI, E. MENZIANI, B. TOSI, (\*Dipartimento di Chimica, via L. Borsari 345, Univ. di Ferrara, I-44100 Ferrara, Italy): **Automated multiple development HPTLC analysis of sugars on hydrophilic layers: II. Diol layers.** J. Planar Chromatogr. **10**, 31-37 (1997). HPTLC of sugars (i.a. glucose, isomaltotetraose, isomaltotriose, isomaltose, raffinose, nystose, 1-kestose, lactose, lactulose, sucrose, galactose, fructose, arabinose, xylose, ribose, rhamnose) on diol with AMD using a fifteen-step ACN - water gradient with water concentration decreasing linearly from 35 to 15%. Detection by absorbance at 515 nm after derivatization with 4-aminobenzoic acid reagent or a-naphthol reagent by immersion for 2 min. After drying for 2 min finally heating at 100-120°C. Quantification by densitometry at 365 nm (fluorescence) and at 400 resp. 515 nm (absorbance).

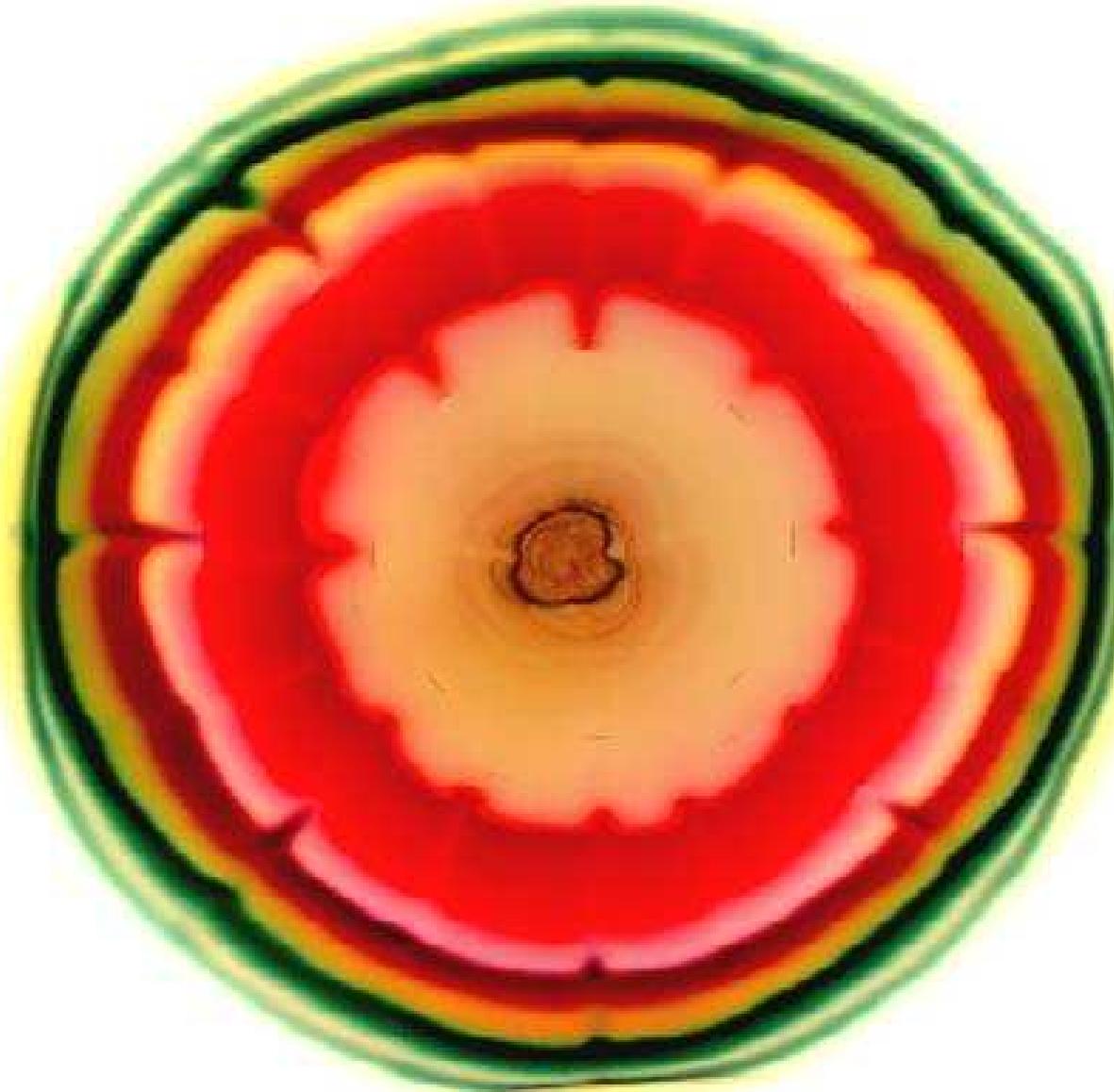
Food analysis, quantitative analysis, densitometry, AMD

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