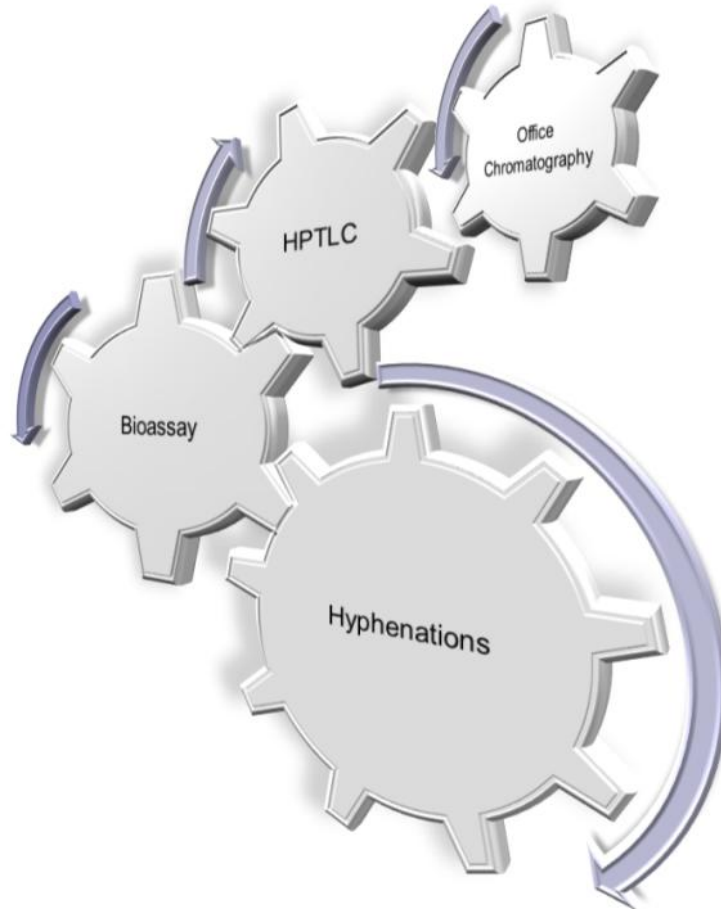


# Overview on the power of hyphenated HPTLC analysis of food and herbs



Gertrud Morlock, Chair of Food Science  
 Justus Liebig University Giessen

# Analysis of wax ester in fish (*Gempylidae*)

Stearyl stearate  
Oleyl oleate

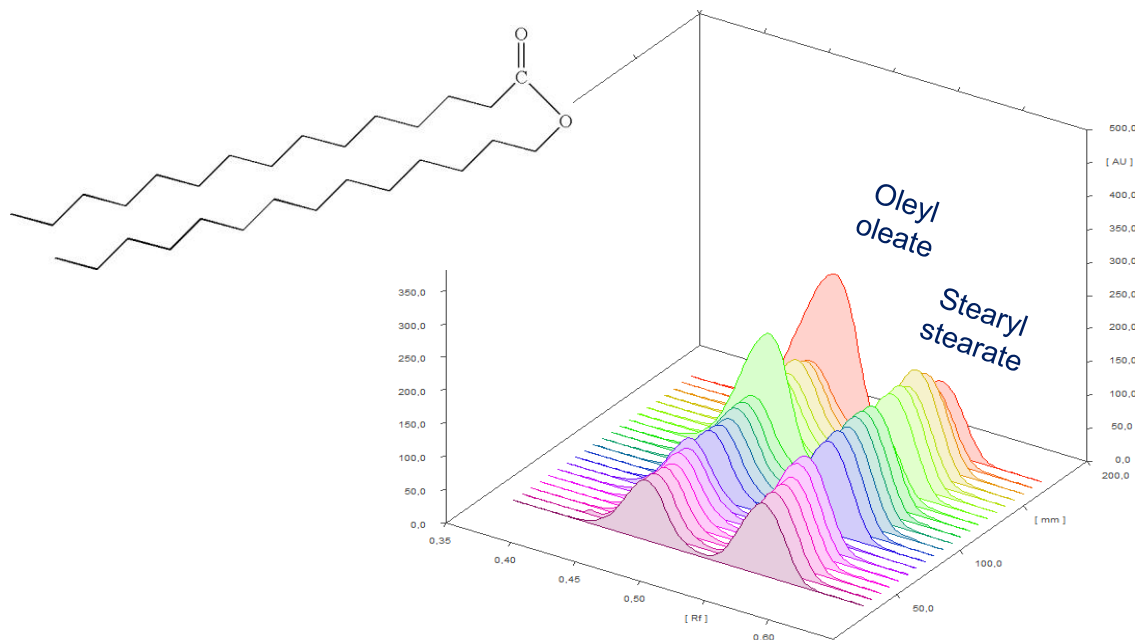
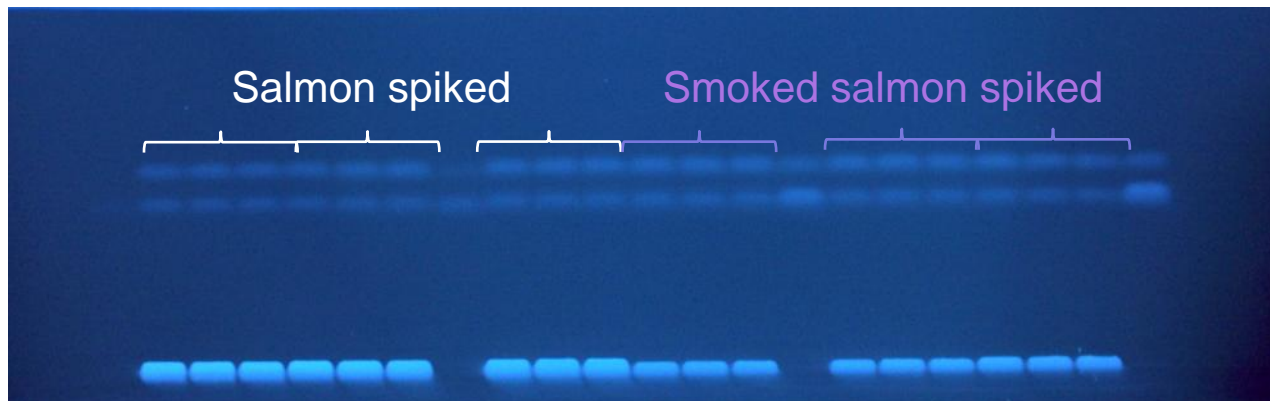


Figure 1. Oilfish, *Ruvettus pretiosus*



Figure 2. Escolar, *Lepidocybium flavobrunneum*



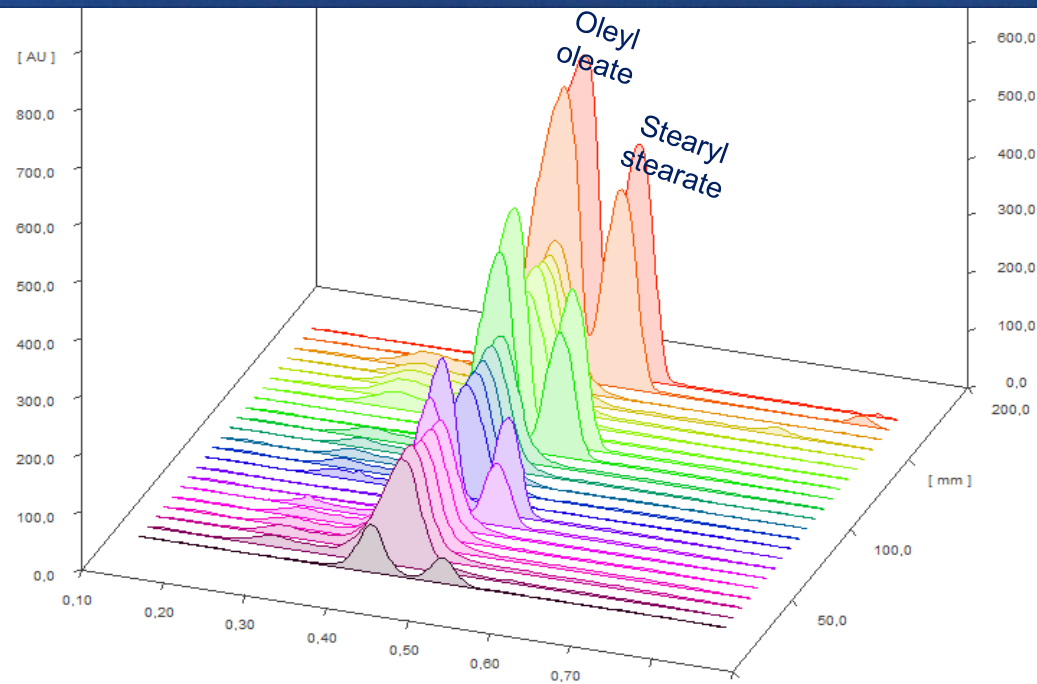
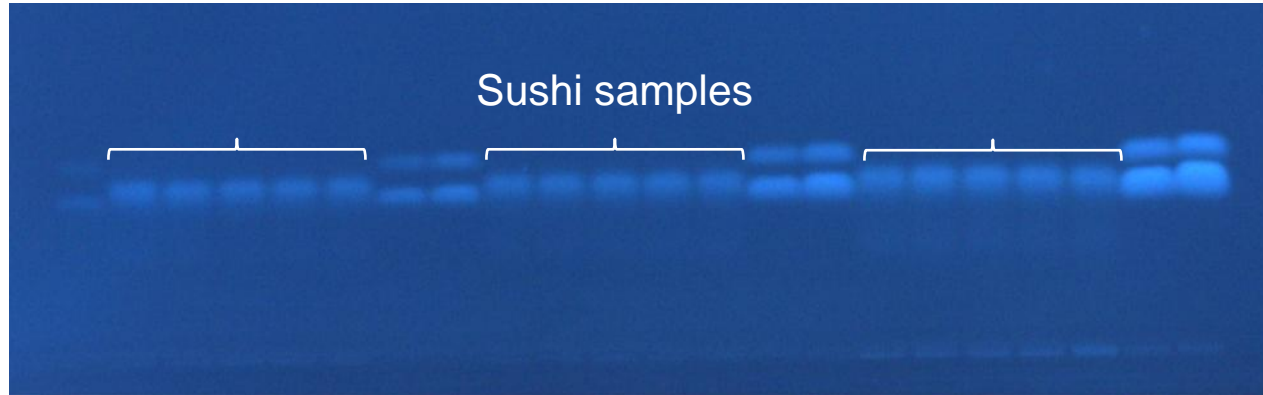
Figure 3. Rudderfish, *Centrolophus niger*



Figure 4. Rudderfish, *Tubbia* sp.

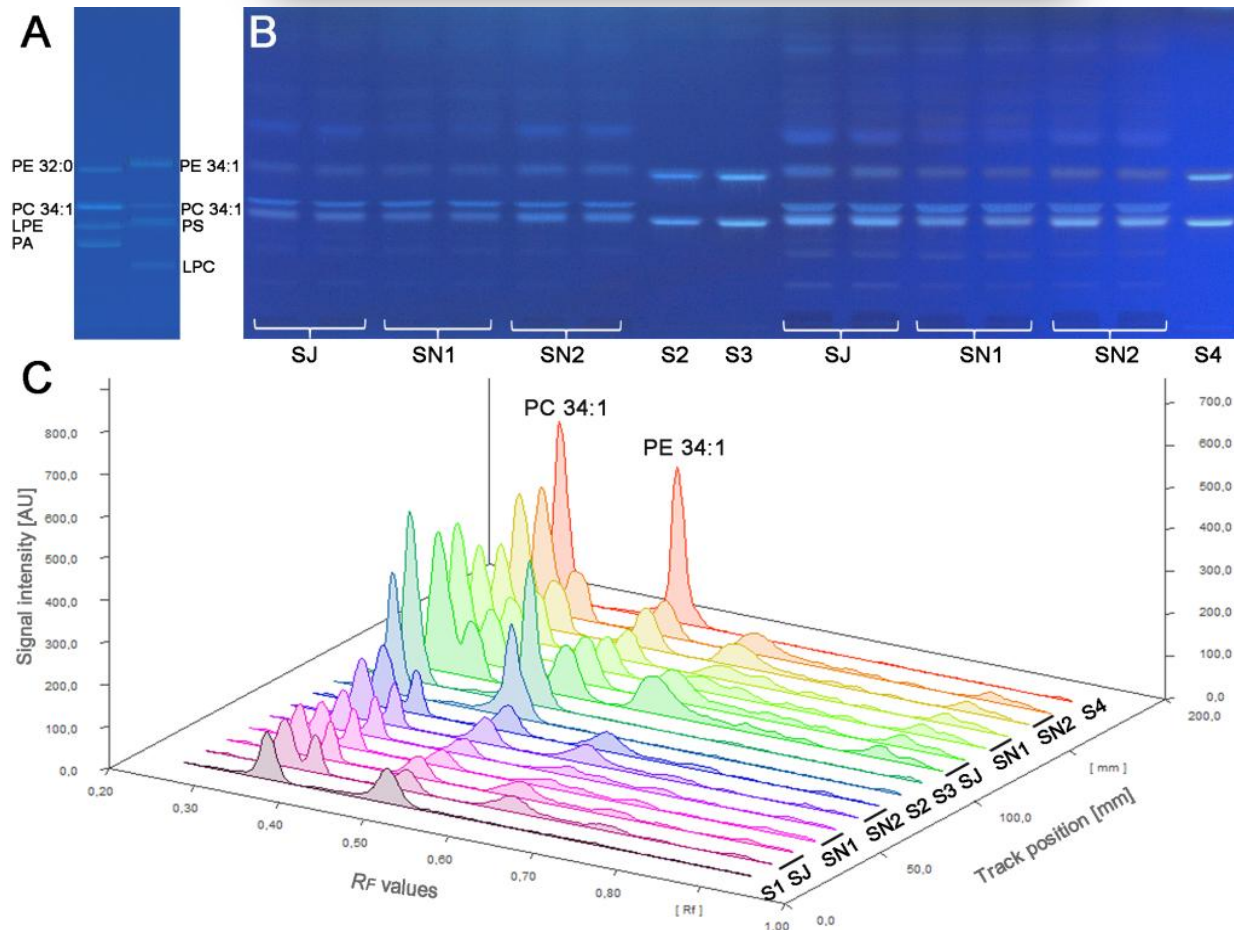


# Analysis of wax ester in fish (*Gempylidae*)

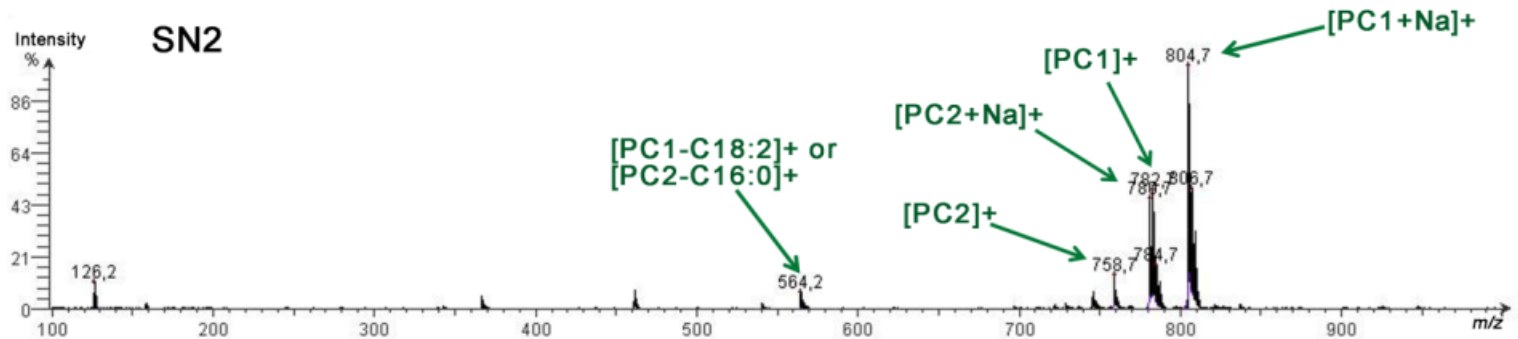
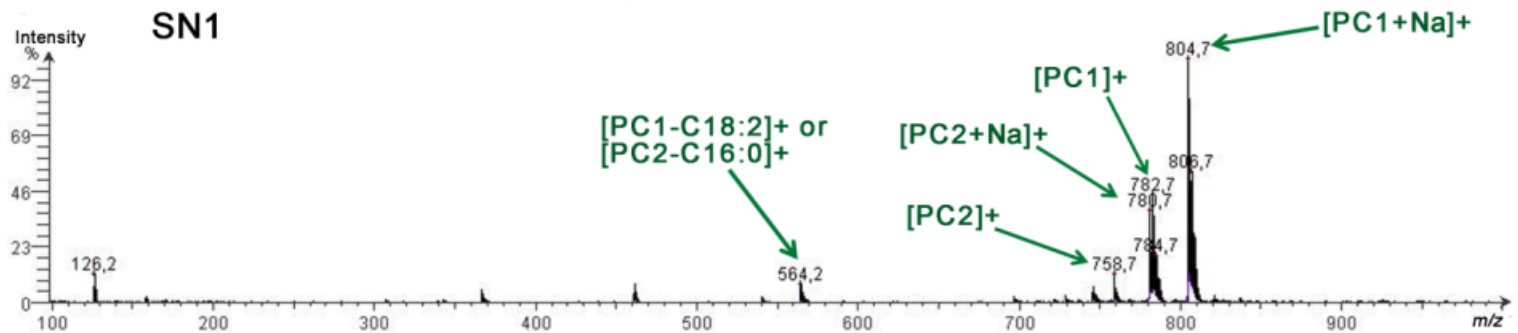
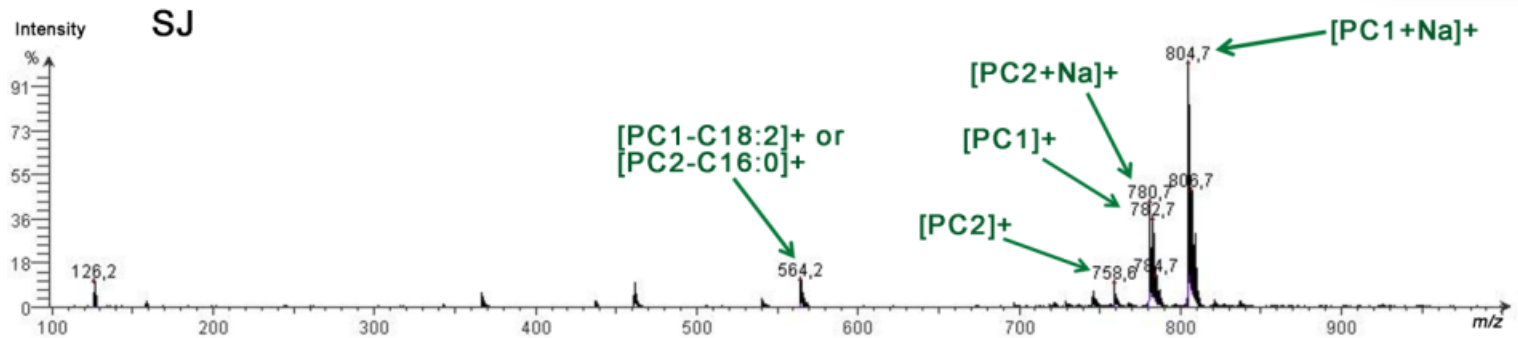


L. Winheim, A. Miller, G. Morlock, in preparation

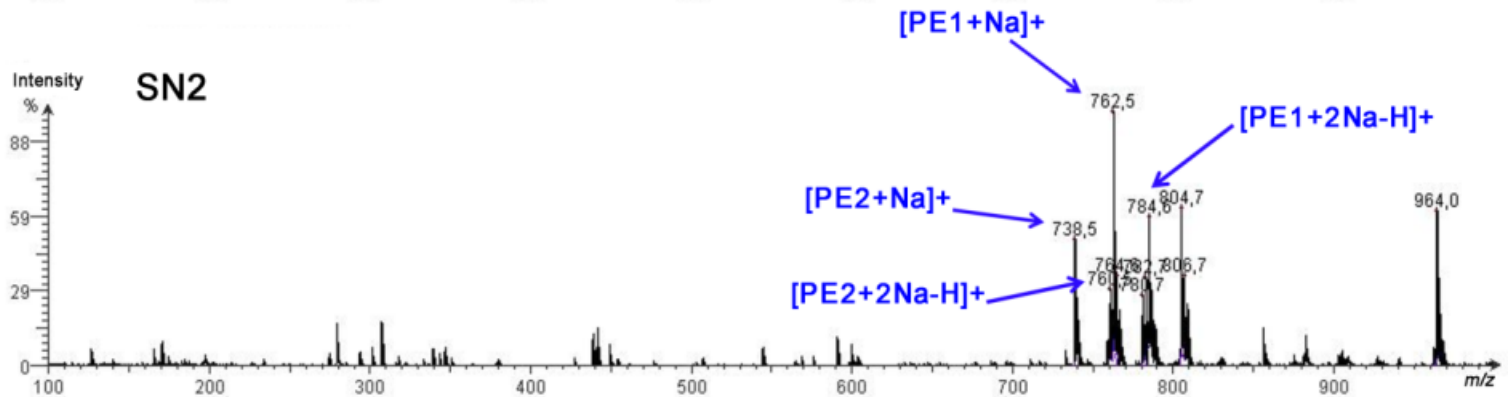
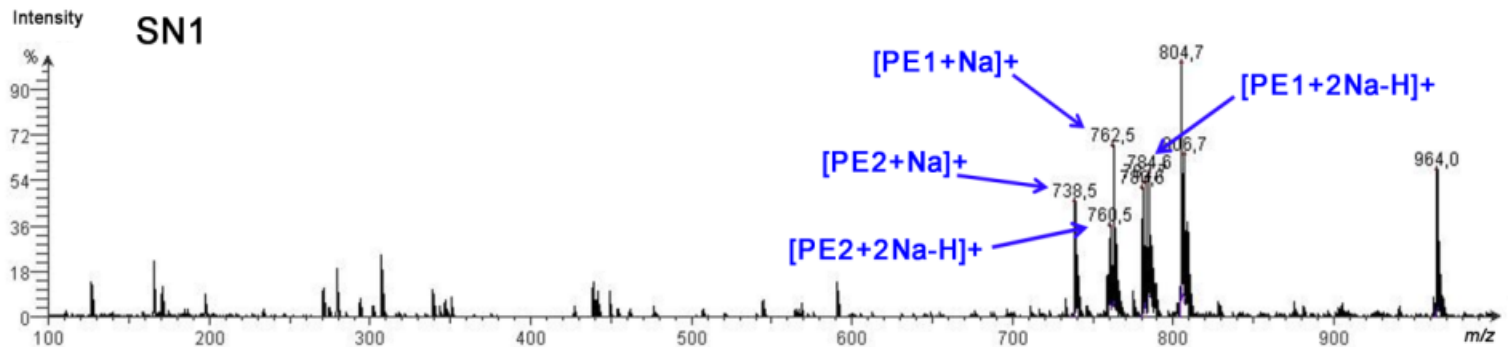
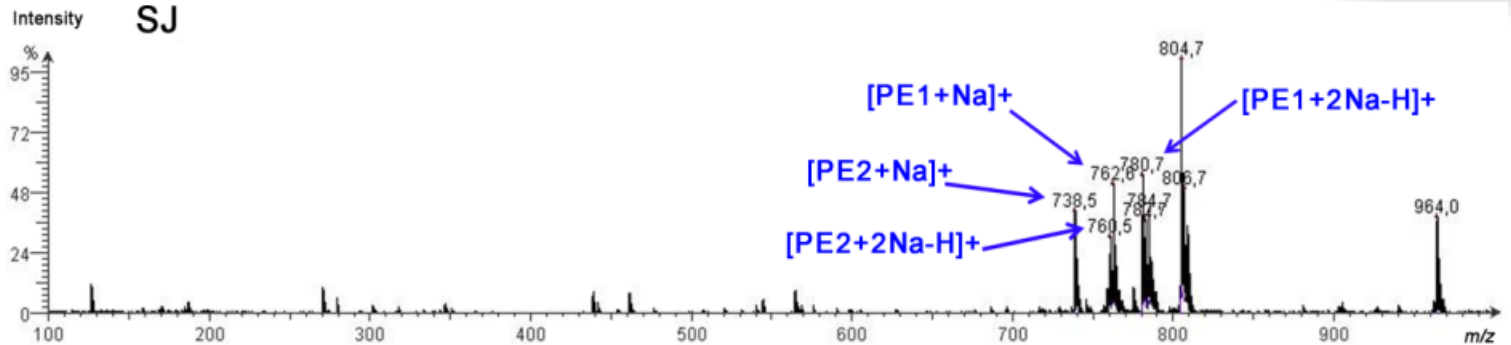
# Analysis of phospholipids in lecithins



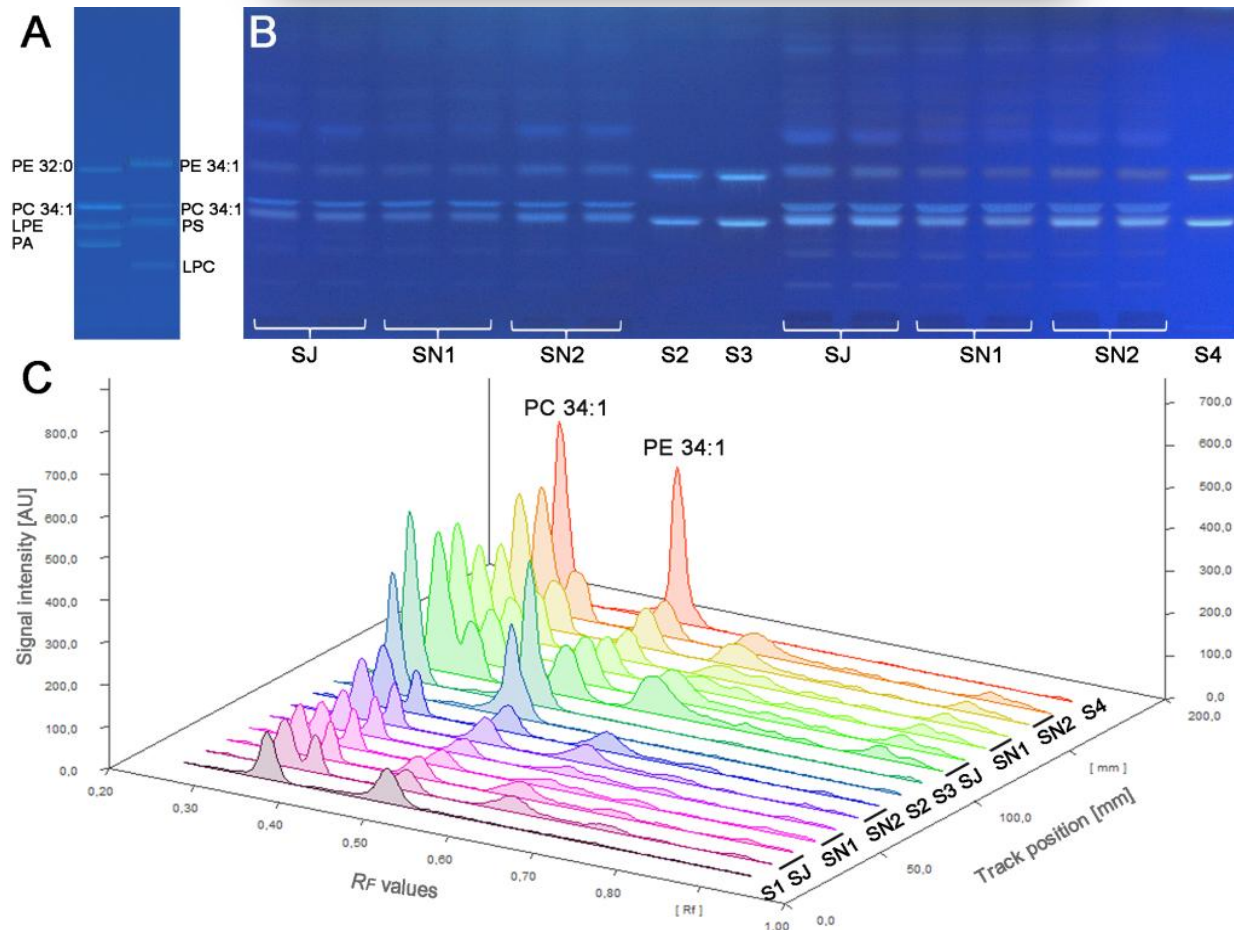
# Characterization: soy bean vs. sunflower lecithin



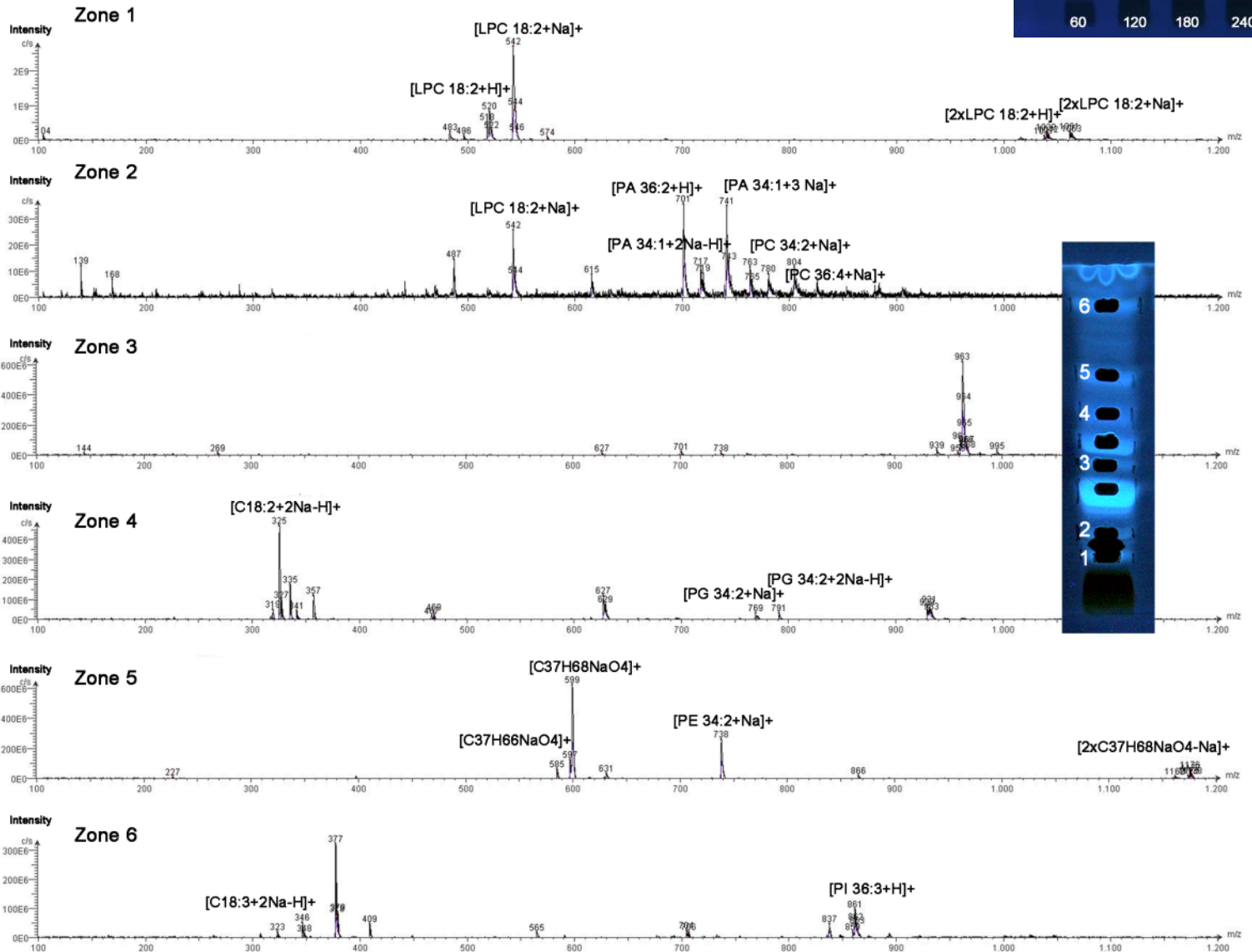
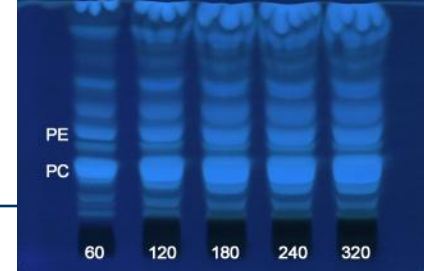
# Characterization: soy bean vs. sunflower lecithin



# Analysis of phospholipids in lecithins



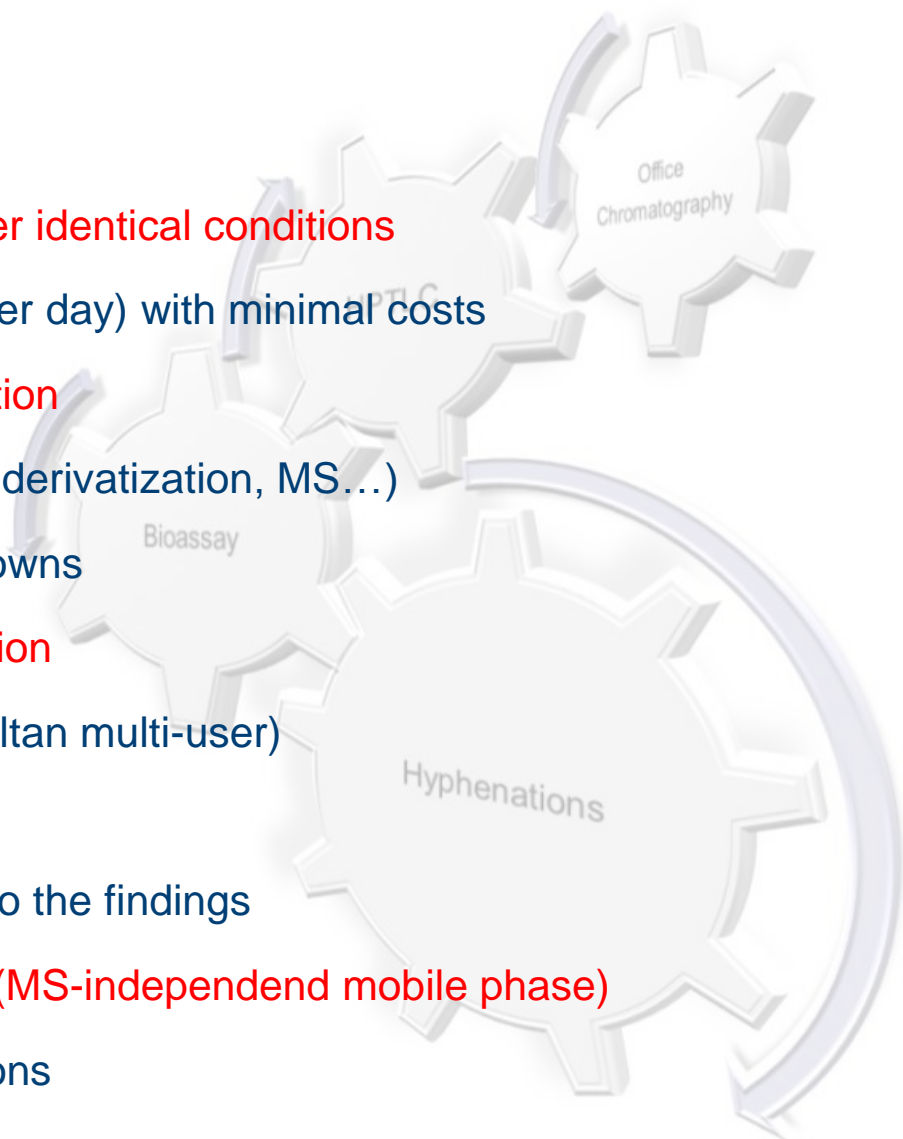
# Minor phospholipids?





# The power of HPTLC

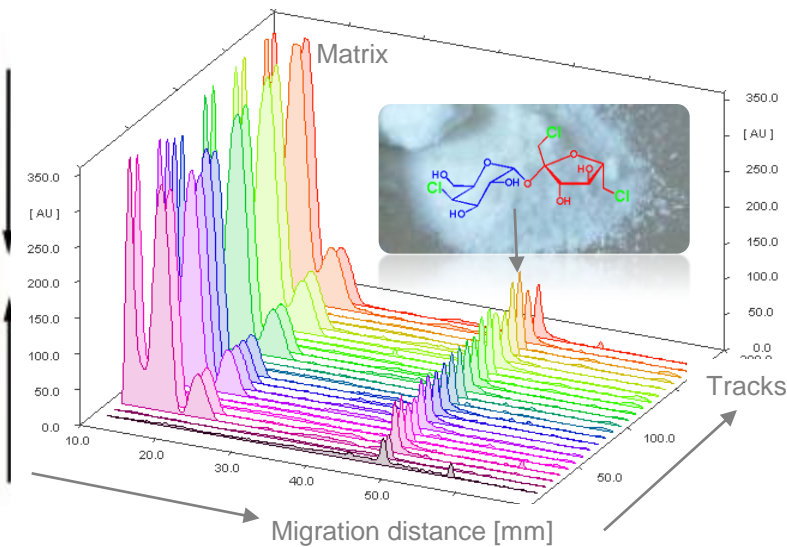
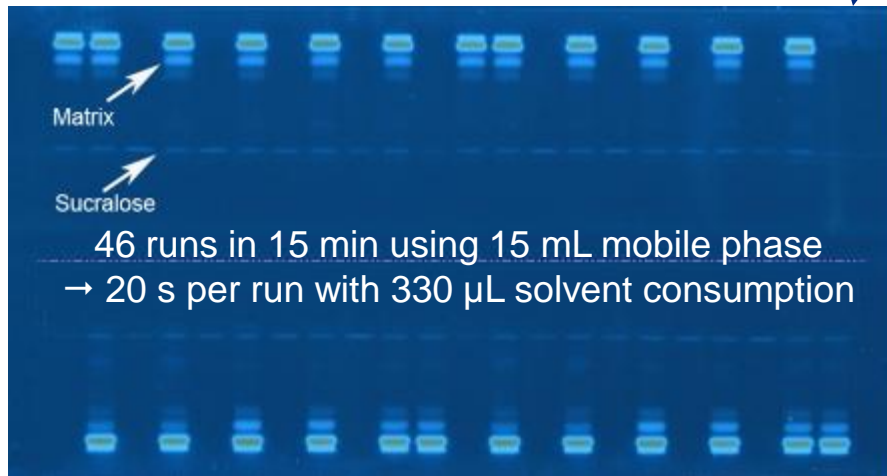
1. Reduced sample preparation
2. Matrix-tolerant method
3. Parallel chromatography under identical conditions
4. High throughput (1000 runs per day) with minimal costs
5. Selective, simultan derivatization
6. Multi-detection (UV/Vis, FLD, derivatization, MS...)
7. More information about unknowns
8. Concentration during application
9. Flexible working station (simultan multi-user)
10. Effect-directed analysis
11. Analytical workflow adjusted to the findings
12. Targeted mass spectrometry (MS-independend mobile phase)
13. The ease of super-hyphenations



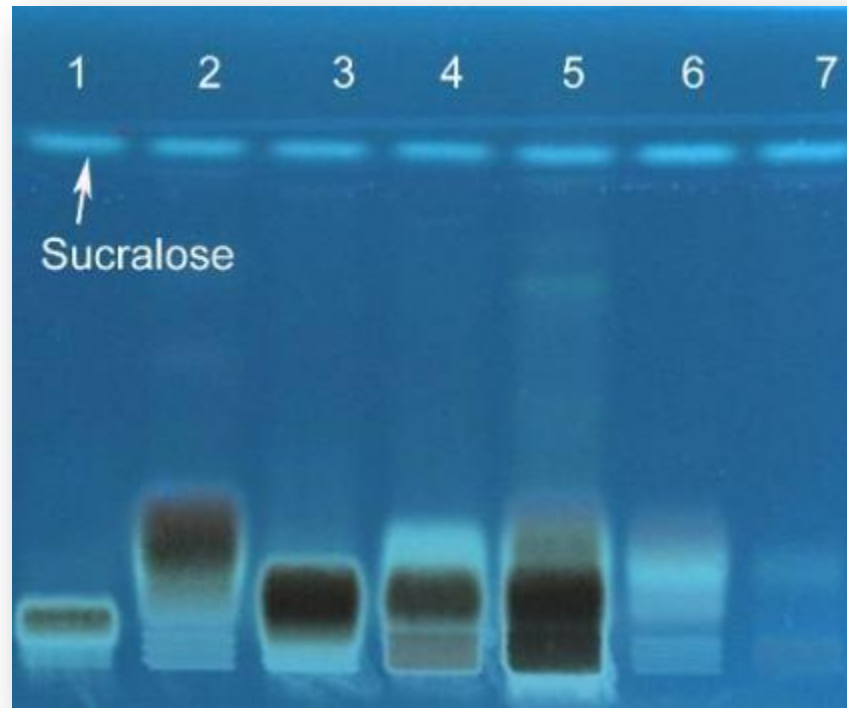
# Sucralose in milk-based confection (*Burfi*)



extracted in MeOH, filtered

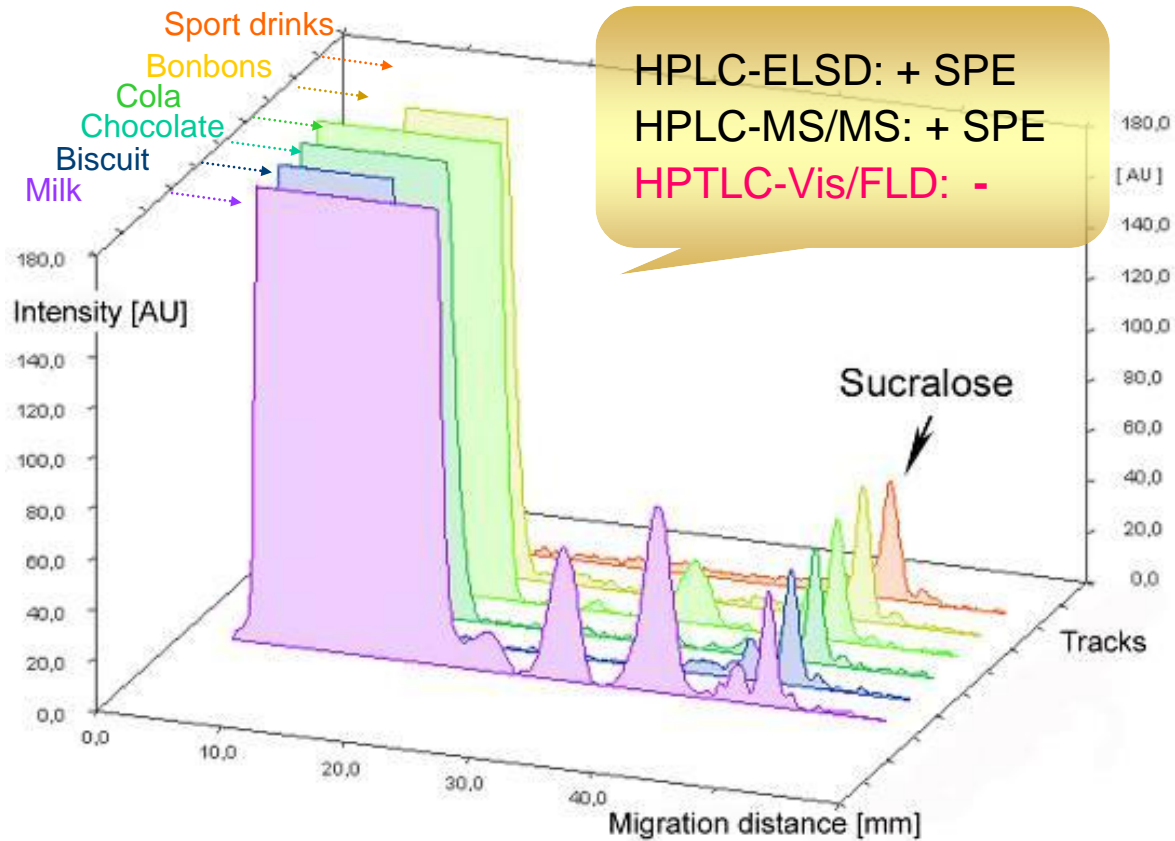


# ... in further matrices



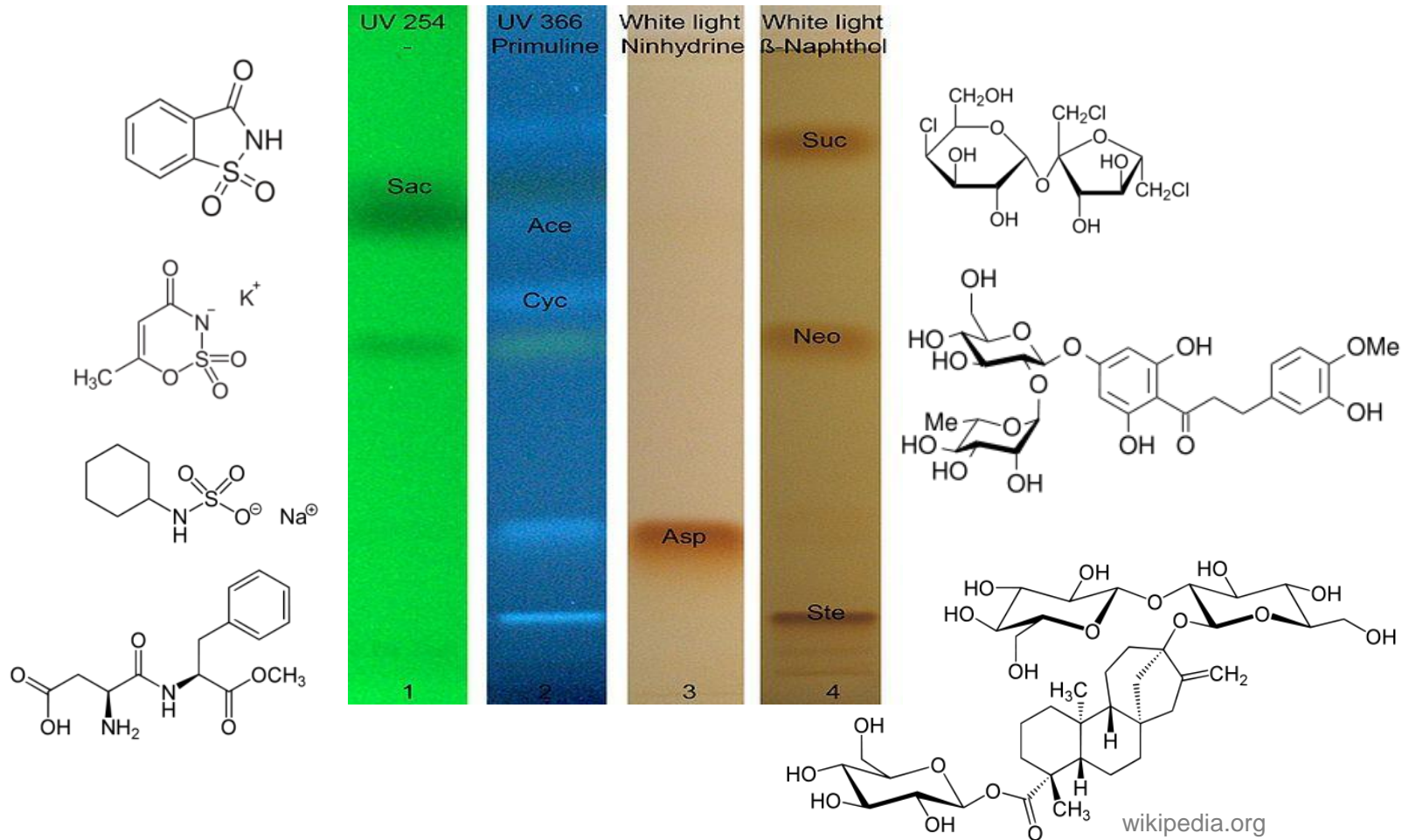
Milk, biscuit, chocolate, cola, bonbons, energy/sport drinks

# Sample preparation and chromatography

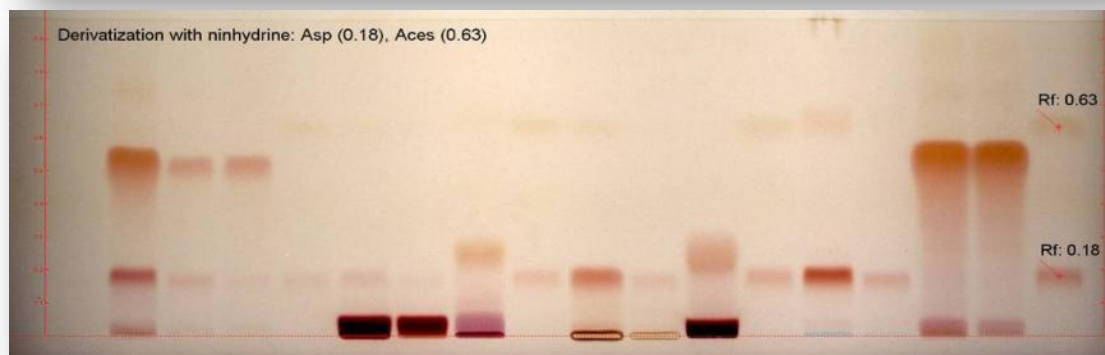
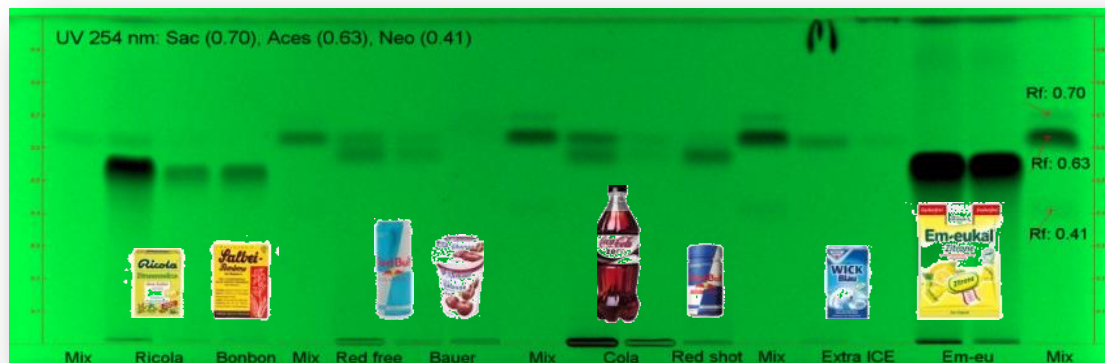


# Analysis of sweeteners

→ Reagent sequence: 4 reagents on the same plate

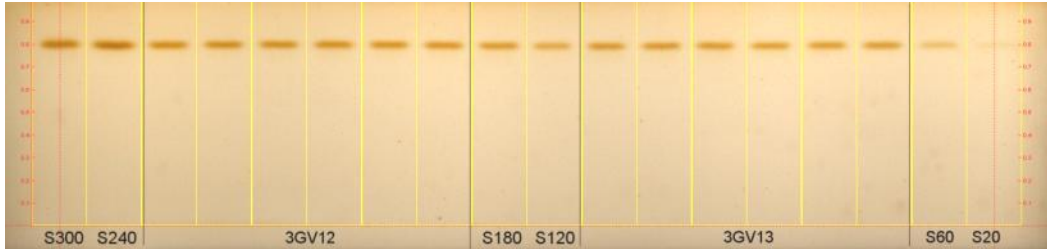


# Food safety → Sweeteners

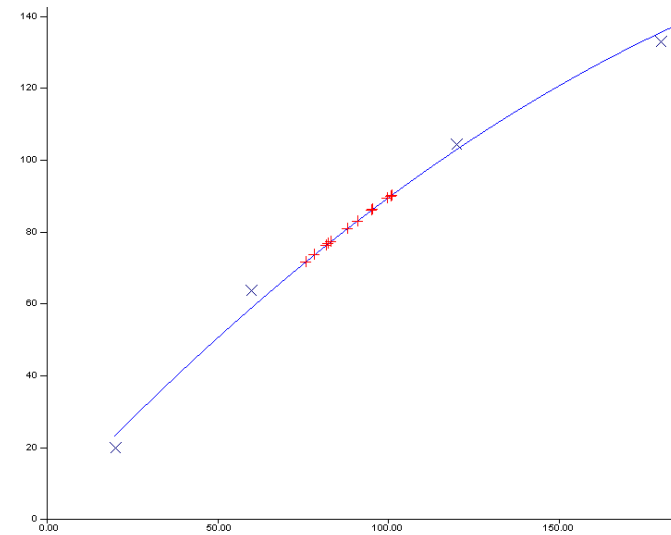
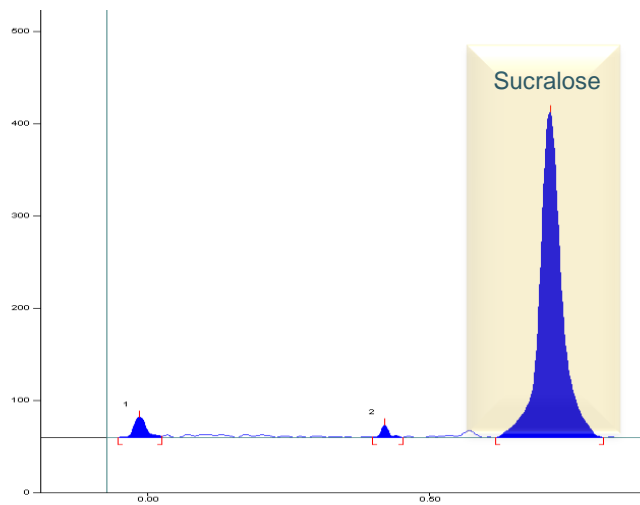


# Sucralose in saliva

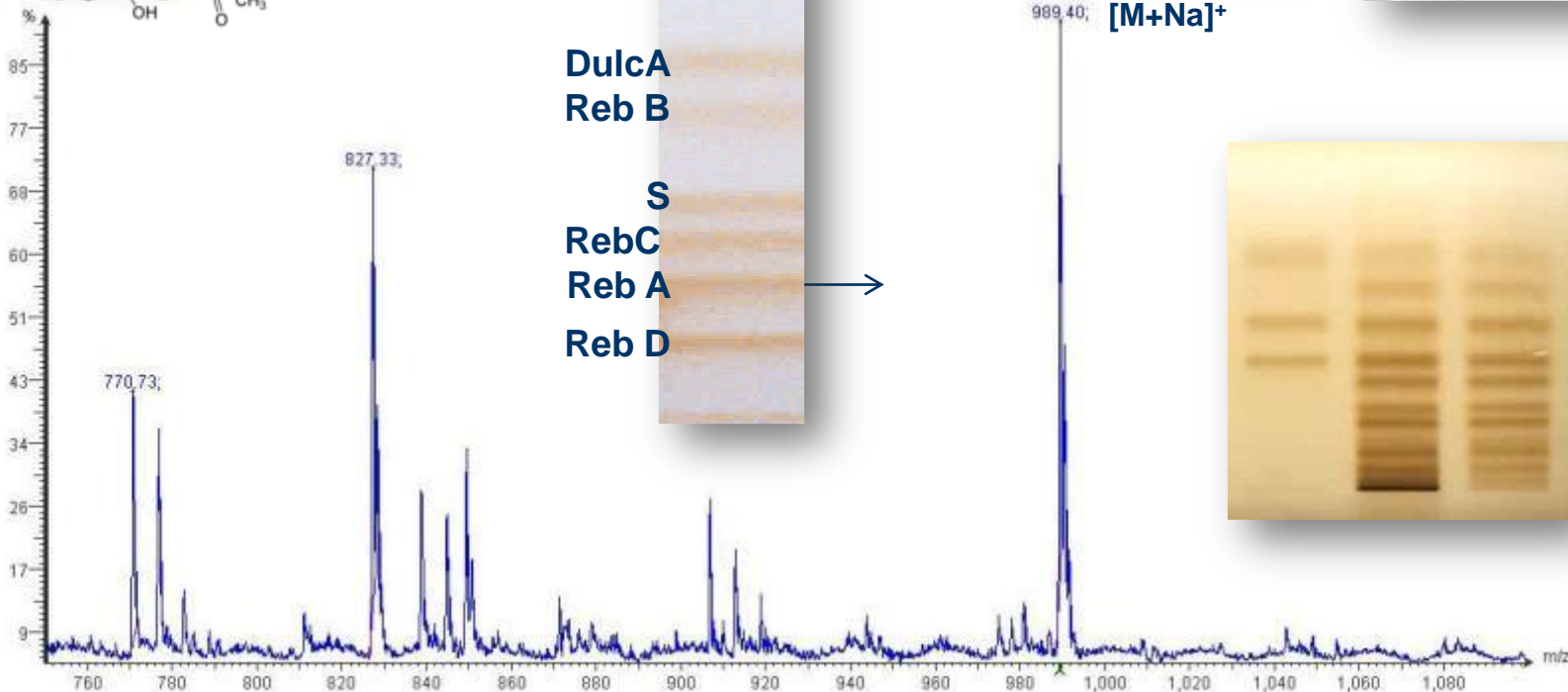
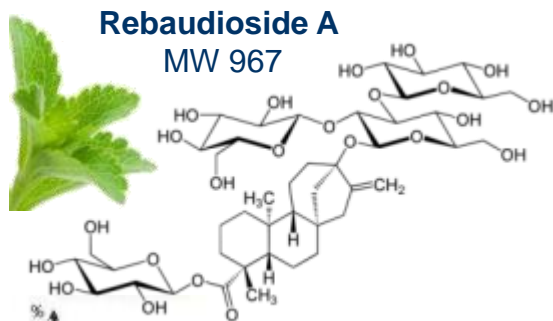
→ Release study from chewing gum



[www.media.news.de/resources](http://www.media.news.de/resources)



# Analysis of steviol glycosides

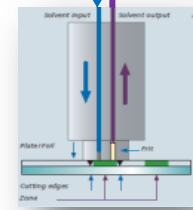


SB

DulcA  
Reb B

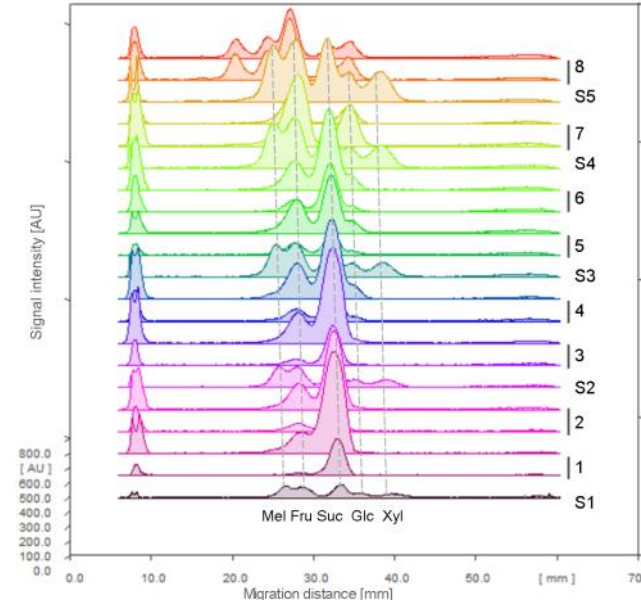
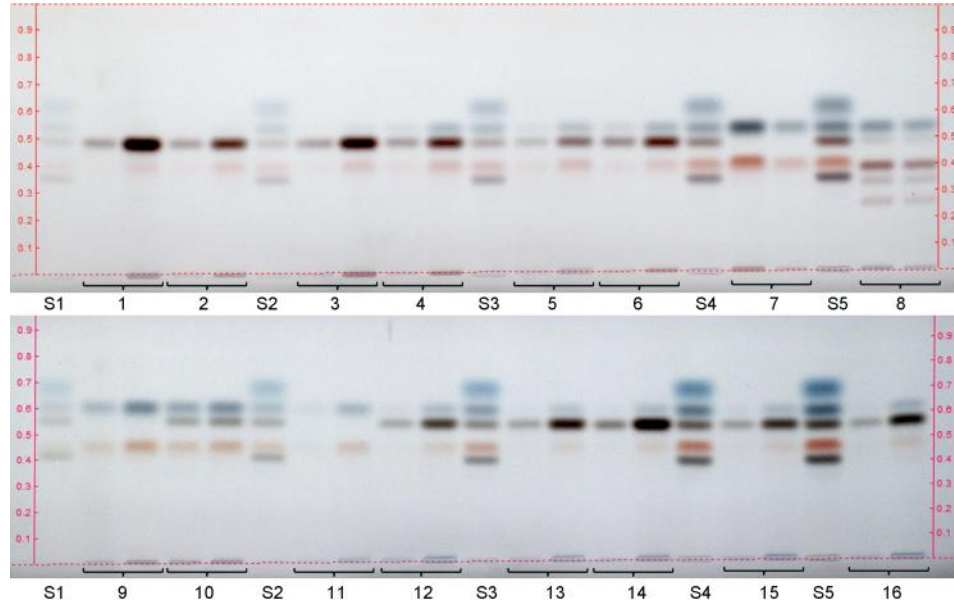
S

RebC  
Reb A  
Reb D



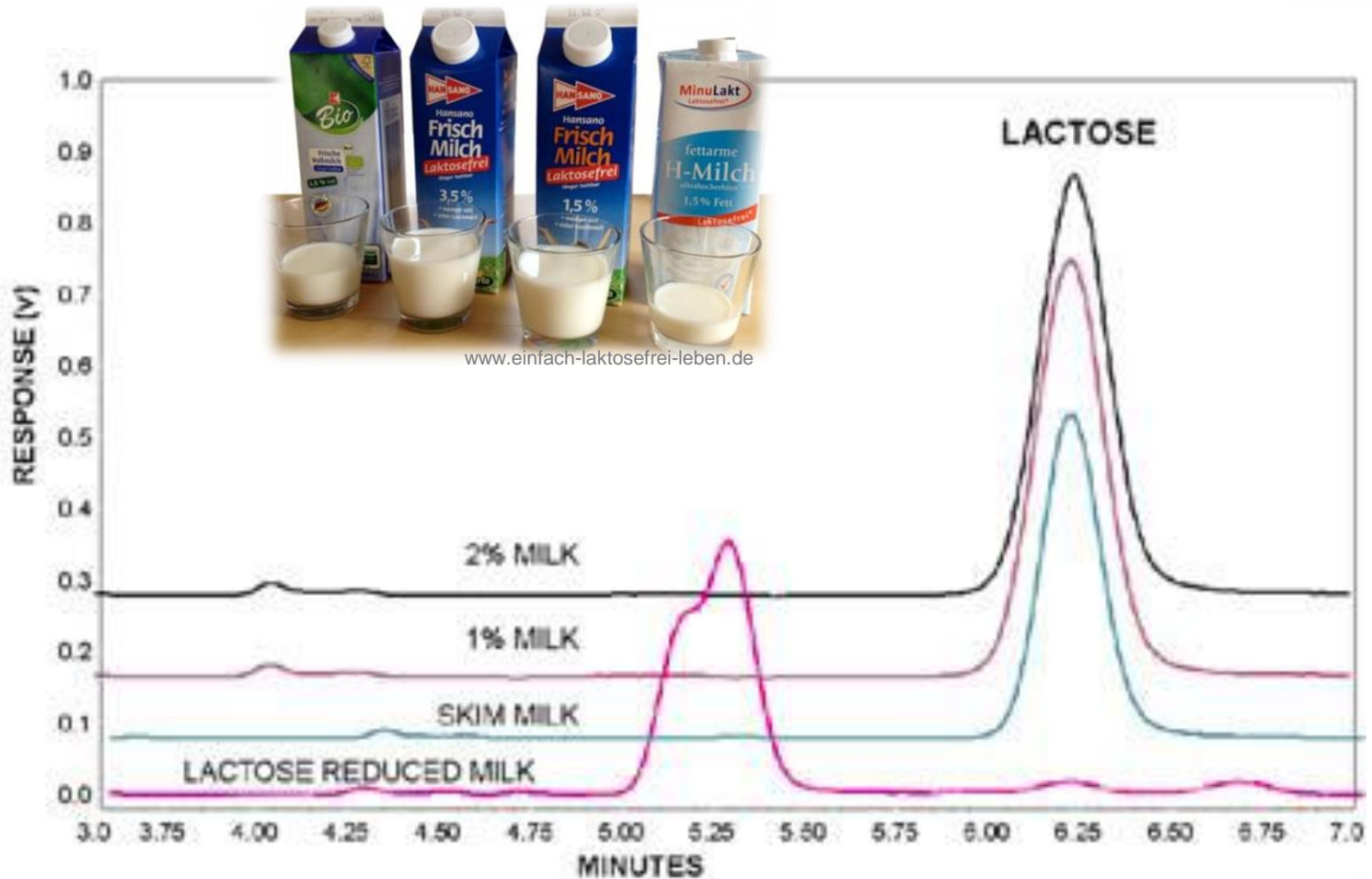


# Carbohydrates in *Marantaceae* nectars



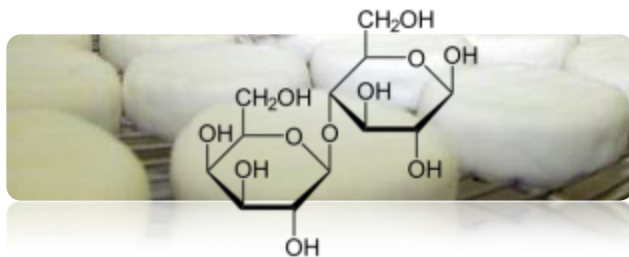
	<i>C. lutea</i> So	<i>C. lutea</i> S	<i>C. crotalifera</i> So	<i>C. crotalifera</i> S	<i>C. picturata</i> S	<i>C. hylaeanthoides</i> S	<i>C. platystachya</i> S	<i>C. lasiophylla</i> So	<i>C. lasiophylla</i> S	<i>C. donnell-smithii</i> S	<i>C. marantifolia</i> S	<i>H. hofmannii</i> S	<i>Ct. Dasycarpa</i> S	<i>Ma. Purpurea</i> S	<i>P. pruinosa</i> So	<i>P. pruinosa</i> S
N	10	12	5	6	1	1	6	5	7	6	7	7	3	6	6	7
Suc	24.85	28.92	22.57	18.47	3.21	16.32	26.38	17.34	21.52	41.55	43.21	41.14	24.03	19.87	41.4	31.25
Glc	8.06	6.10	2.99	3.06	0.00	1.42	1.41	1.83	1.63	1.13	1.36	0.00	1.02	0.00	4.22	2.68
Fru	10.66	7.22	4.40	4.40	0.00	3.17	1.90	2.56	2.25	2.58	3.10	0.00	3.62	0.00	8.05	4.76
Suc / Glc + Fru	1.327	2.170	3.053	2.473	/	3.548	7.970	3.945	5.527	11.157	9.659	/	5.167	/	3.371	4.196
Glc / Fru	1.322	1.183	1.471	1.440	/	2.225	1.348	1.399	1.399	2.279	2.279	/	3.534	/	1.905	1.778

# Analysis of lactose-free food: HPLC-CAD

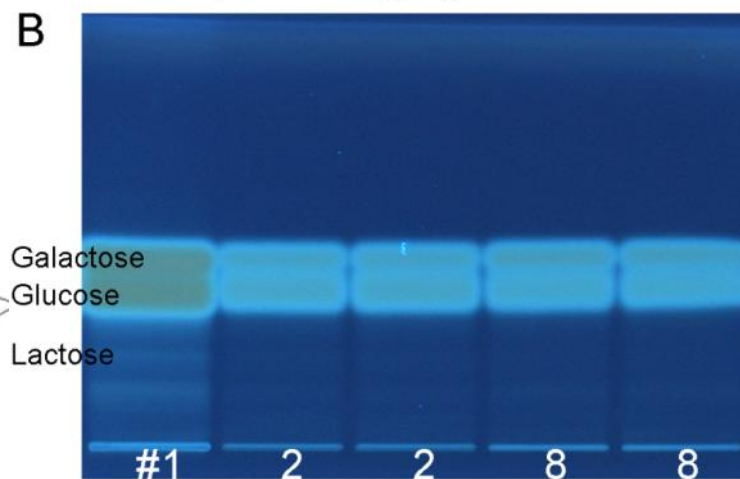
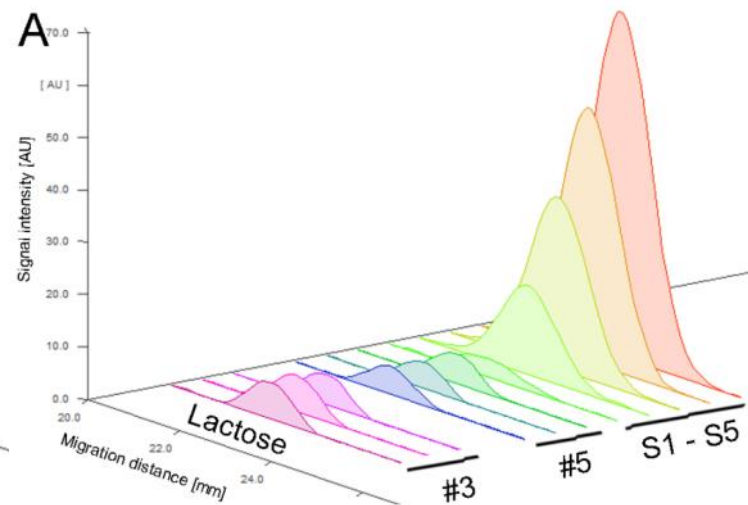
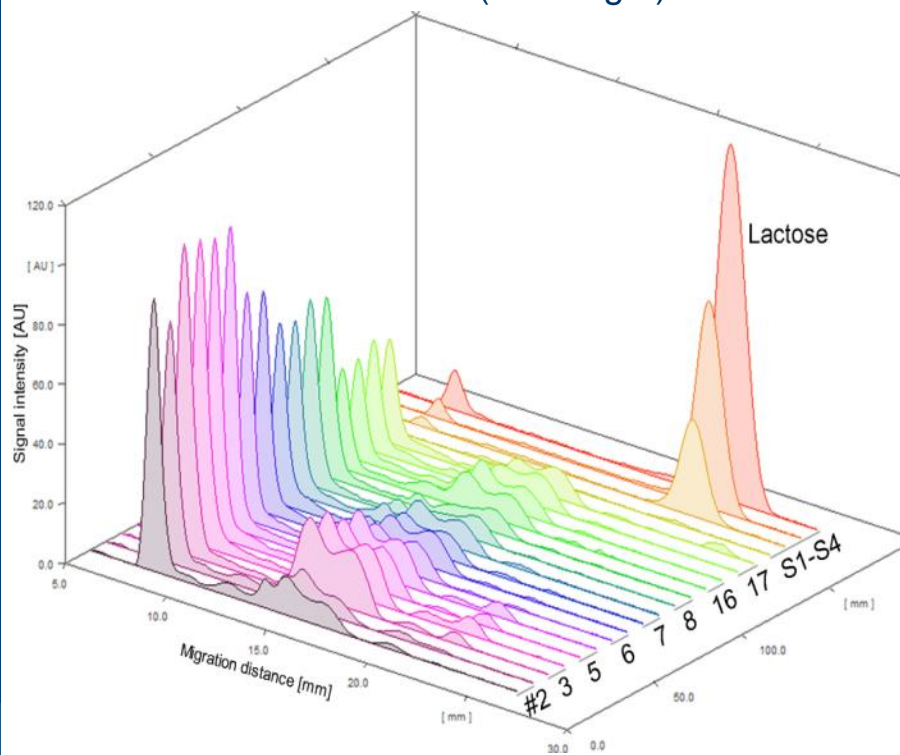


ESA, Dionex, application note 2011: Lactose in milk - a validated method

# Analysis of lactose-free food



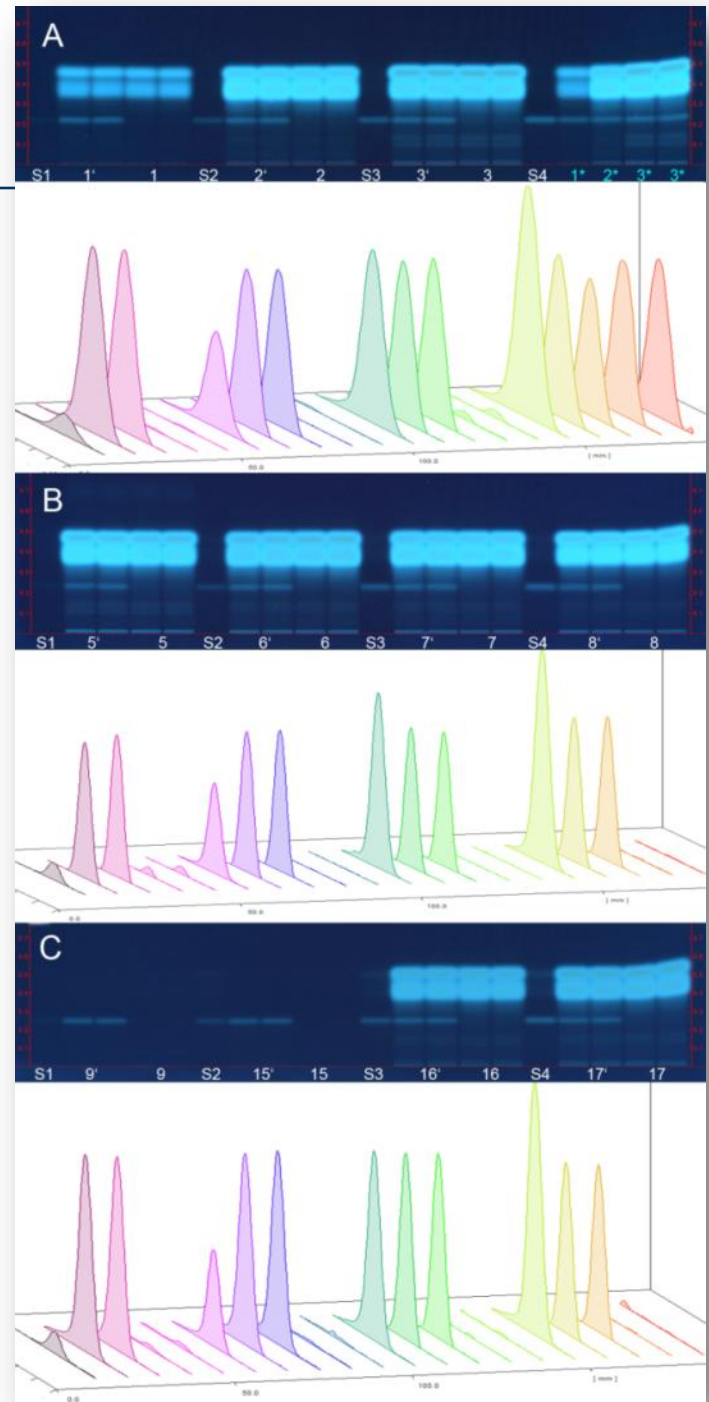
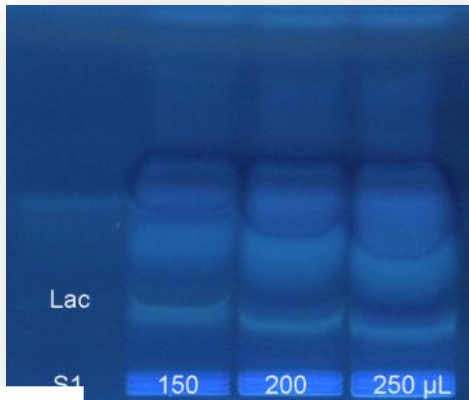
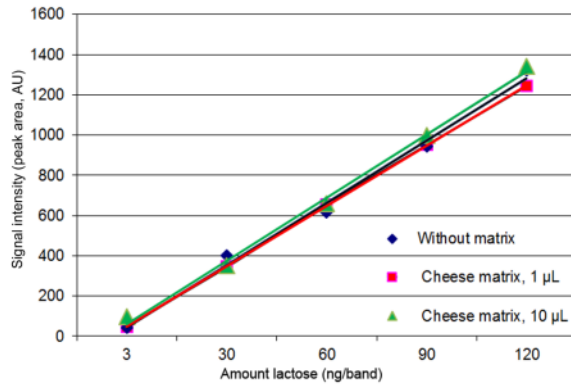
LOD  $\rightarrow$  0.000004 % (0.04 mg/L)



# Rectangular application

Lactose-free dairy products

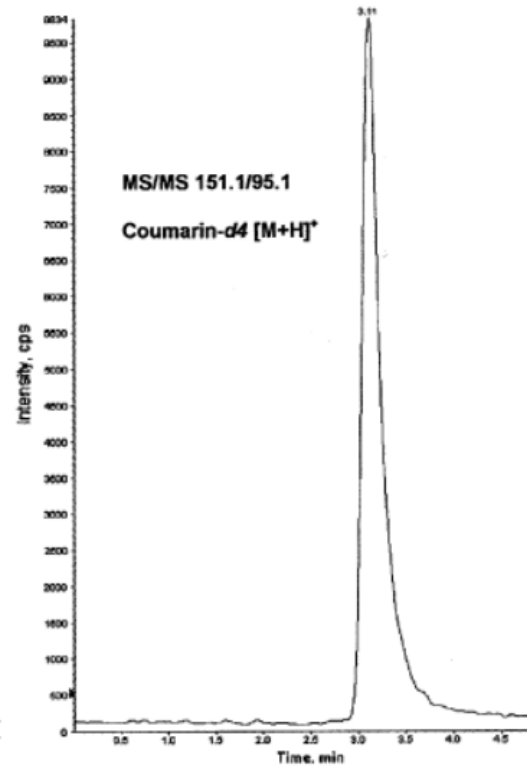
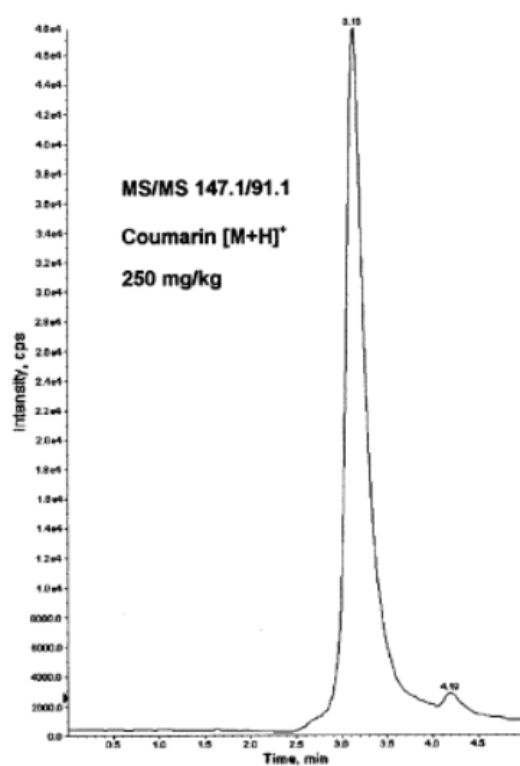
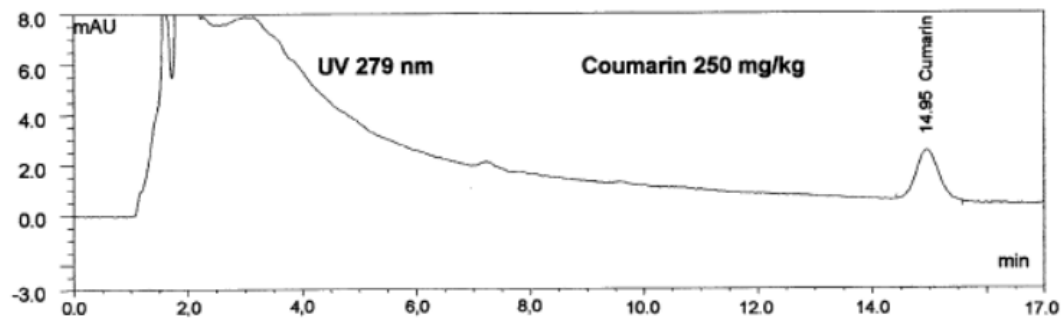
LOD  $\rightarrow$  0.000004 % (0.04 mg/L)



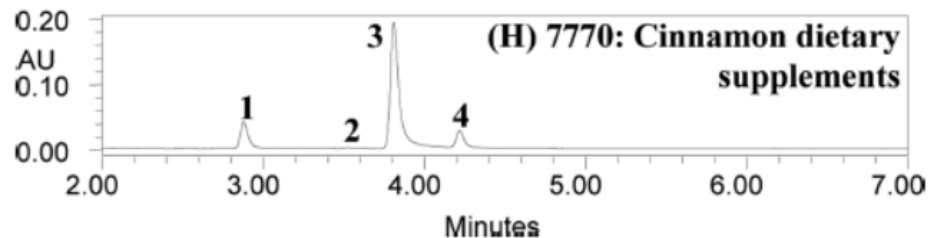
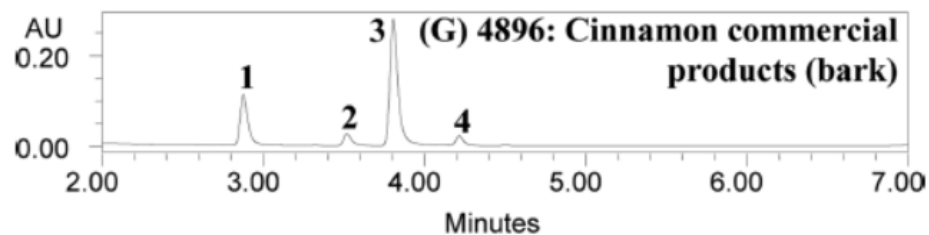
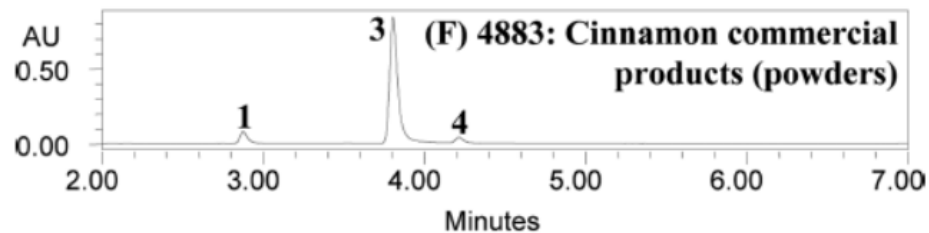
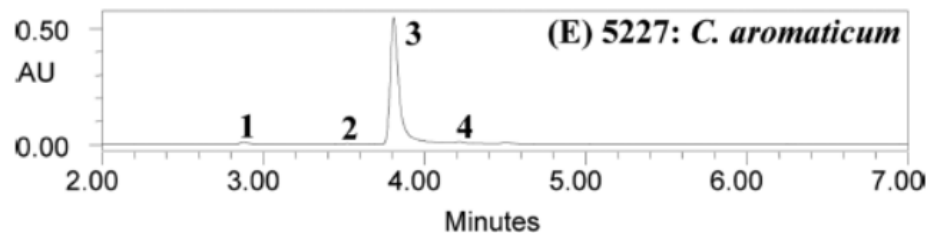
# Recovery rates in lactose-free food

Lactose-free dairy product	#	Mean recovery of lactose (%)	Repeatability (%RSD, n = 2)
Butter	1	105.3	1.9
Yoghurt	2	87.7	0.7
Milk	3	87.7	0.6
Evaporated milk	5	76.2	3.1
Buttermilk	6	81.2	0.2
Sour cream	7	78.9	2.9
Cream	8	82.8	0.2
Goat cheese	9	103.6	1.4
Cheese	15	102.4	1.2
Cream cheese	16	99.2	0.5
	17	90.3	1.8
Mean ± %RSD (n = 11)		90.5 ± 10.5	1.3 ± 1.0

# Cumarine analysis: HPLC-UV *and* -MS

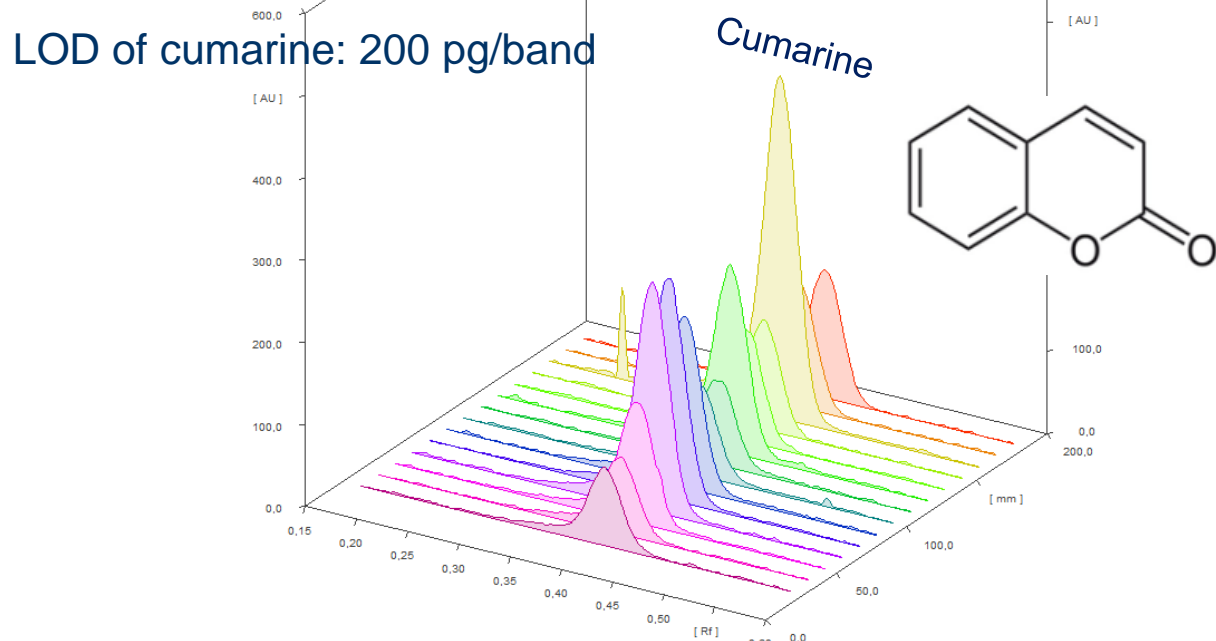


# Cumarine analysis: UPLC-UV



# Analysis of cumarine

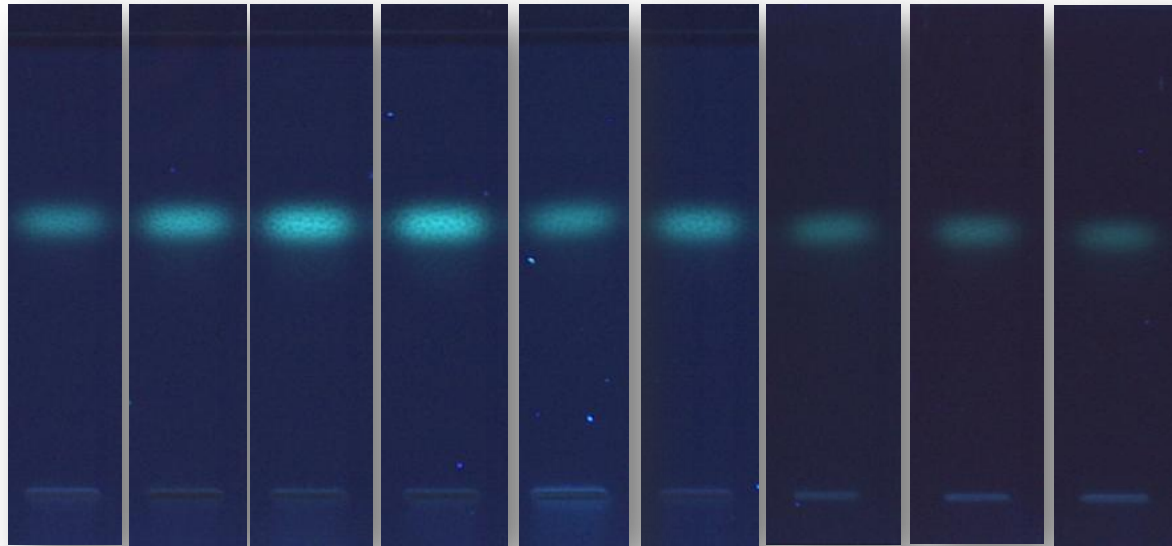
5 different cinnamom spice samples containing cumarine





# Analysis of cumarine

9 different food samples containing cumarine



[www.wikipedia.de](http://www.wikipedia.de)



[www.colourbox.de](http://www.colourbox.de)



[www.deutsche-wirtschafts-nachrichten.de](http://www.deutsche-wirtschafts-nachrichten.de)

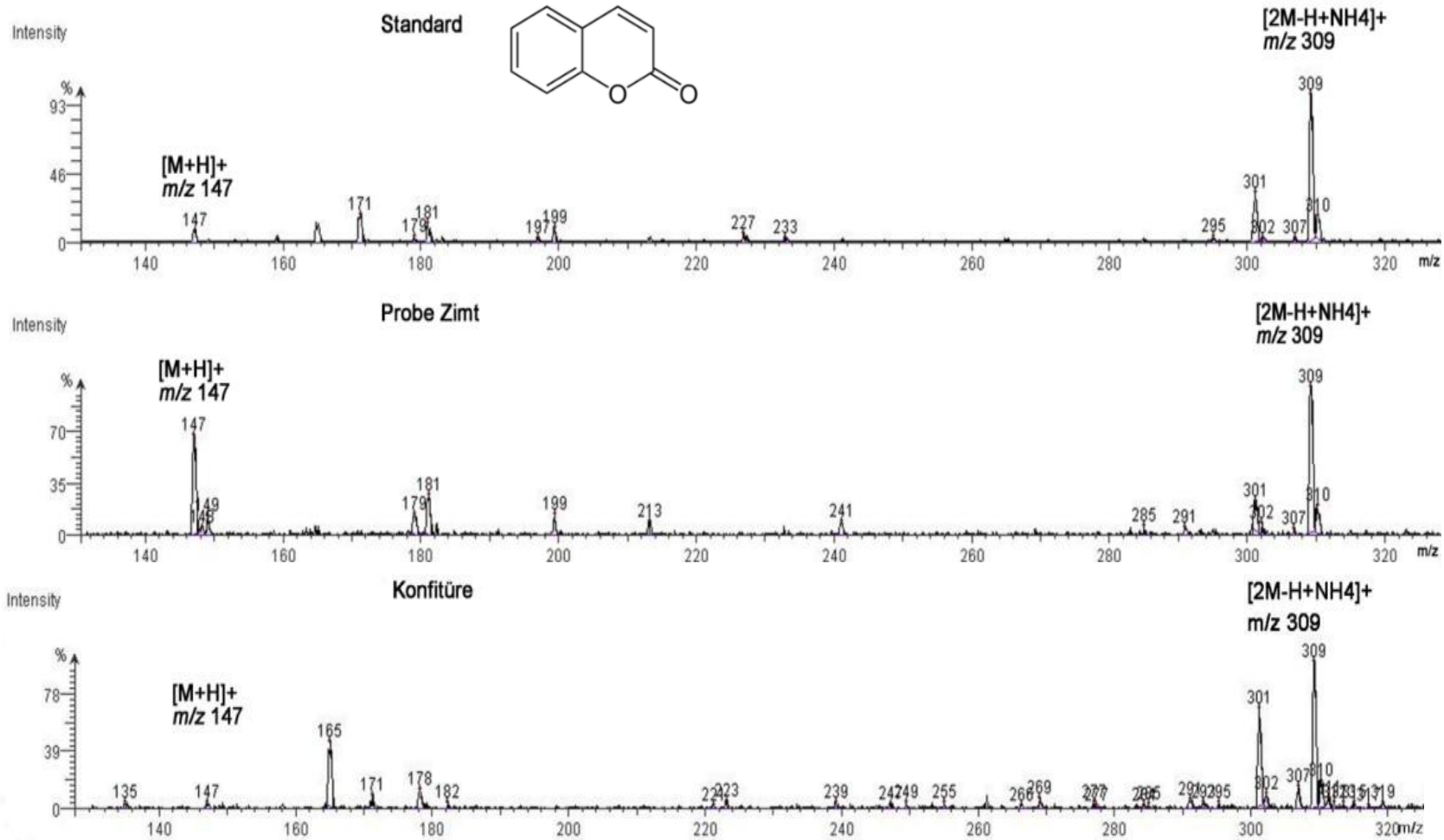


[www.institut-fresenius.de](http://www.institut-fresenius.de)

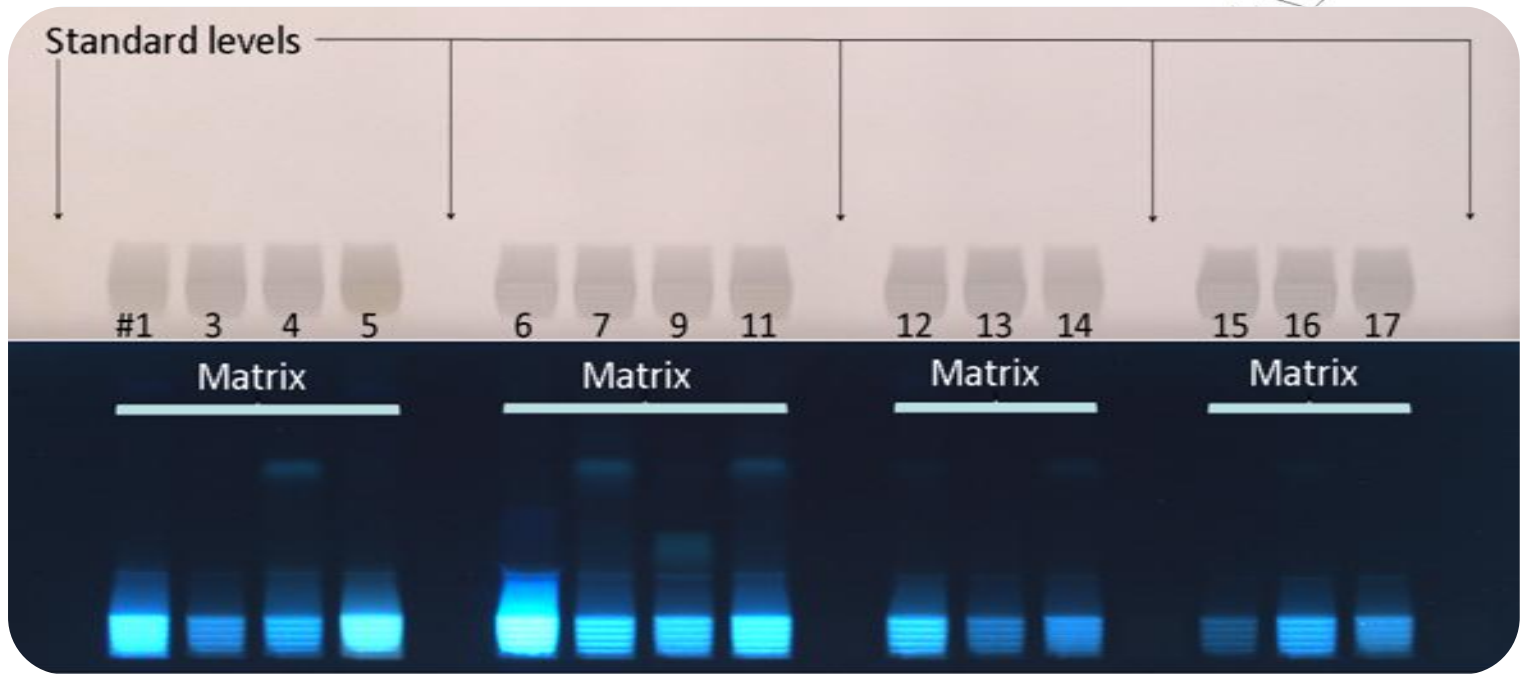
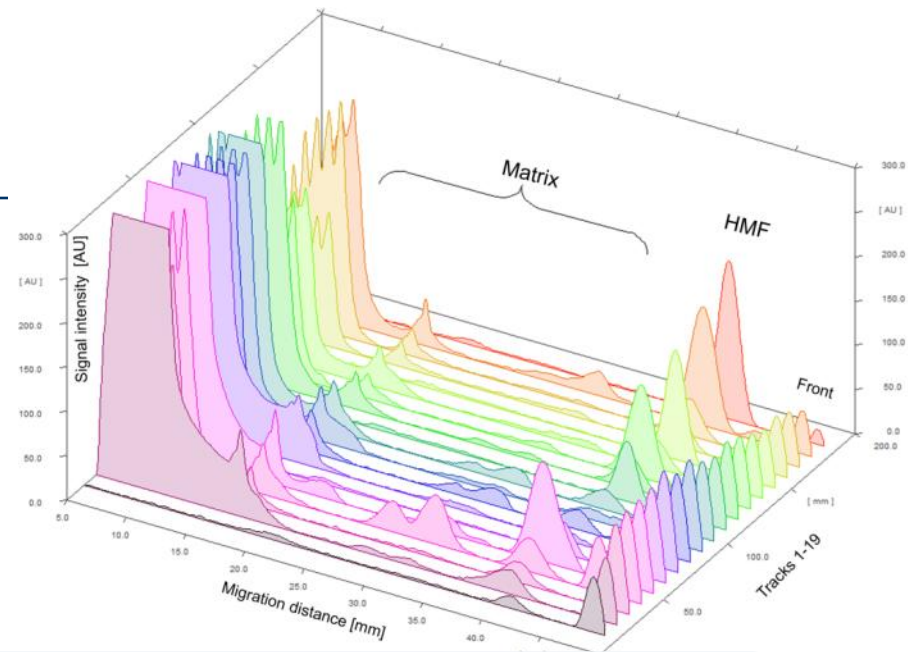
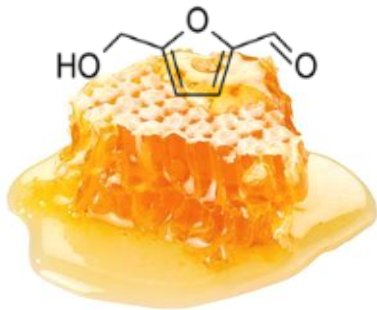
# Cumarine contents

Sample	Manufacturer	Cumarine (mg/kg)	Repeatabilities (%RSD, n=2)	Reproducibility (%RSD, n=2)
Spice	Sonnentor - gemahlen	3732	0.4/1.2	1.0
	Lidl Kania - gemahlen	1615	0.4/0.3	2.4
Tea	YogiTea - Frauenpower	19	1.4/3.2	4.5
	TeeGeschwendner – Kaminfeuer	22	3.3/2.0	0.5
Cereals	Aldi Knusperone - Zimt Chips	1	3.8/5.9	0.4
Cinnamon bun	IKEA - Kanelbullar	16	1.4/4.5	6.6
Jam	Grafschafter Konfitüre – Winterzauber	4	1.6/1.6	1.6

## Confirmation by HPTLC-ESI-MS



# Quality of honey → HMF



# Method comparison

→ Traditional methods (photometry, HPLC) *versus* HPTLC

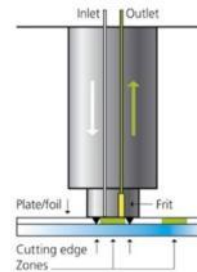
Concentration of HMF in honey						
Honey sample #	Winkler	HPLC-UV		HPTLC-UV		
	HMF in honey, mg/kg	HMF in honey, mg/kg	Deviation to Winkler method, %	Mean HMF in honey, mg/kg	Deviation to Winkler method, %	Deviation to HPLC, mg/kg
1	95.3	n/a	n/a	75.2	-22	n/a
2	41.8	n/a	n/a	30.8	-30	n/a
3	46.1	38.5	-16	39.3	-16	0.8
5	17.6	13.5	-23	13.7	-20	0.2
7	21.6	18.1	-16	18.8	-13	0.7
8	40.2	30.4	-24	28.7	-26	1.7
10	23.9	n/a	n/a	25.1	+5	n/a
Mean			20		19	0.9

Cooperation with Apicultural Institute, Celle and Stuttgart

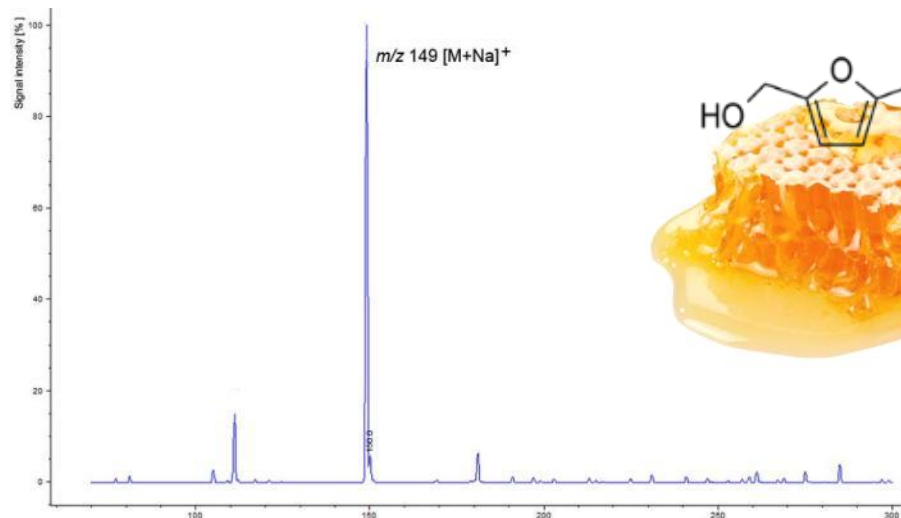
E. Chernetsova, A. Revelsky, G. Morlock, *Anal Bioanal Chem* 401 (2011) 325-332

# Method comparison

→ HPTLC: UV *versus* MS

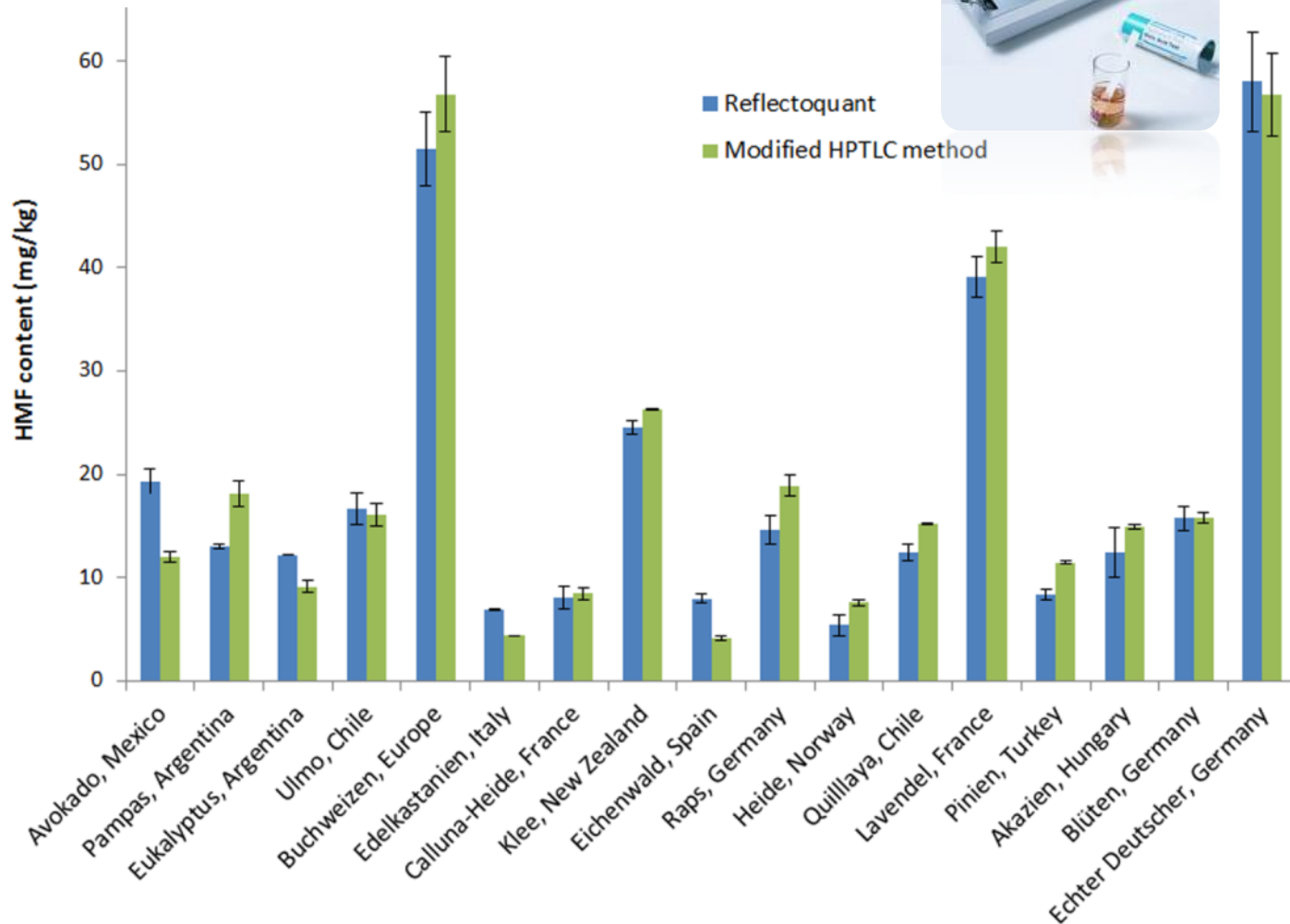


Sample#	HPTLC-UV			HPTLC-ESI-MS, SIM		Deviation UV <i>versus</i> MS HMF in honey, mg/kg	
	Plate 1	Mean HMF in honey, mg/kg	%RSD (n = 3)	Reprod. (%RSD, n=2)	Mean HMF in honey, mg/kg		%RSD (n = 3)
1		37.7	3.7	4.4	36.4	18	1.3
2		50.1	2.8	2.3	57.1	10	-7.0
3		78.4	1.5	1.9	72.0	2.5	6.4
4		35.0	2.0	3.3	29.2	7.5	5.8
Mean				3.0			5.1

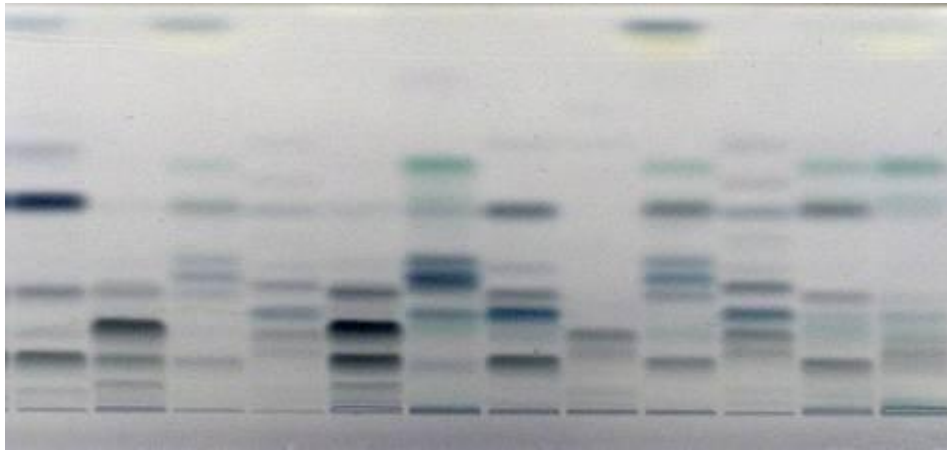


# Method comparison

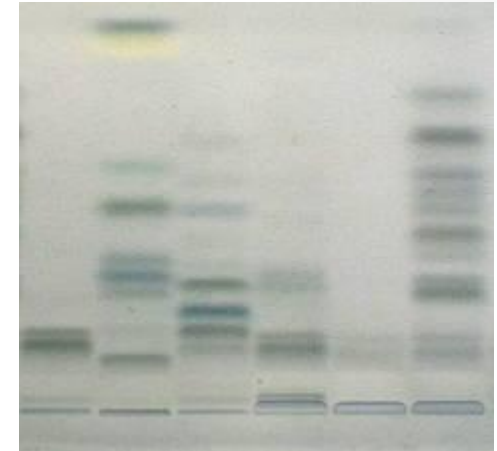
→ Fast methods: HPTLC *versus* Reflectoquant



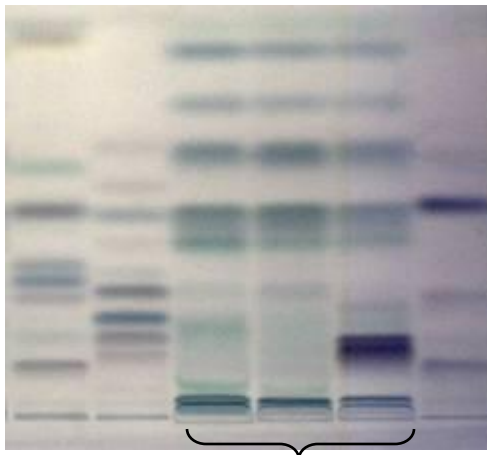
# Analysis of biopolymers → monomeric units



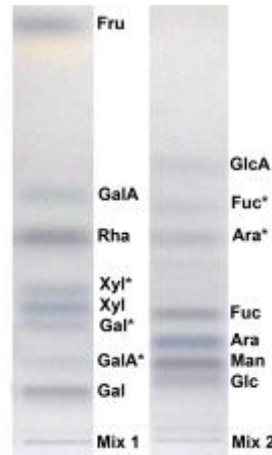
Carr JBKM Mix 1 Mix 2 Guar Trag Gum Xan Mix 1 Mix 2 Kara Pect



Starch Mix 1 Mix 2 CMC Cell HPMC



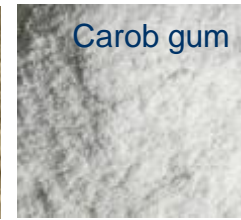
Mix 1 Mix 2 Alginate Agar



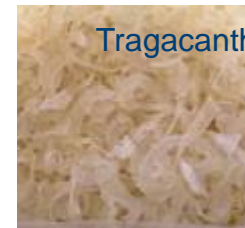
Pectin



Agar-Agar



Carob gum



Tragacanth



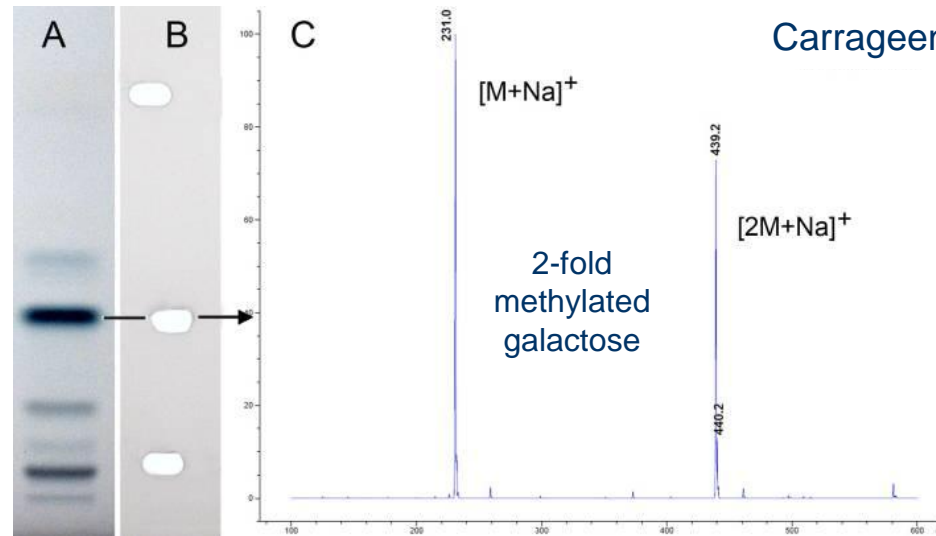
Gummi arabicum



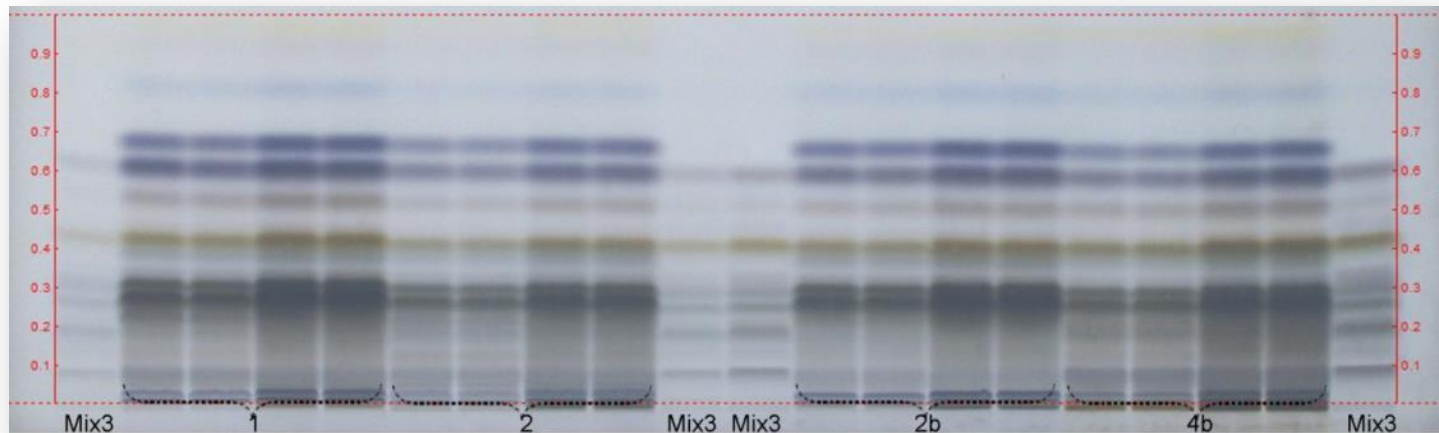
Carrageen



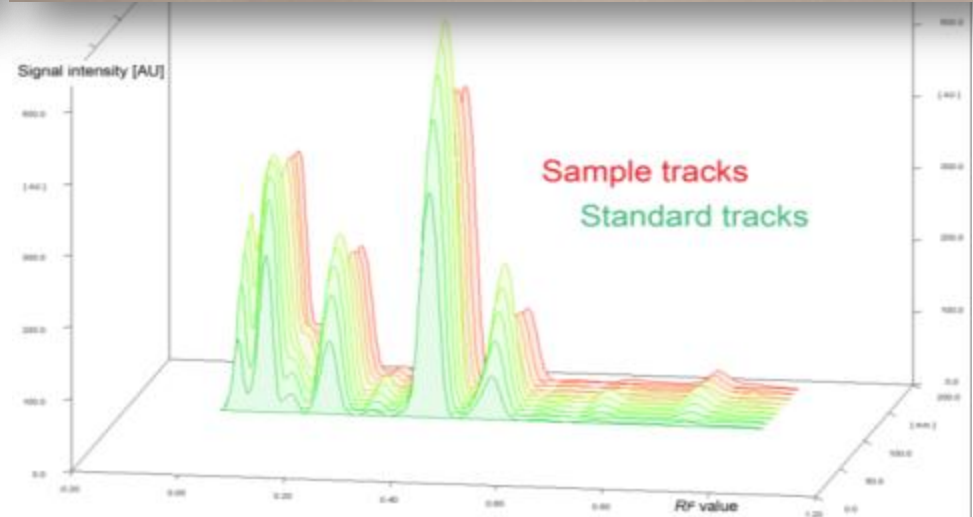
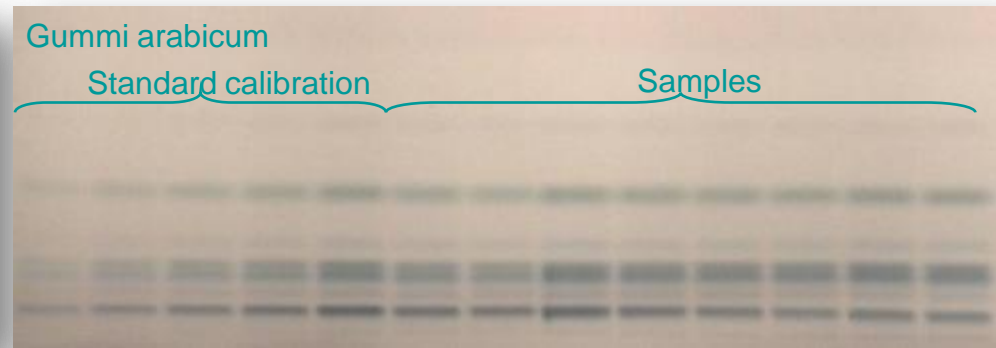
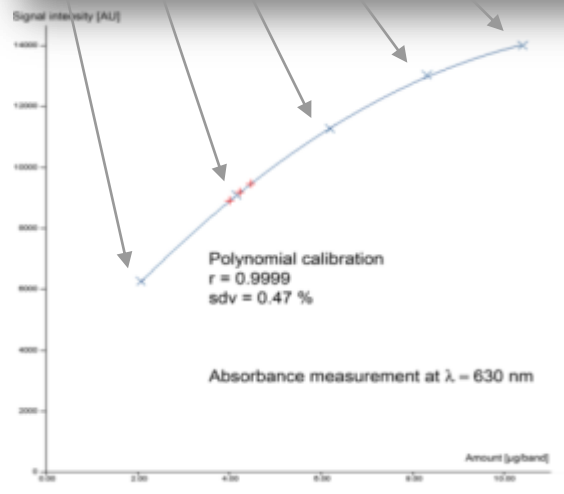
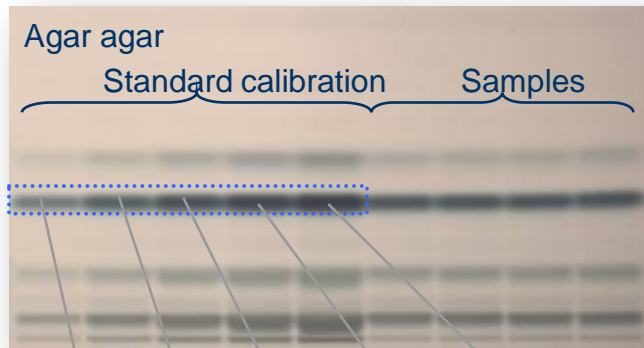
# Analysis of biopolymers → monomeric units



## *Ocimum basilicum*

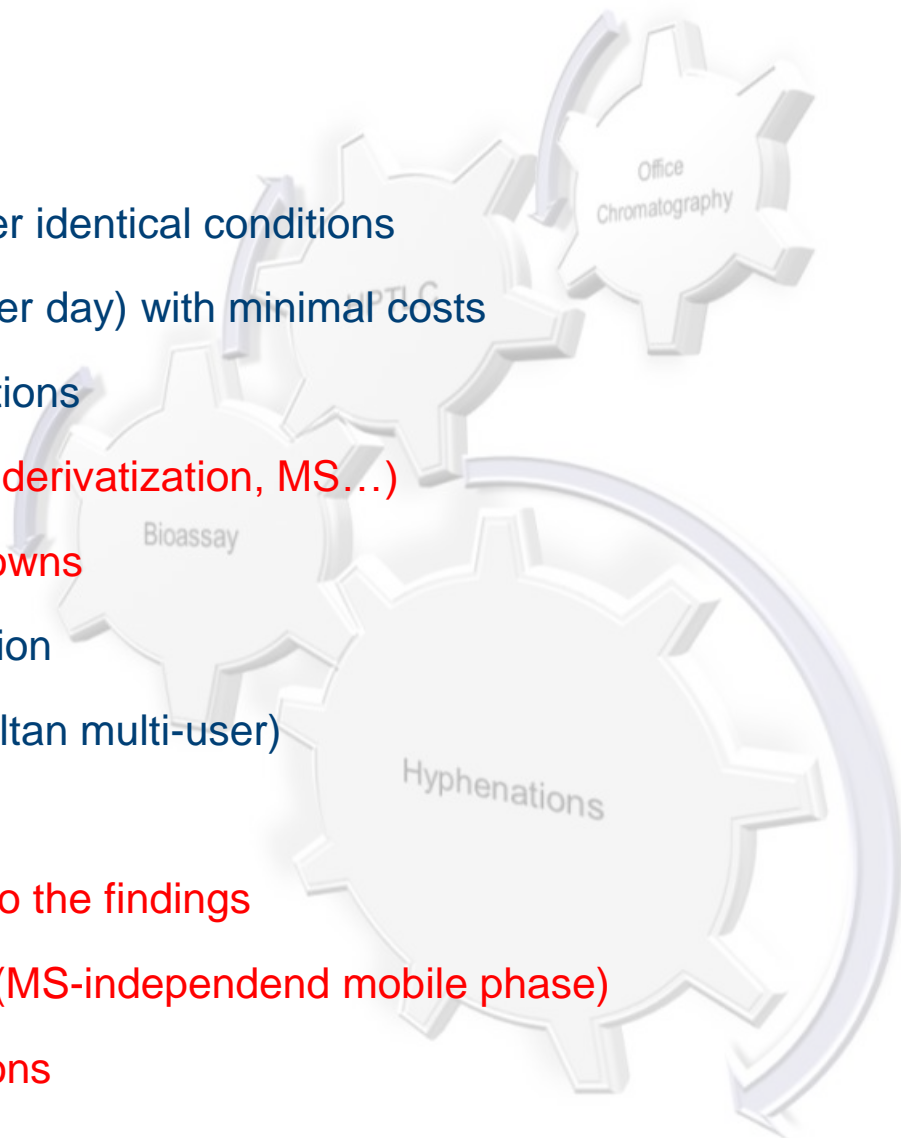


# Quantitation

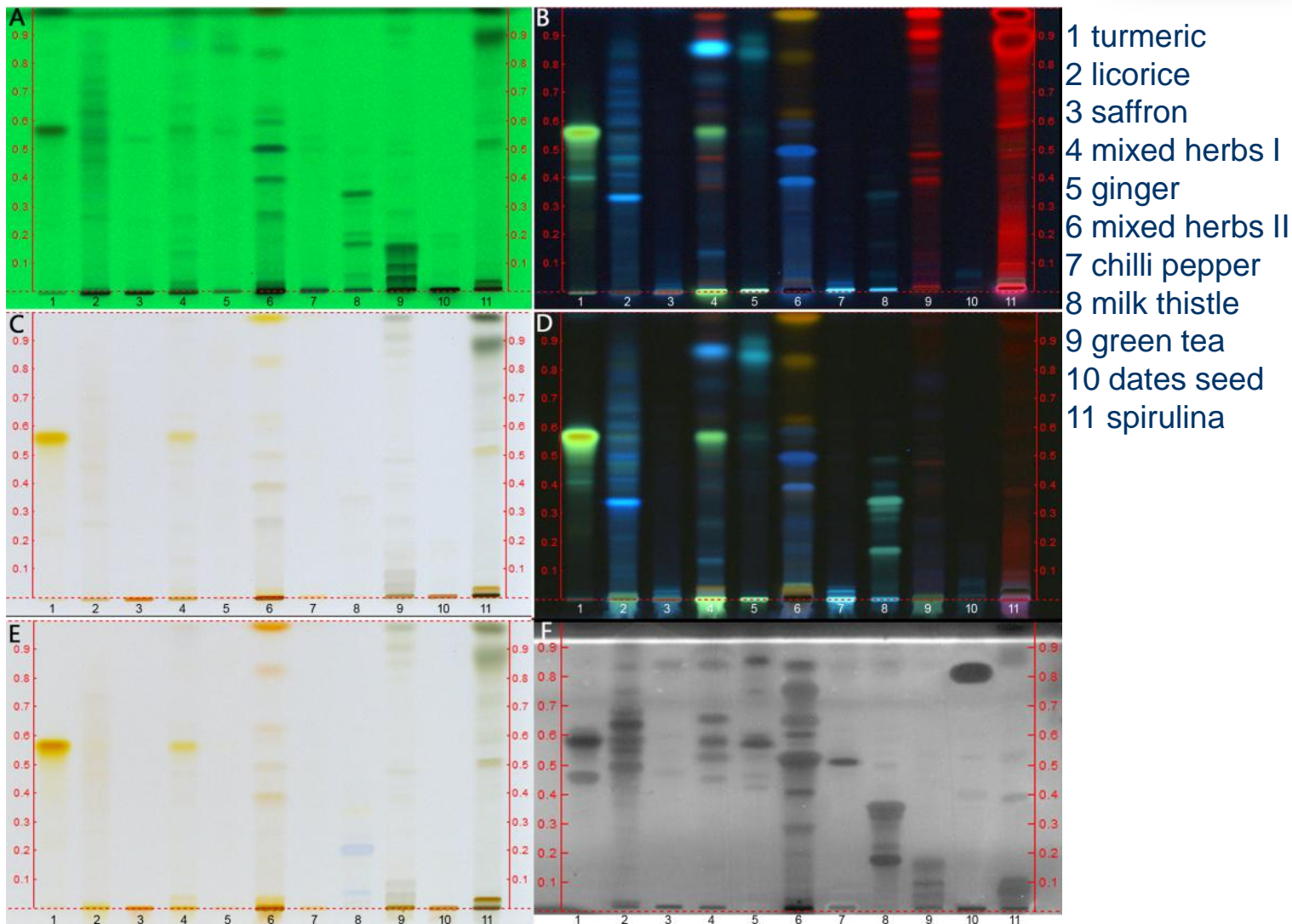


# The power of HPTLC

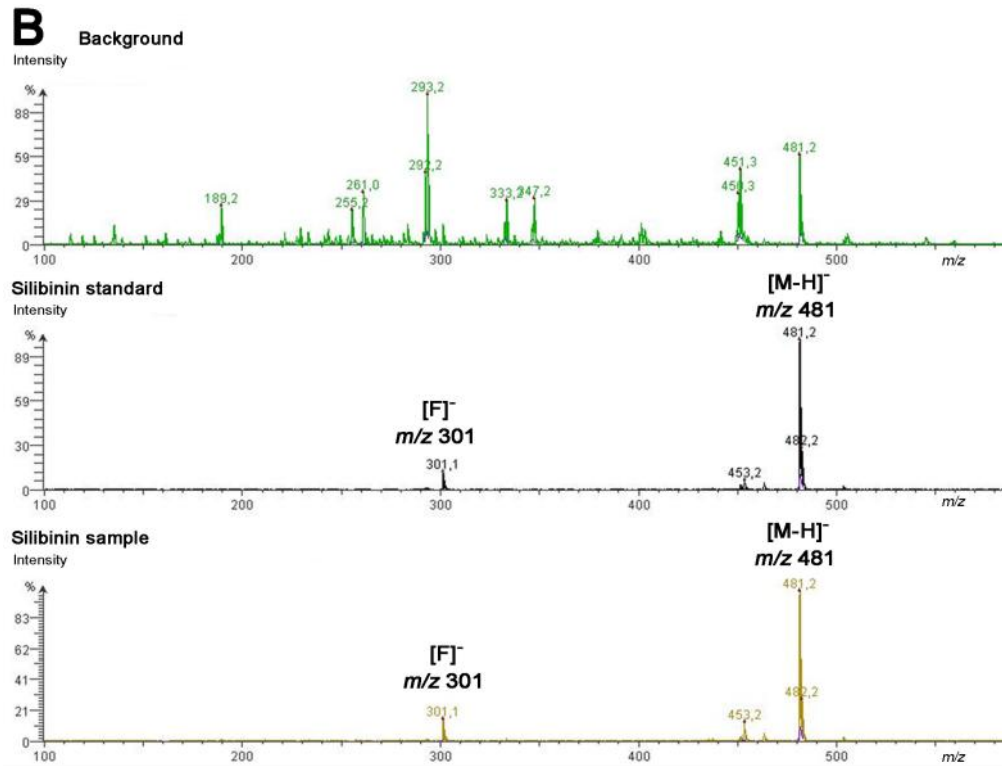
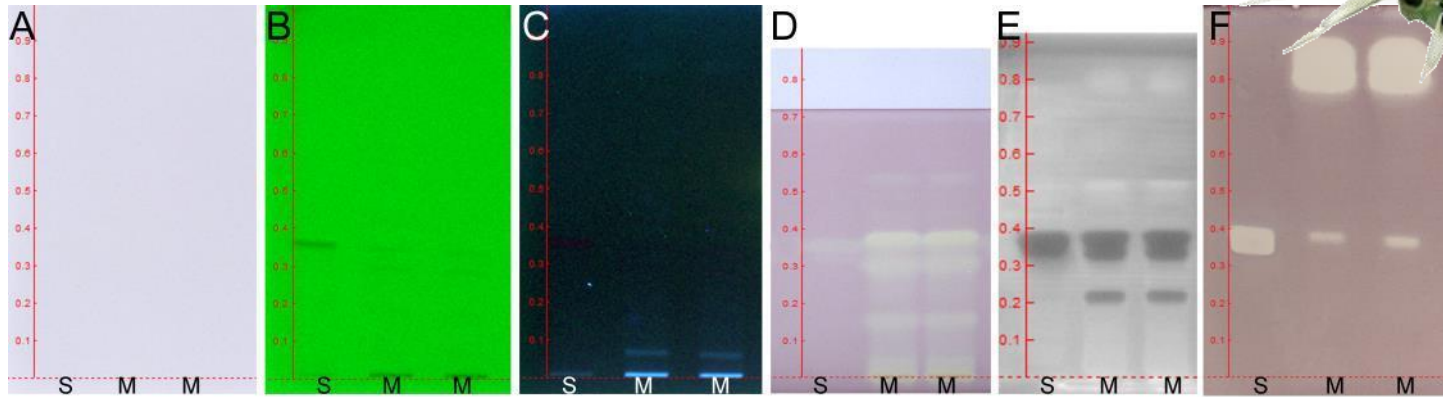
1. Reduced sample preparation
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3. Parallel chromatography under identical conditions
4. High throughput (1000 runs per day) with minimal costs
5. Selective, simultan derivatizations
6. Multi-detection (UV/Vis, FLD, derivatization, MS...)
7. More information about unknowns
8. Concentration during application
9. Flexible working station (simultan multi-user)
10. Effect-directed analysis
11. Analytical workflow adjusted to the findings
12. Targeted mass spectrometry (MS-independend mobile phase)
13. The ease of super-hyphenations



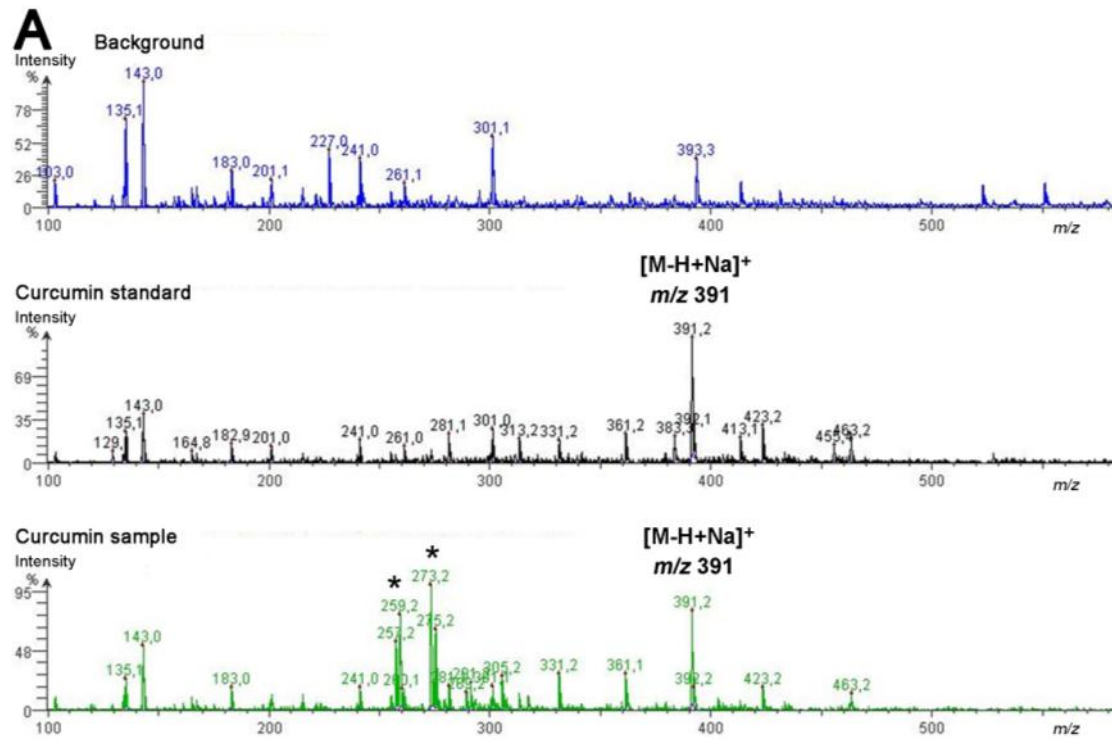
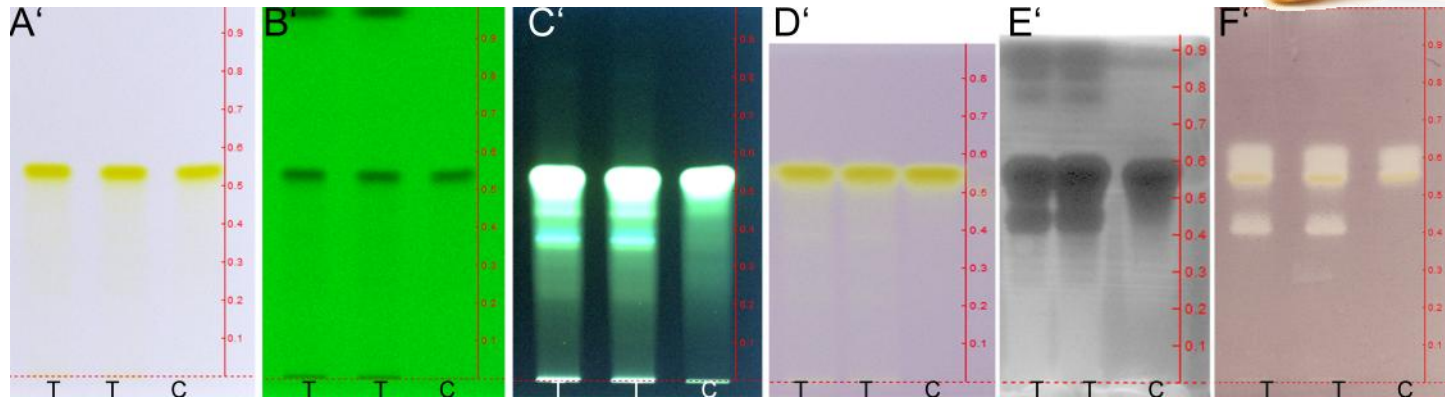
# EDA of 11 plant species



# EDA of milk thistle (*Silybum marianum*) seeds

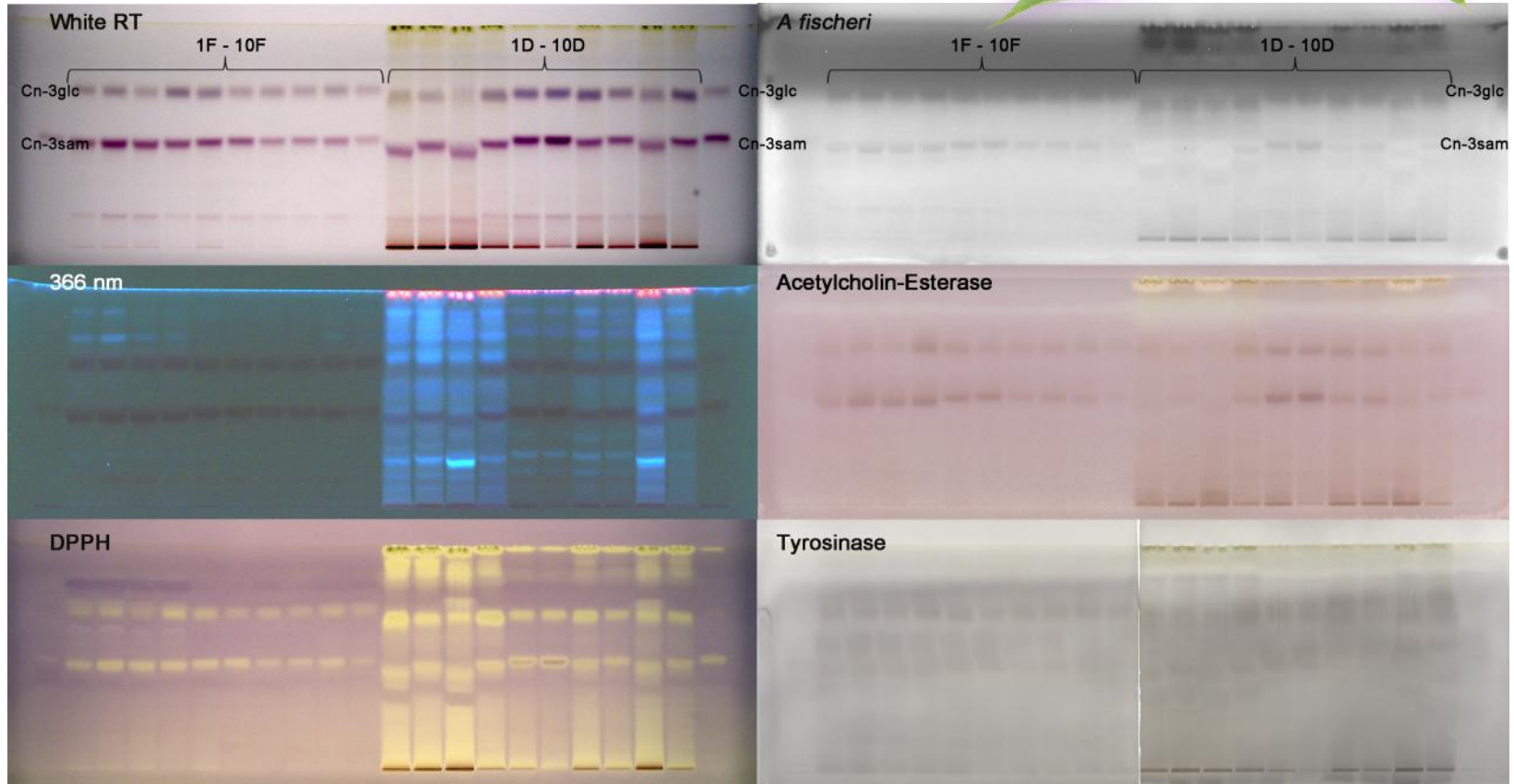


# EDA of turmeric (*Curcuma longa*) rhizomes

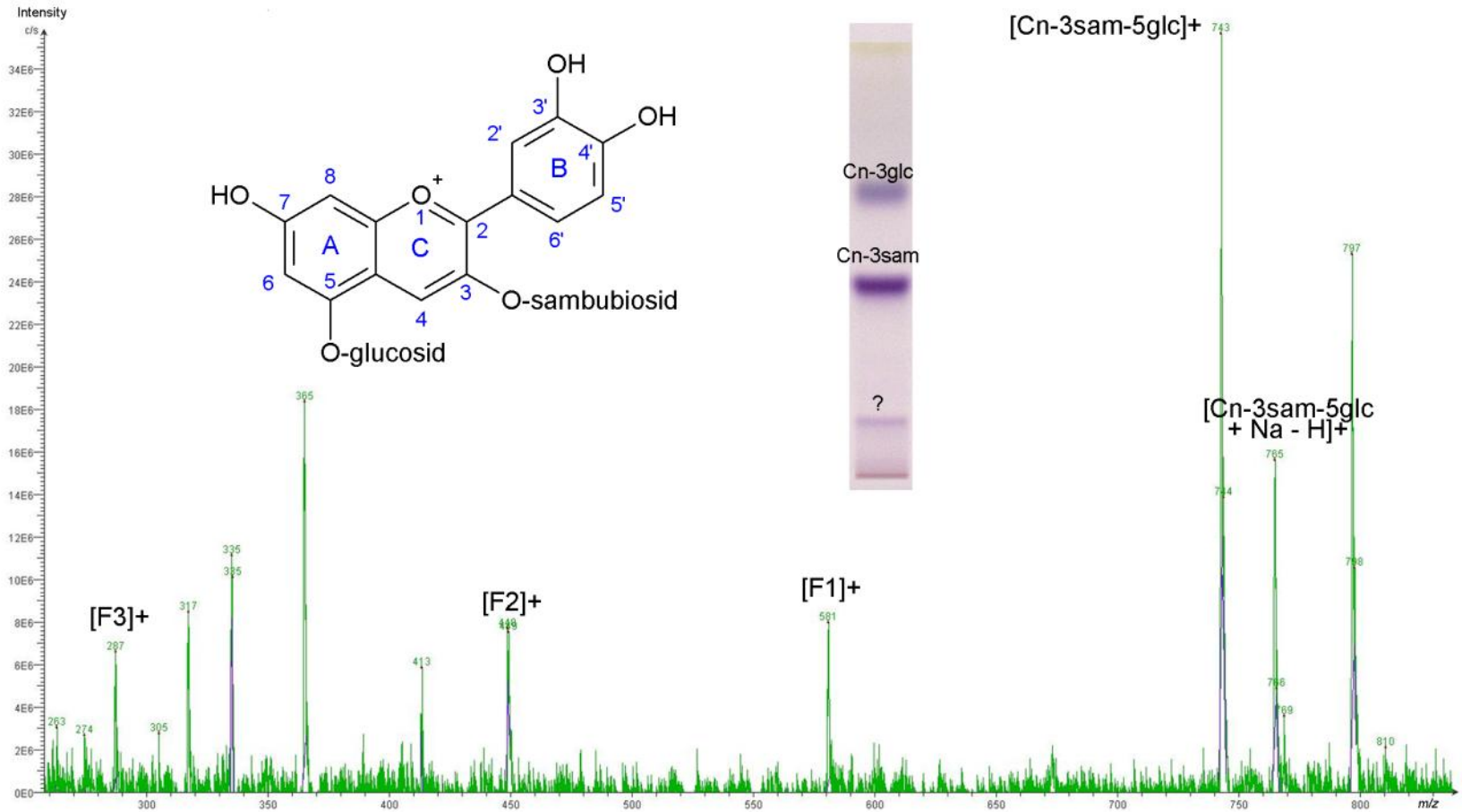


# EDA of elderberry (*Sambucus nigra* L.)

→ fresh *versus* dried fruit samples

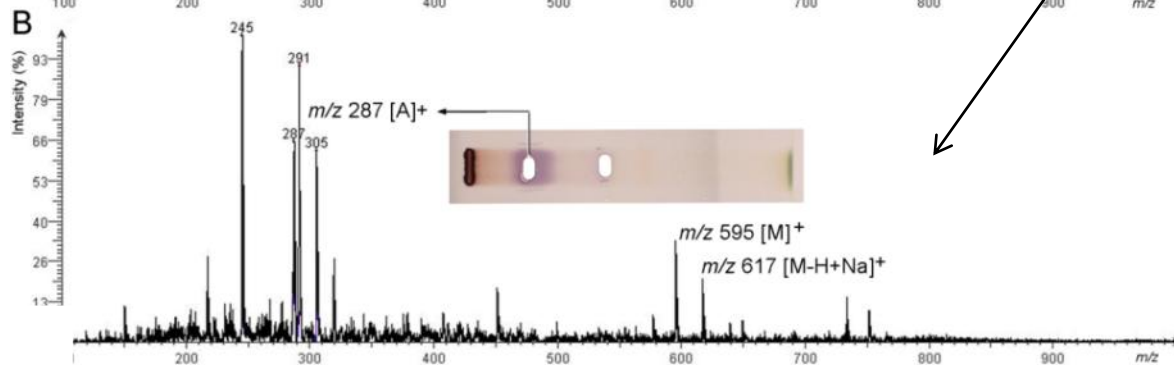
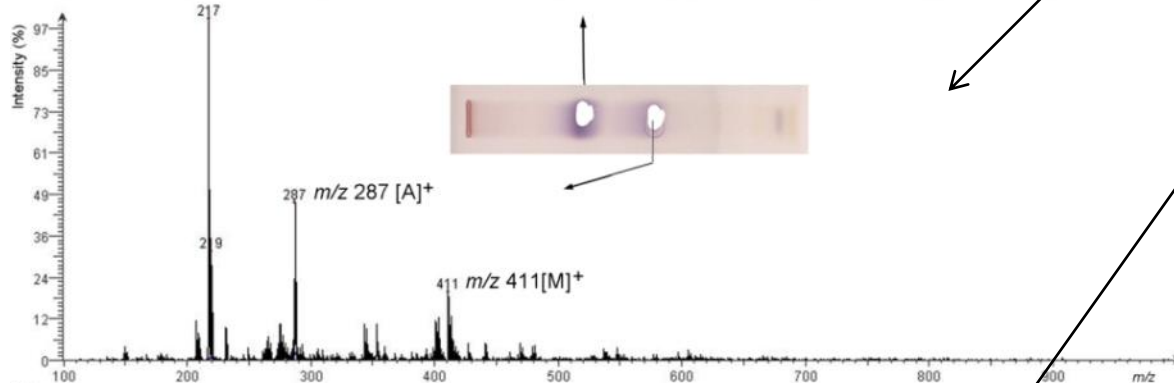
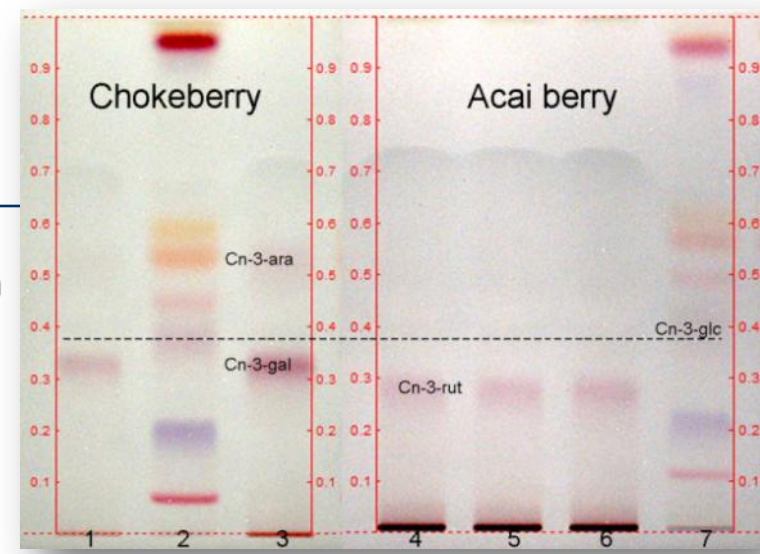
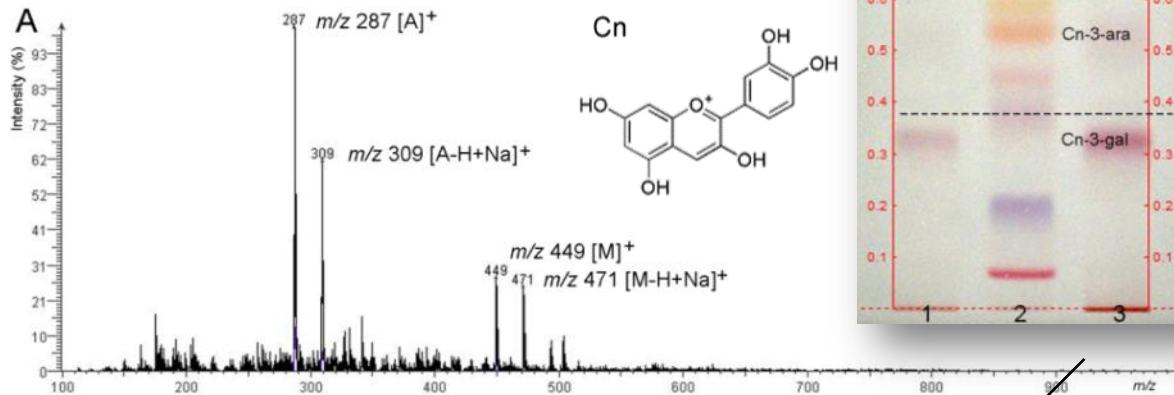


# HPTLC-ESI<sup>+</sup>-MS of unknown zone at $hR_F$ 10





# Analysis of anthocyanes

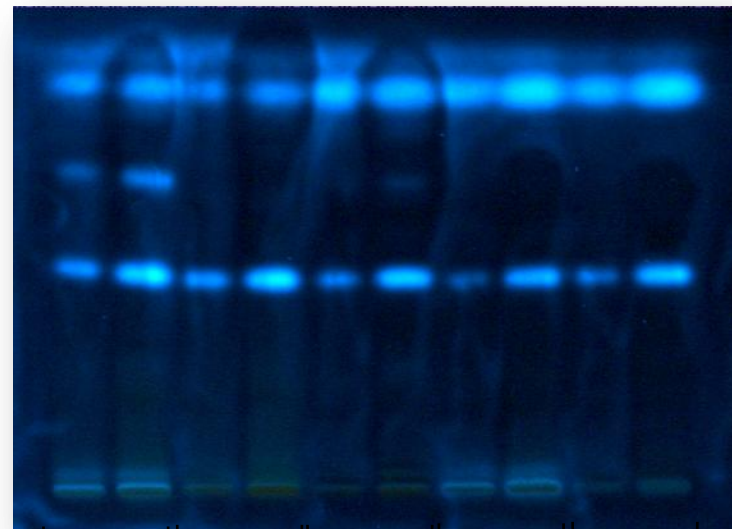


# EDA of estrogen-effective compounds

→ Spices



→ Beer samples

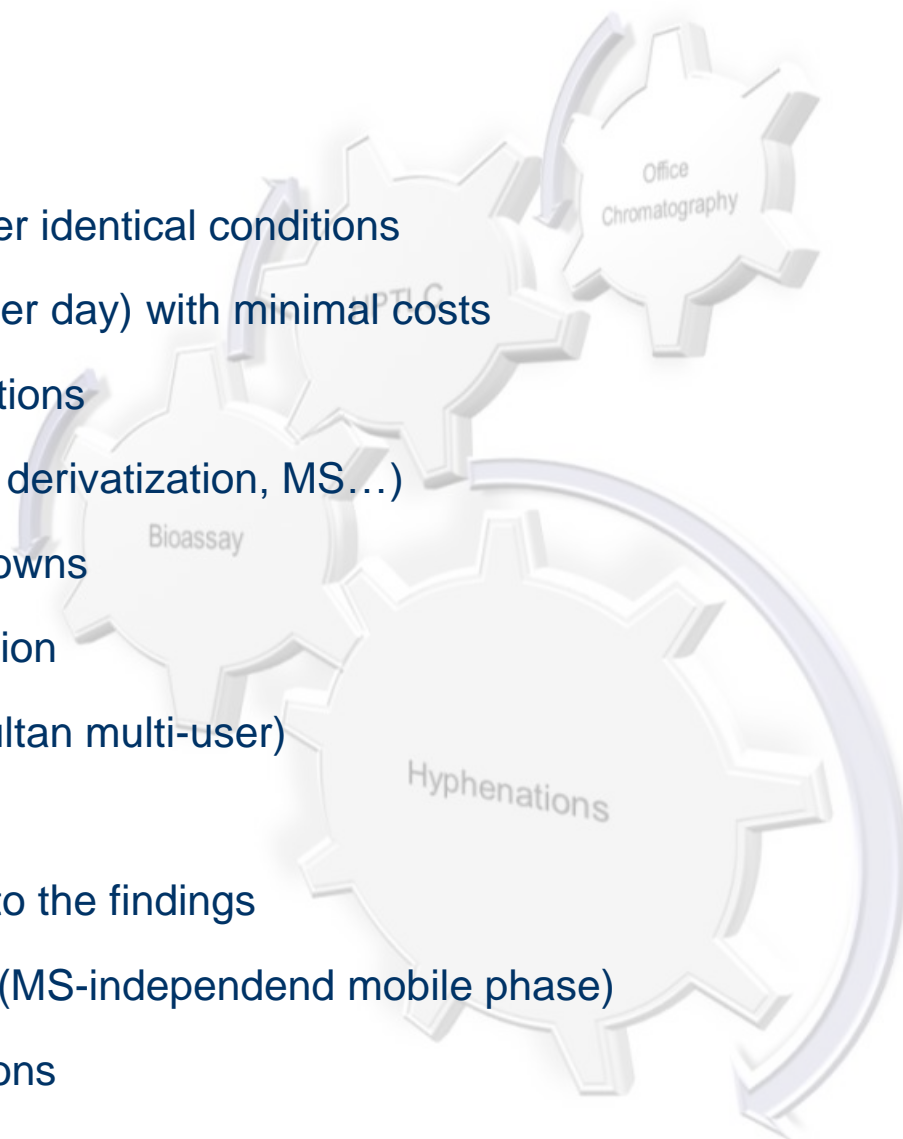


Köstritzer Guinness Licher Porter Karlskrone



# The power of HPTLC

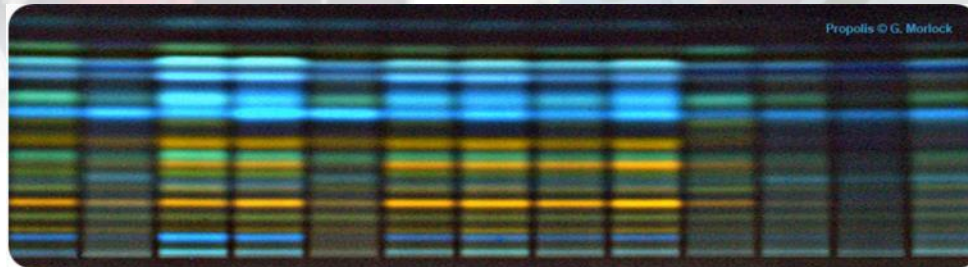
1. Reduced sample preparation
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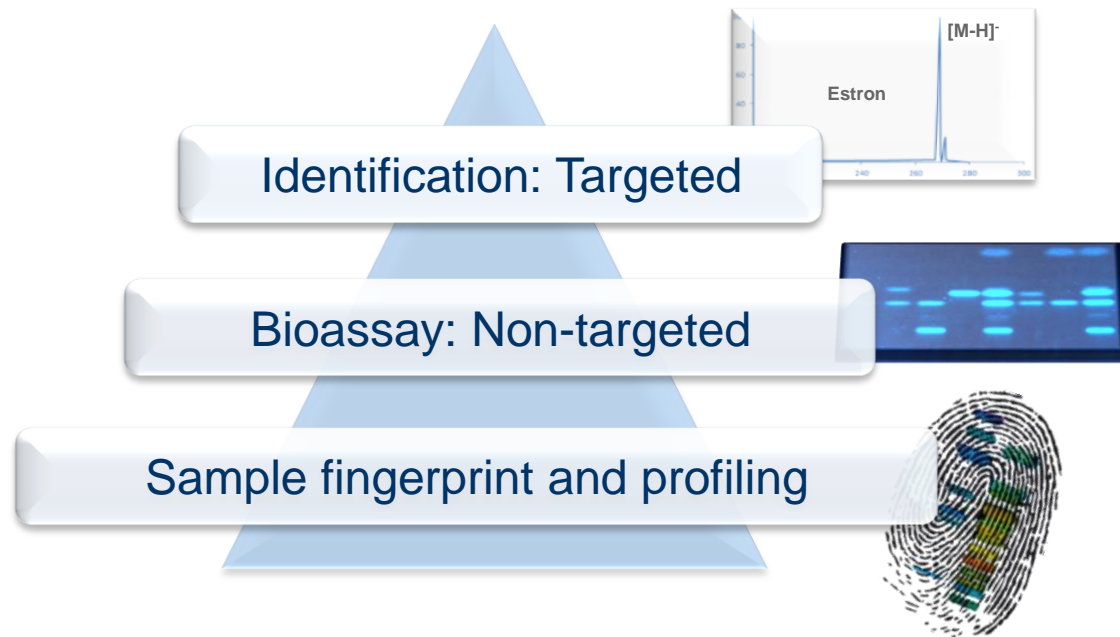


Thank you!

JUSTUS-LIEBIG-  
UNIVERSITÄT  
GIESSEN  
Food Science 

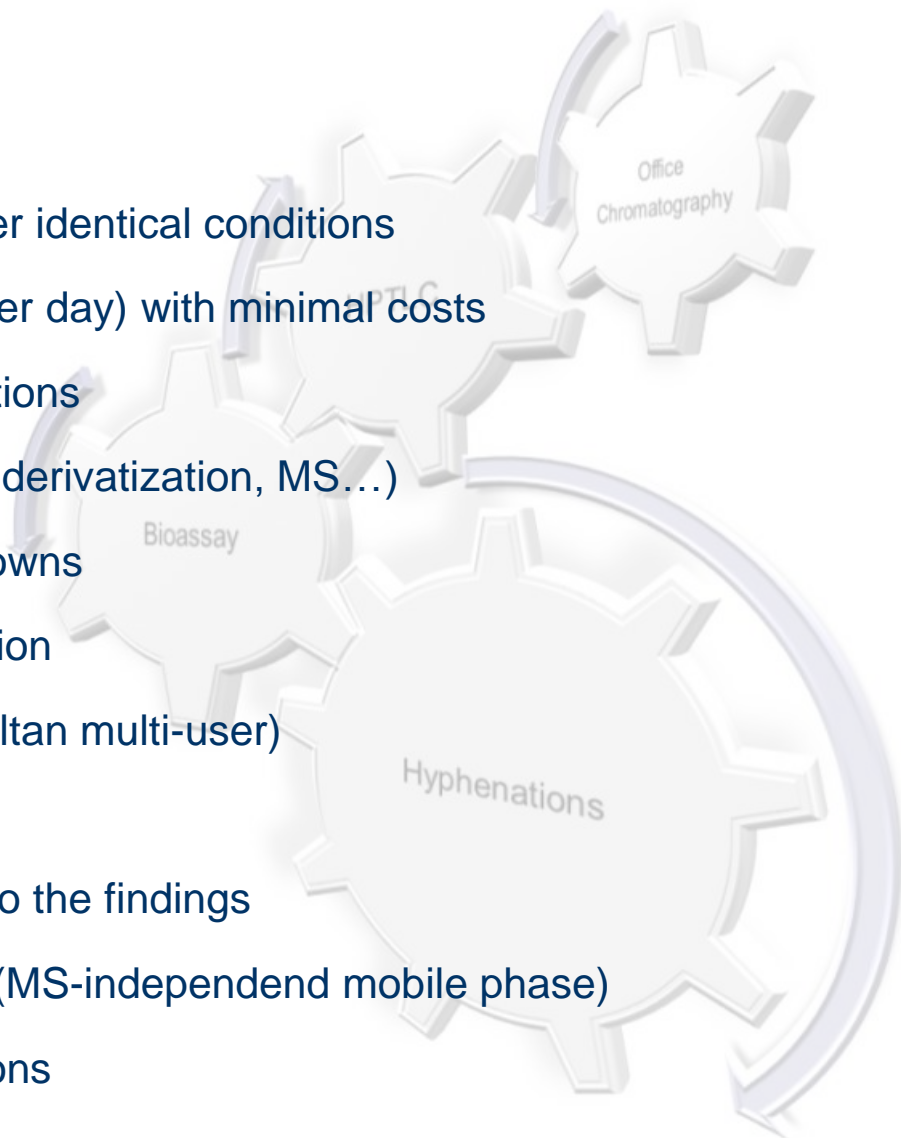


# Breakthrough in the hyphenation HPTLC-EDA-MS – a must-have method in the analyst's toolbox



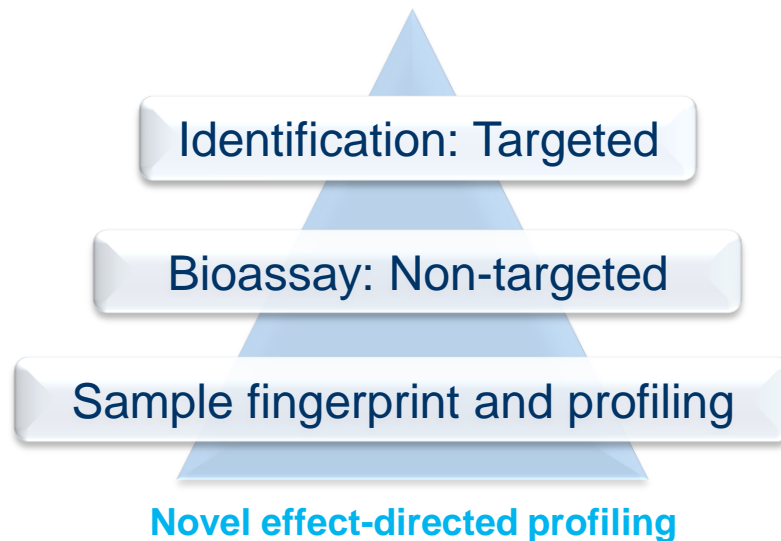
# The power of HPTLC

1. Reduced sample preparation
2. Matrix-tolerant method
3. Parallel chromatography under identical conditions
4. High throughput (1000 runs per day) with minimal costs
5. Selective, simultan derivatizations
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8. Concentration during application
9. Flexible working station (simultan multi-user)
10. **Effect-directed analysis**
11. Analytical workflow adjusted to the findings
12. Targeted mass spectrometry (MS-independend mobile phase)
13. The ease of super-hyphenations

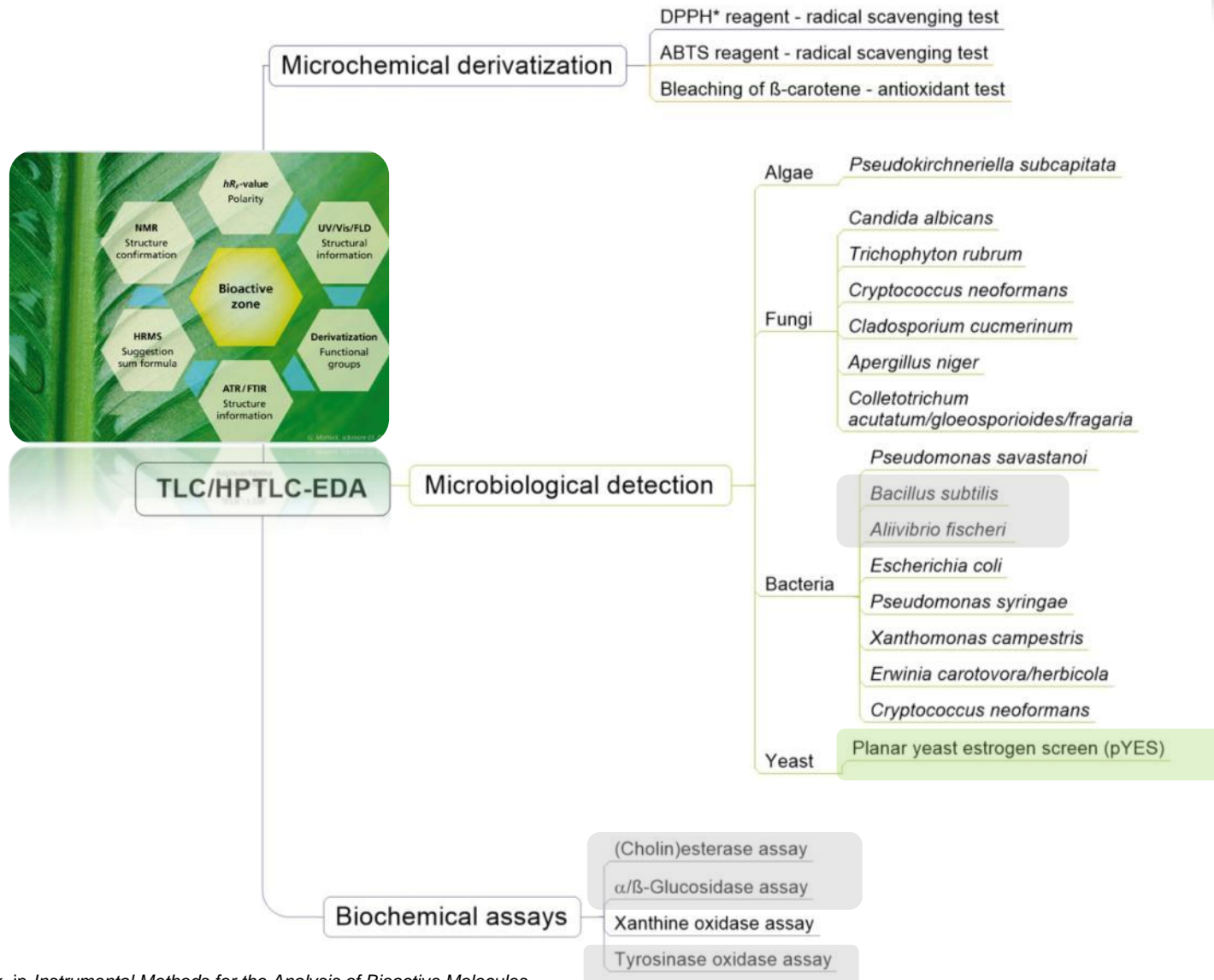


# Impact

- Only a part is represented by target analysis or multi-methods
- Complex samples contain thousands of compounds
- X It remains largely unclear which peaks are bioactive
- X Samples to be identified as risky or bioactive



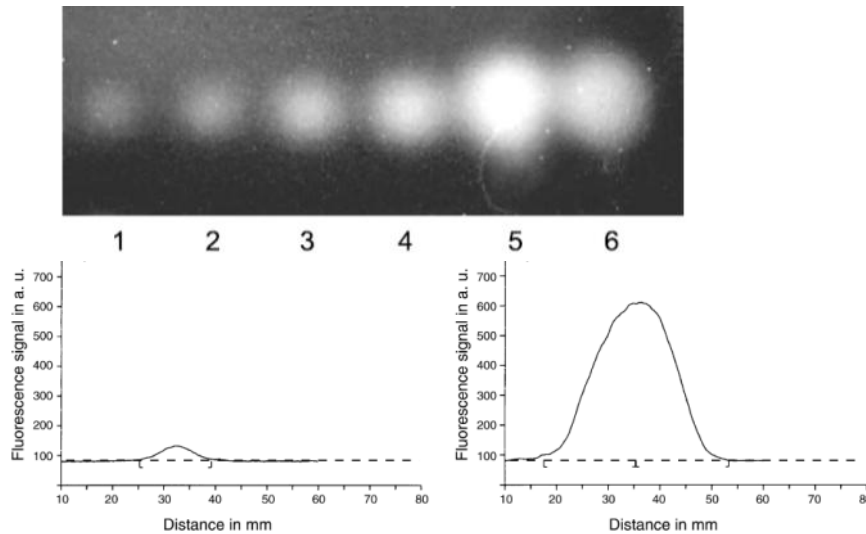
# Effect-directed link to the compound



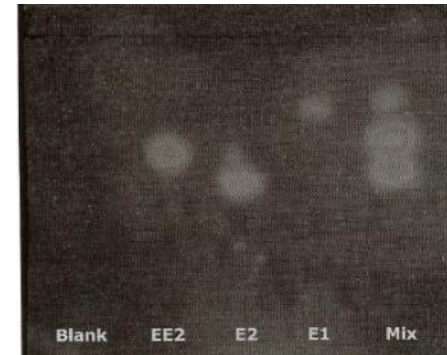


# Challenge: Avoid zone broadening

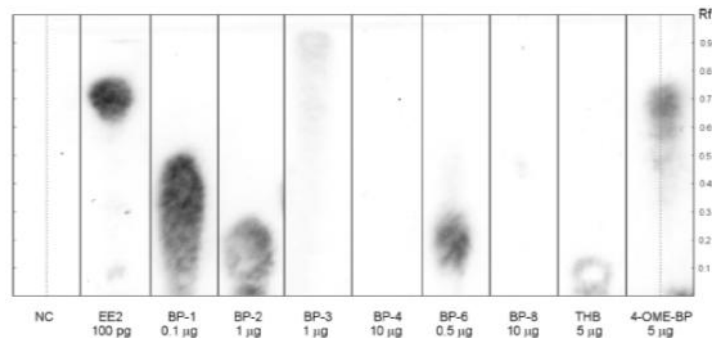
Goodall & Levi, Nature 158, 675–676 (1946)



Mueller *et al.* *Chromatographia* 60, 207-211 (2004)

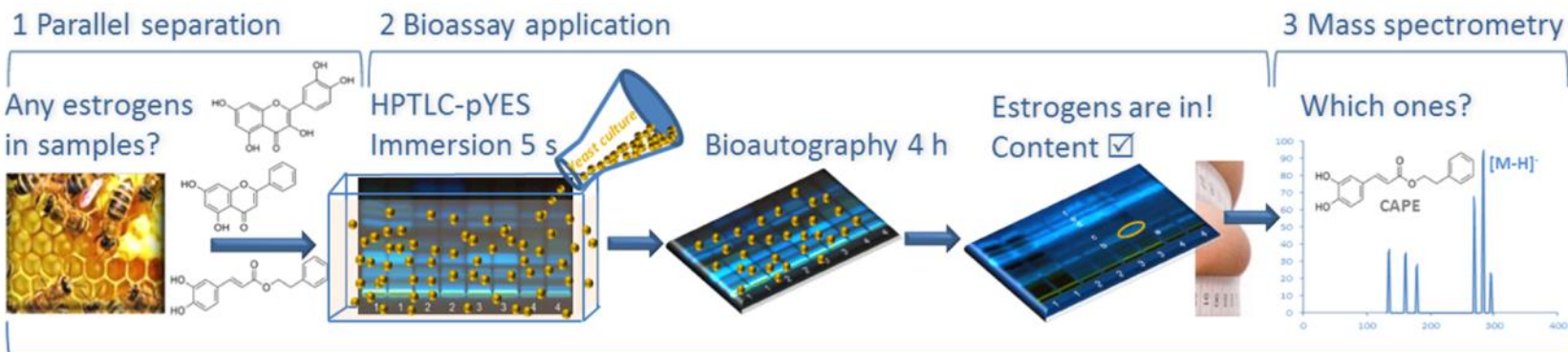


Schönborn & Grimmer,  
*J. Planar Chromatogr.* 26, 402-408 (2013)

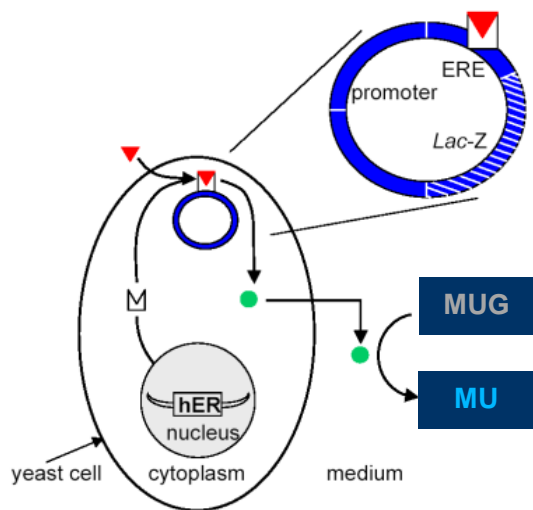


Buchinger *et al.*  
*Anal. Chem.* 85, 7248-7256 (2013)

# Detection of endocrine disrupting compounds (EDCs)



→ LC-bioassay-MS workflow for 20 samples in parallel within 5 h (15 min per sample)



Modified from draft of pYES expert group

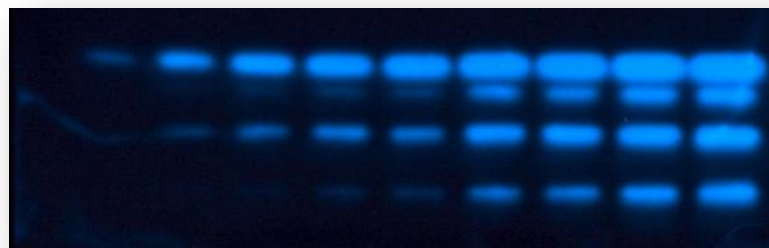
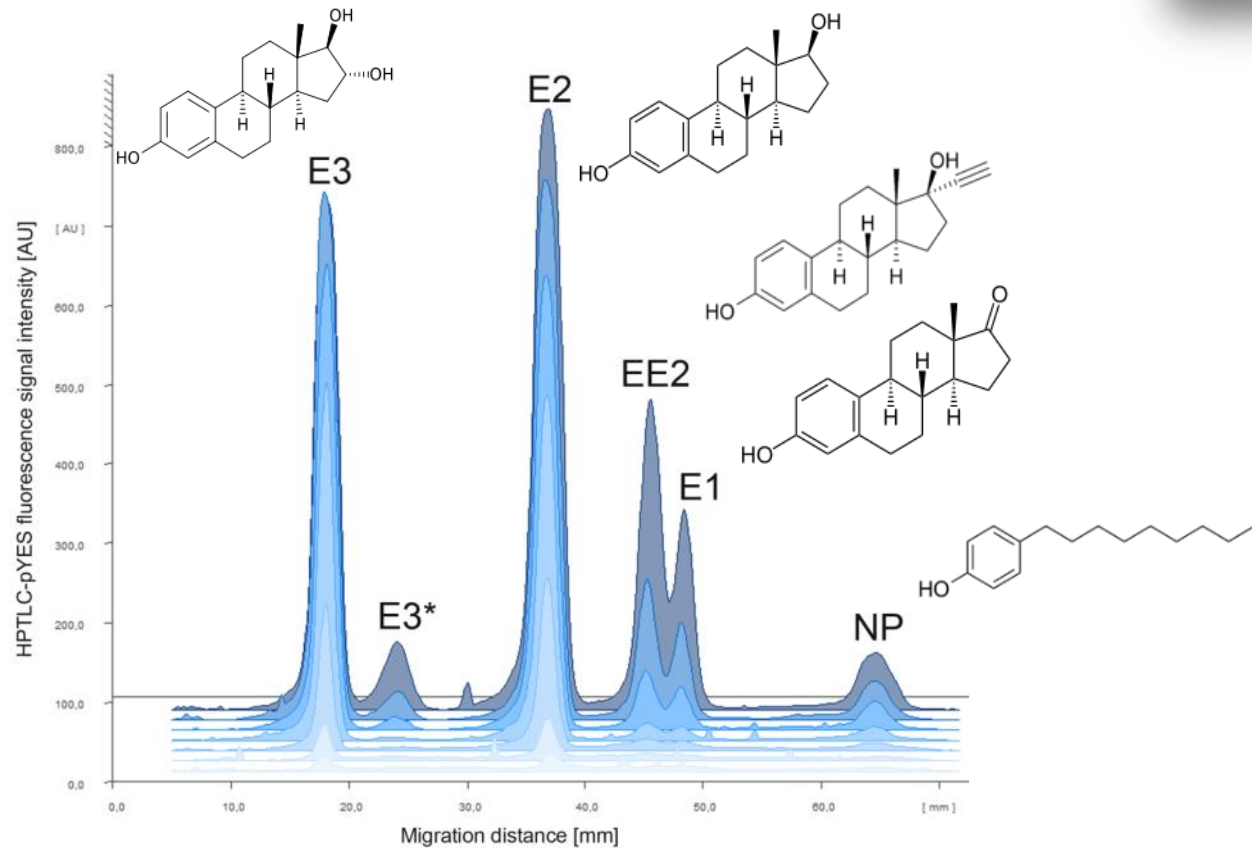
## Planar yeast estrogen screen (pYES)

- using human estrogen receptor hER $\alpha$
- in *Saccharomyces cerevisiae*

1. Routledge & Sumpter, Environ. Toxicol. Chem. 15 (1996) 241
2. McDonnell *et al.*, J. Steroid Biochem. Mol. Biol. 39 (1991) 291

→ blue fluorescent 4-methylumbelliferone

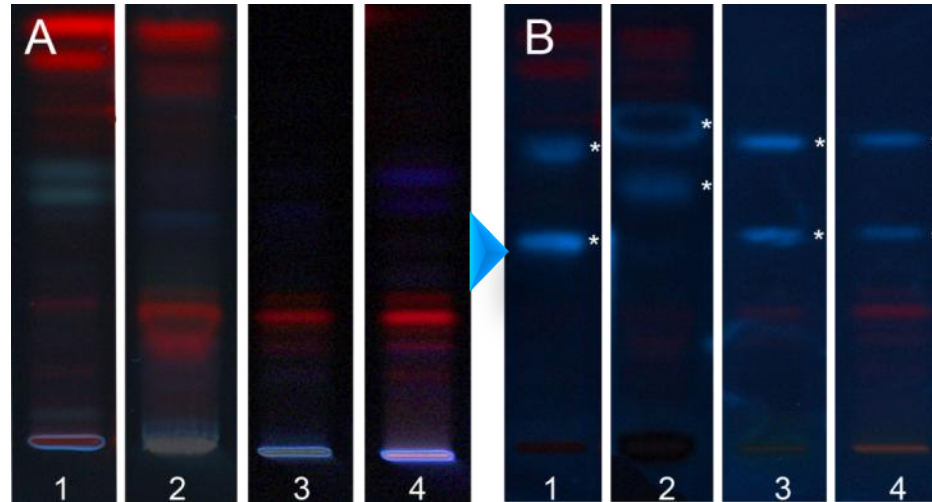
# Detection of EDCs



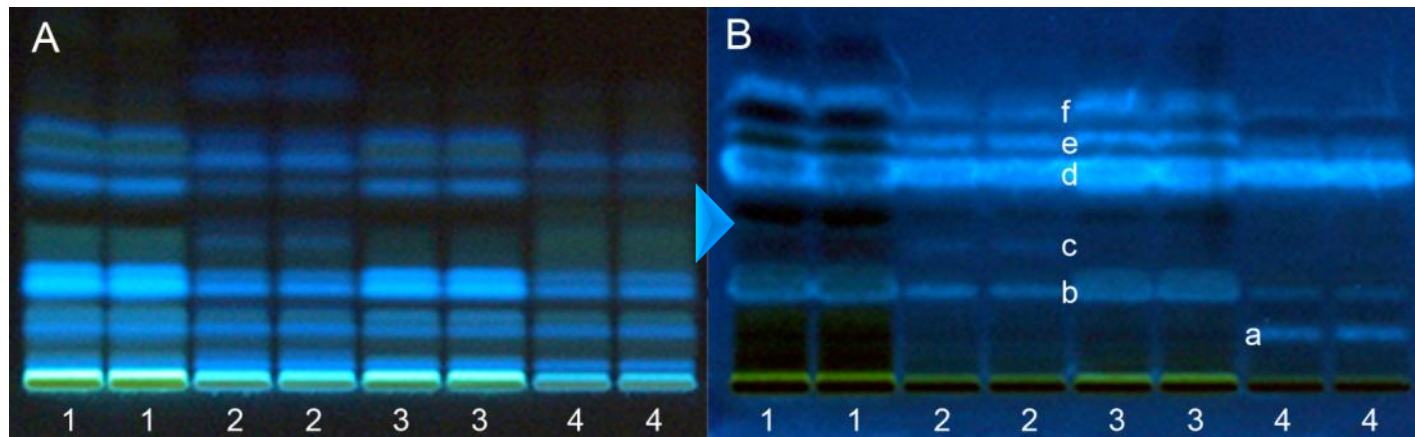
	pg/band
E1	25 – 1000
EE2	0.5 – 20
E2	0.5 – 20
E3	25 – 1000

# Detection of EDCs

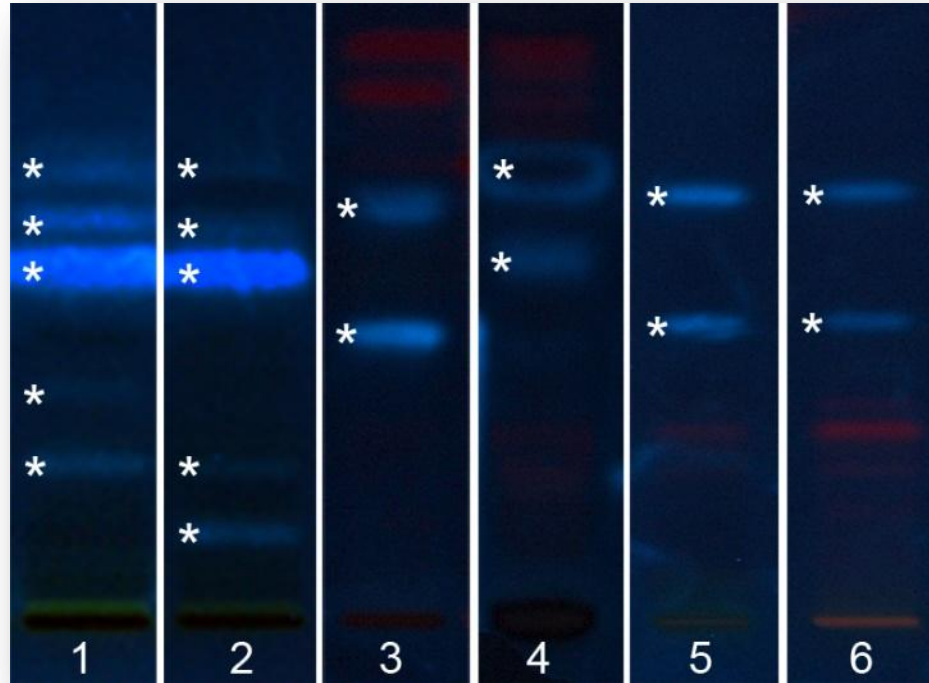
→ Spices/tea



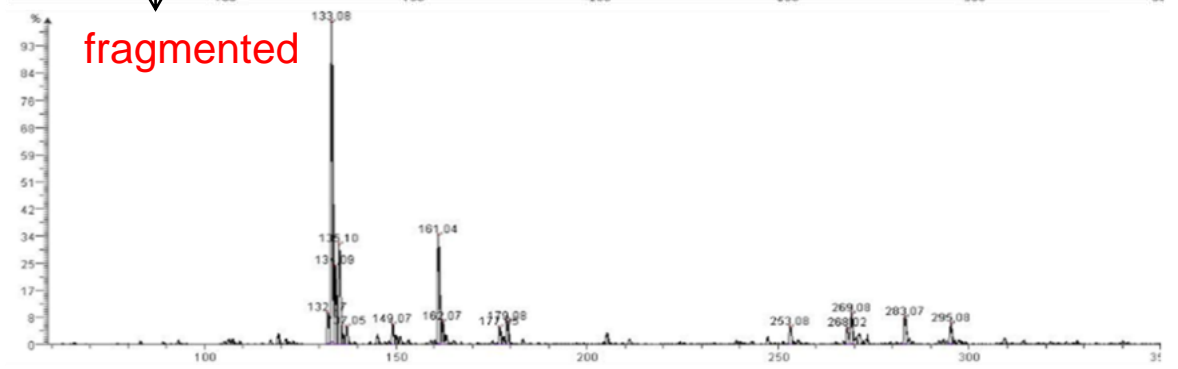
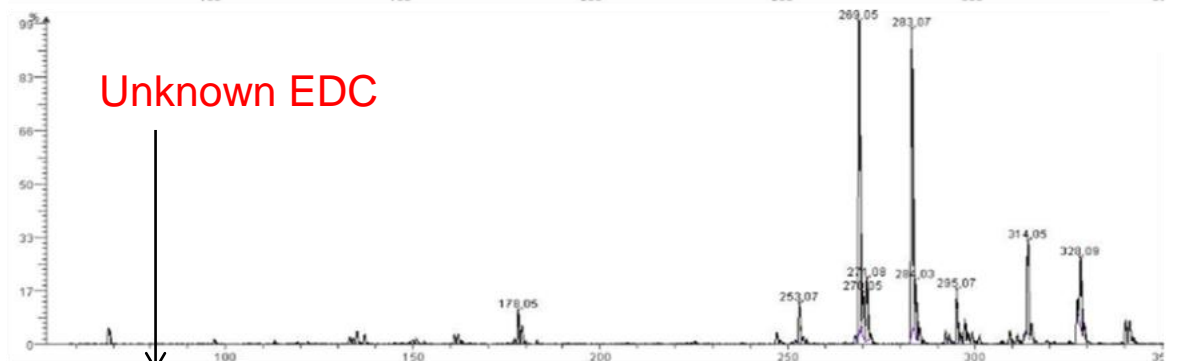
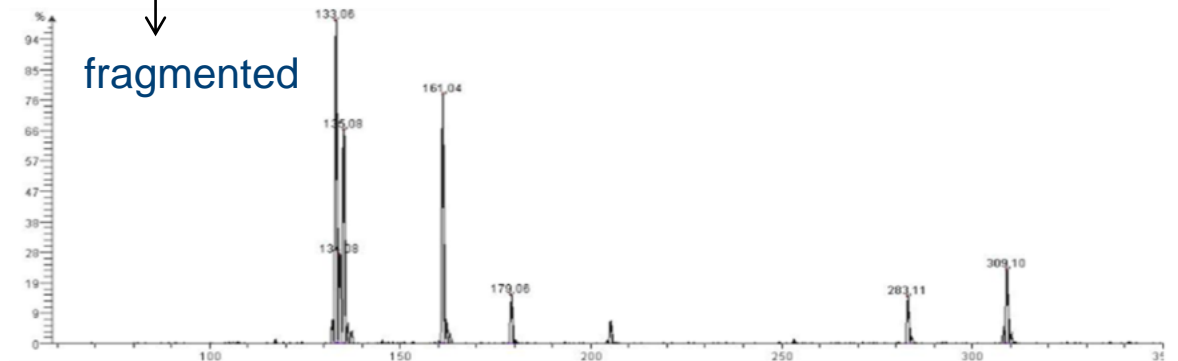
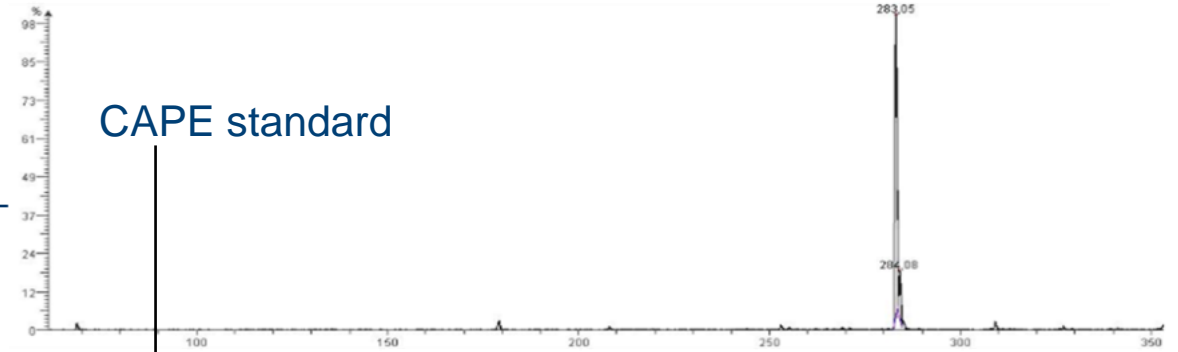
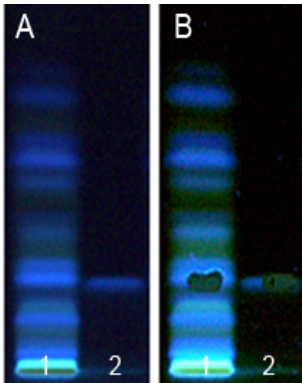
→ Propolis samples



# What is it?



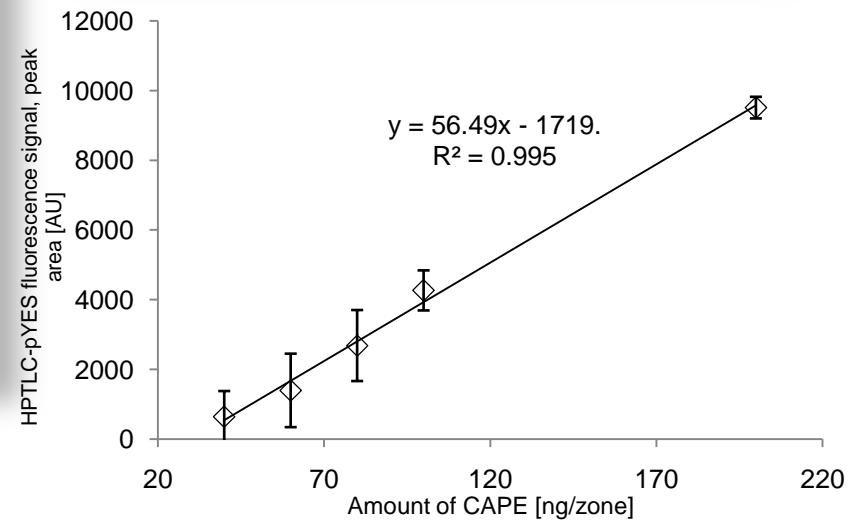
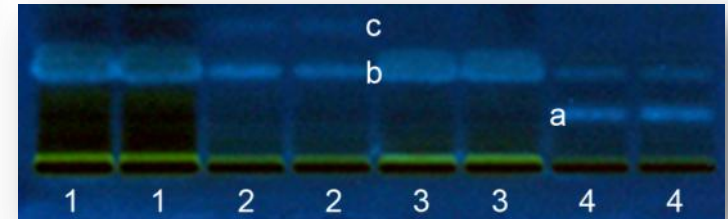
# CAPE?



# Quantitation of CAPE in propolis

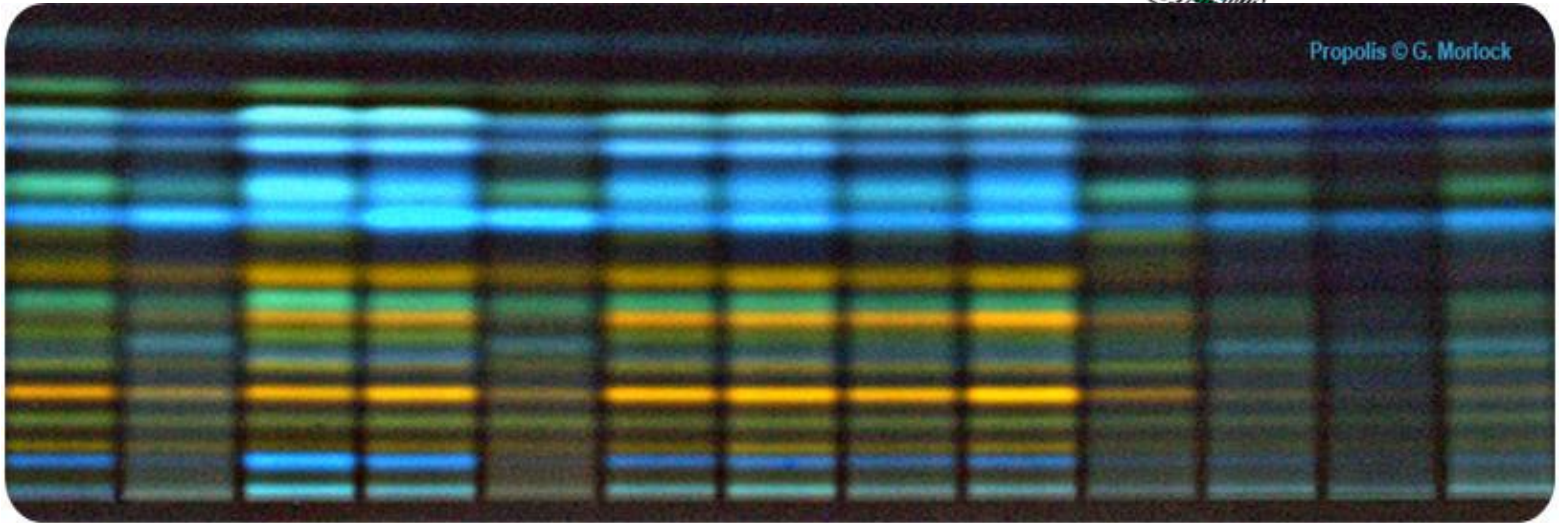
Propolis sample	CAPE content in sample [ $\mu\text{g/mL}$ ]	CAPE content [ $\mu\text{g/g}$ ] referred to propolis weight (n=2)
P1 (30 %)	481	2028
P2 (30 %)	476	2009
P3 (25 %)	471	2387
P4 (62 %)	348	710
P5 (not specified)	380	380 <sup>3</sup>
P6 (250 mg/capsule)	359 <sup>1</sup>	1435
P7 (30 mg/lozenge)	22 <sup>2</sup>	1089

<sup>1</sup> $\mu\text{g/capsule}$ , <sup>2</sup> $\mu\text{g/pastille}$ , <sup>3</sup> $\mu\text{g/mL}$



# Fingerprint of phenolic compounds in propolis

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

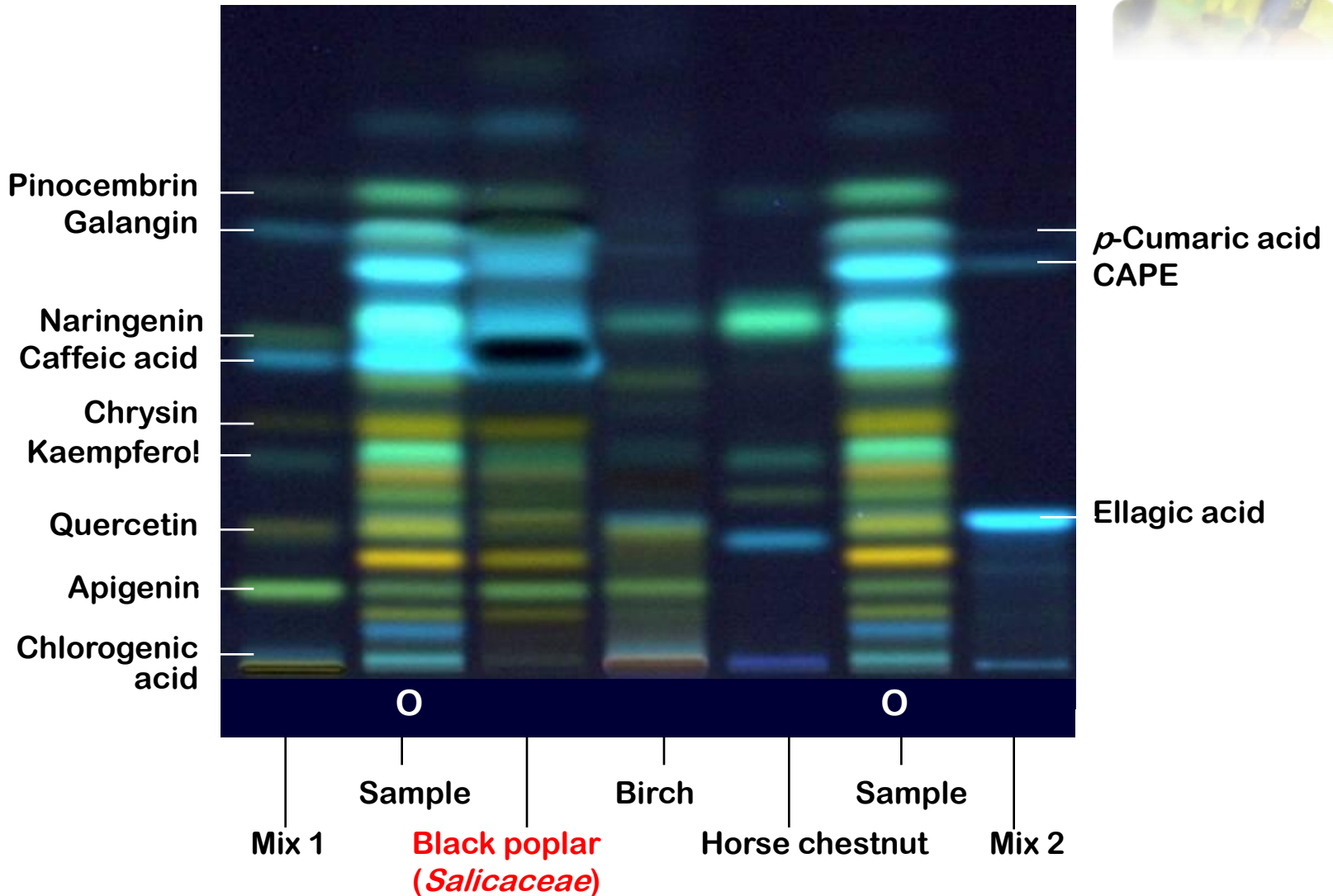


G. Morlock *et al.*, in preparation

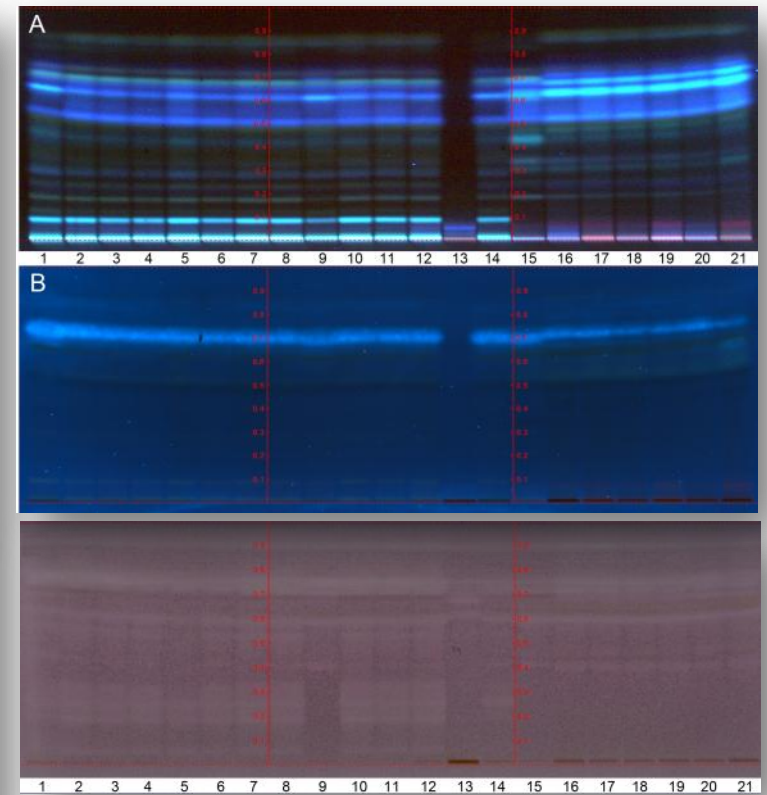
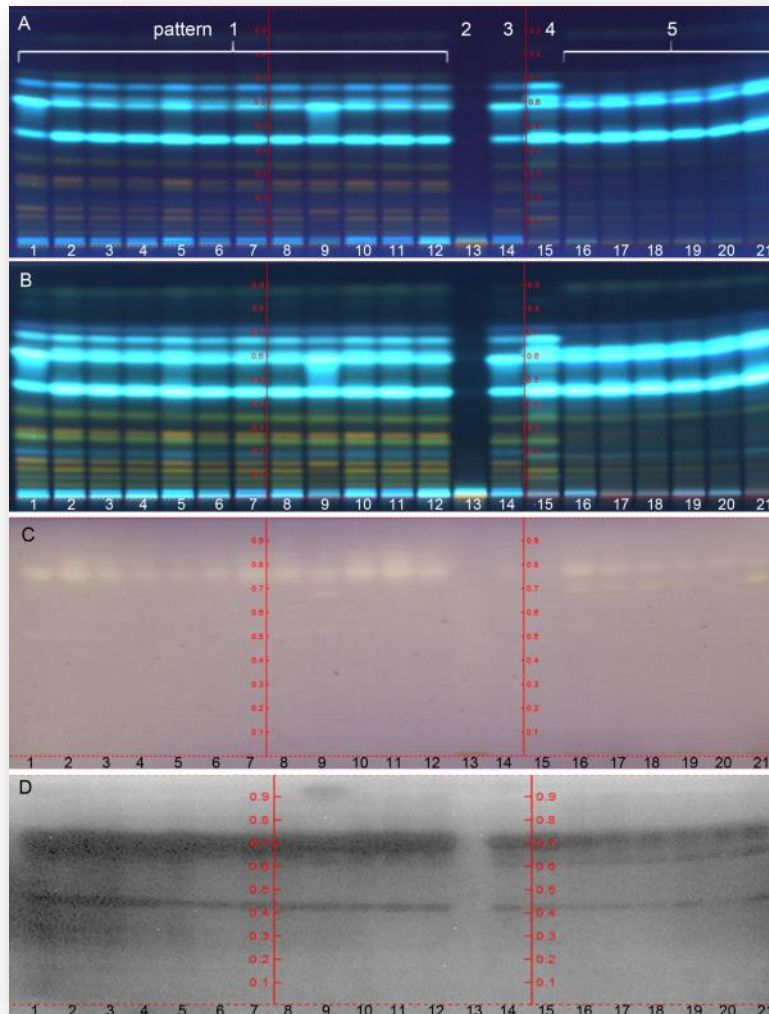
Cooperation with Wala and Apicultural State Institute, Stuttgart



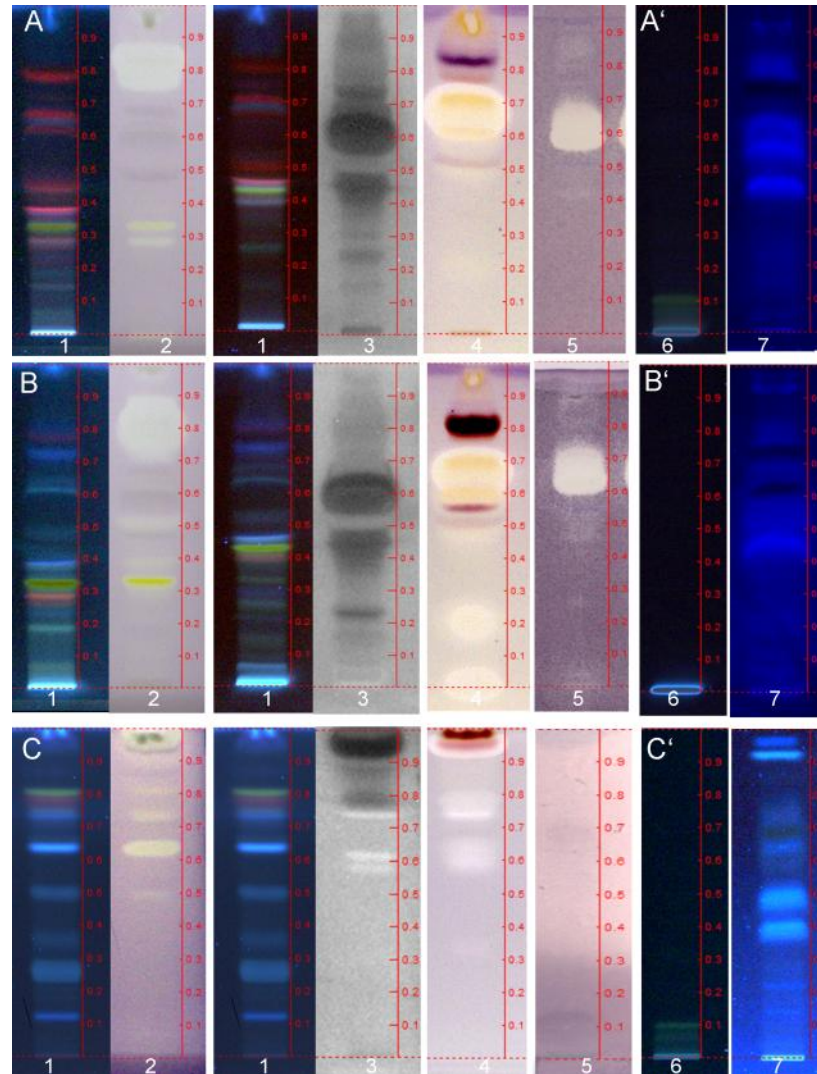
# Plant origin of O-type propolis?



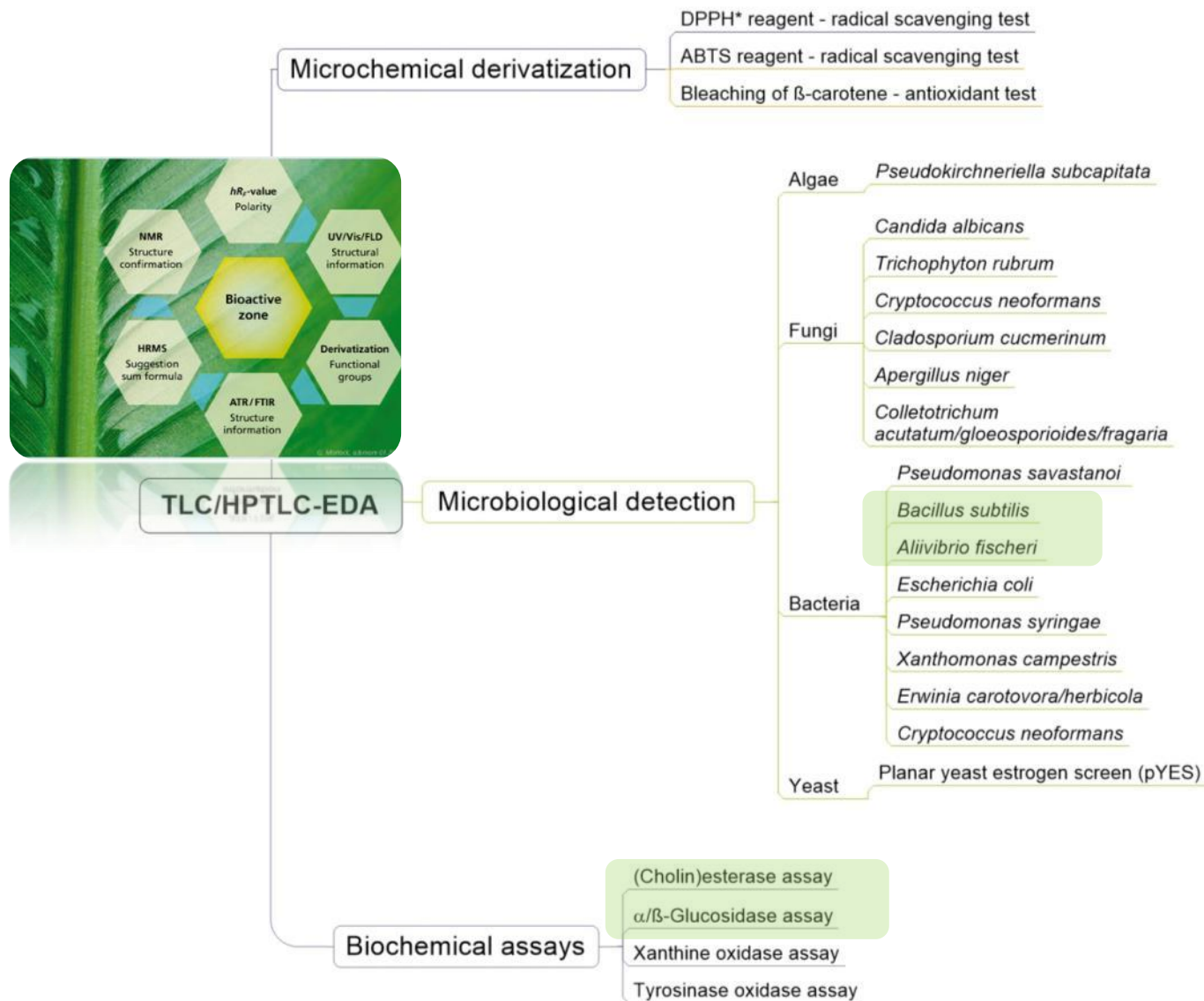
# EDA of *Salicaceae* bud extract samples



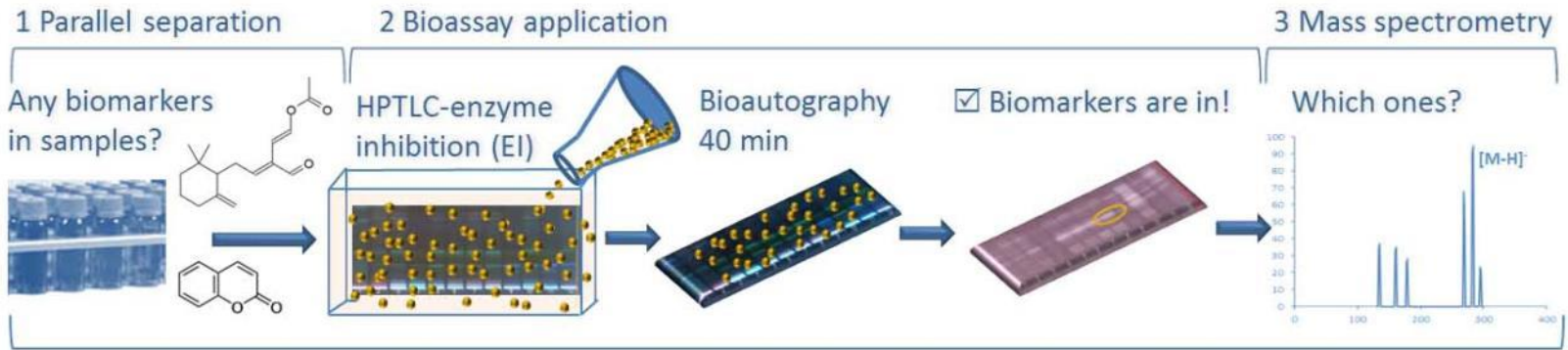
# EDA of oilseed extracts



# Effect-directed link to the compound



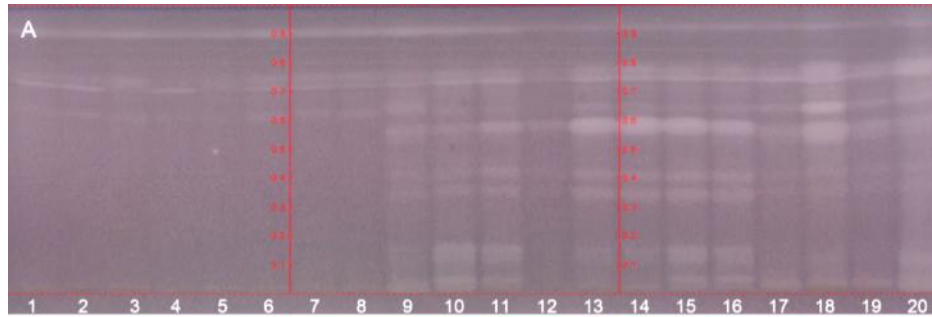
# Detection of cholinesterase inhibitors



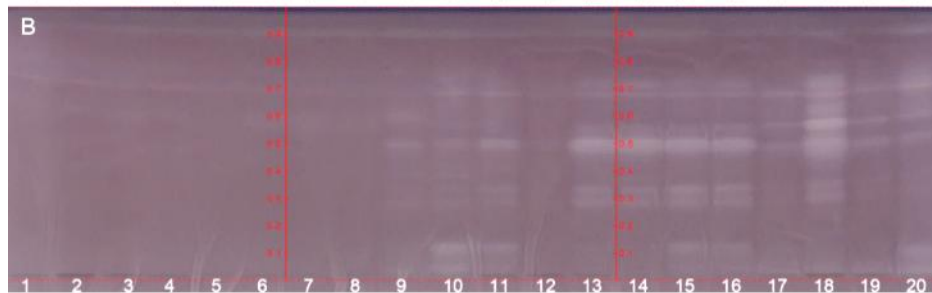
→ LC-EI-MS workflow for 20 samples in parallel within 2 h (6 min per sample)

*Rosaceae*

AChE

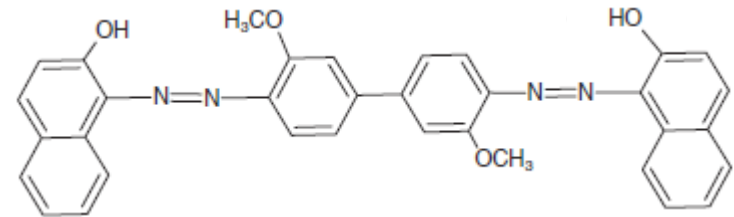


BChE



# Detection of $\alpha$ -/ $\beta$ -glucosidase inhibitors

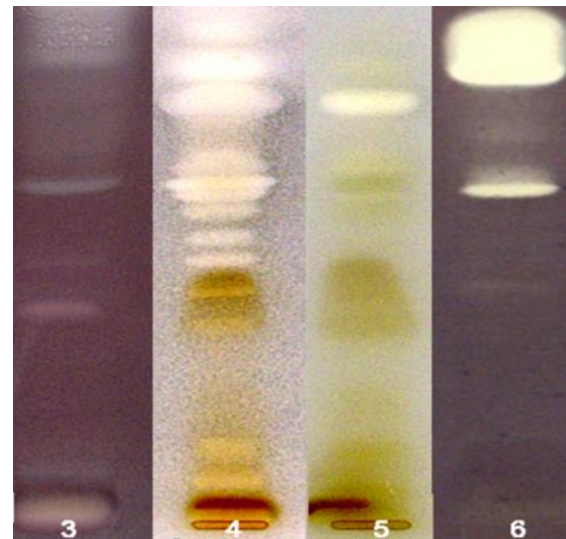
- Immersion in  $\alpha$ -/ $\beta$ -glucosidase solution and incubation
- Immersion in 2-naphthyl- $\alpha$ / $\beta$ -D-glucopyranoside and Fast Blue Salt as substrate for visualization (5 min)
- White zones on a violet background



*Salvia officinalis*



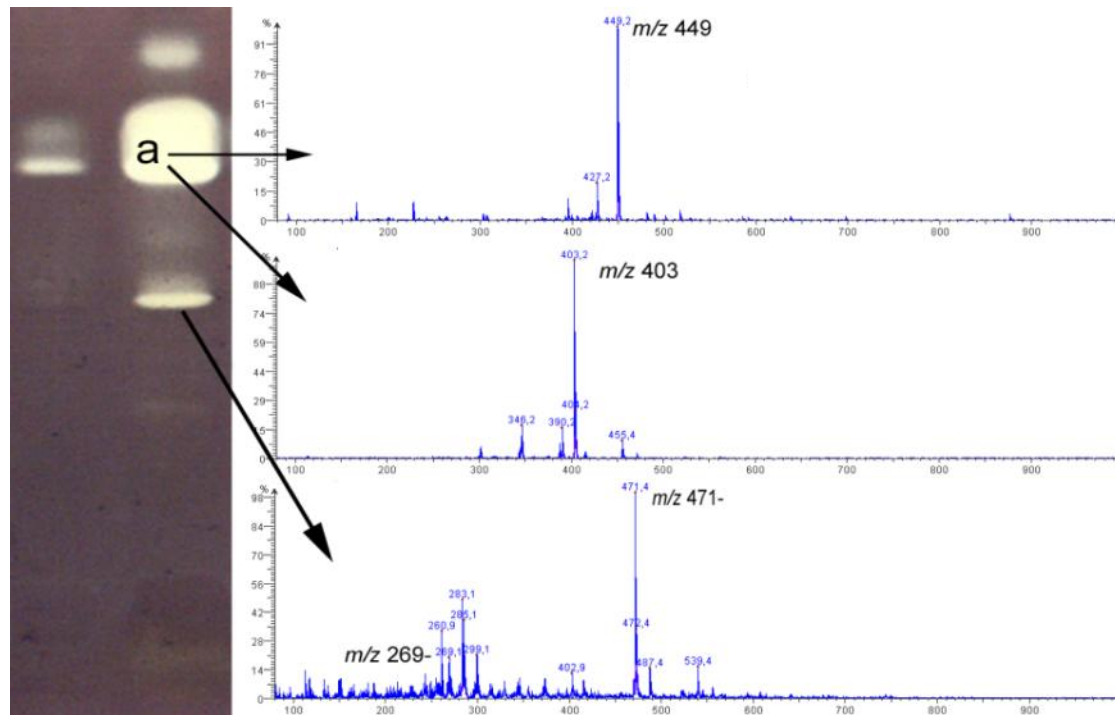
UV366 nm



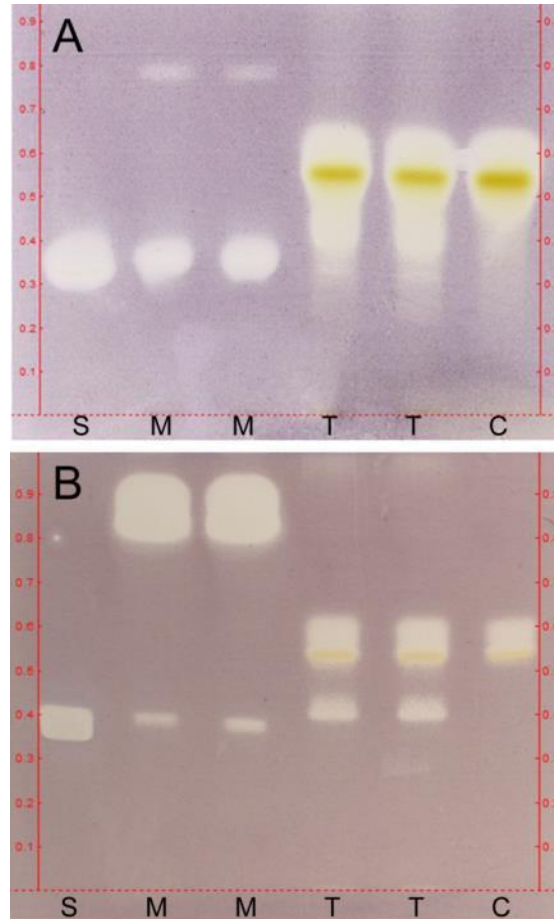
AChE  $\alpha$ -/ $\beta$ -glucosidase *B. subtilis*

# Detection of antibiotics

- Immersion in *Bacillus subtilis* bacteria suspension and incubation
- Immersion in tetrazolium salt as substrate for visualization
- White zones on a pink background



# Current vs. optimized detection of antibiotics

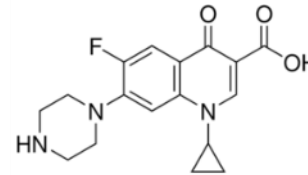




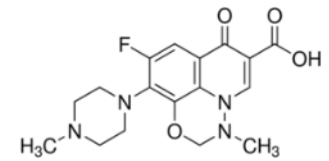
# Detection of antibiotics

Antibiotics in milk down to the  $\mu\text{g}/\text{kg}$  range

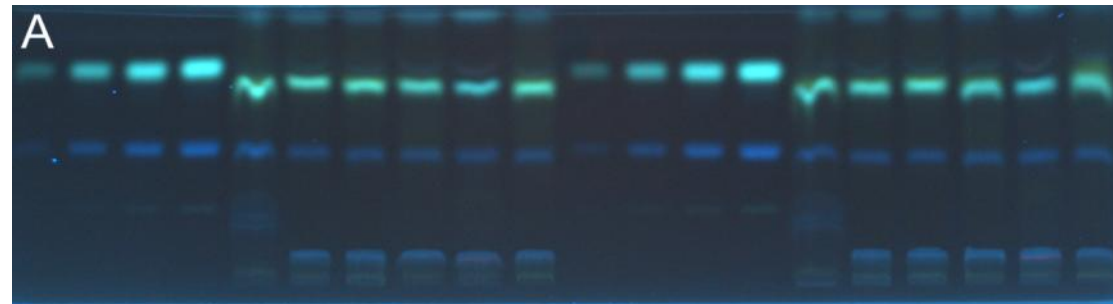
Ciprofloxacin



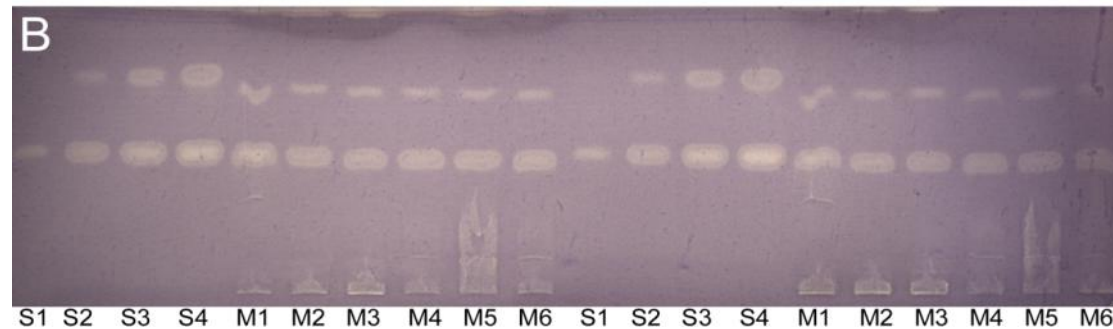
Marbofloxacin



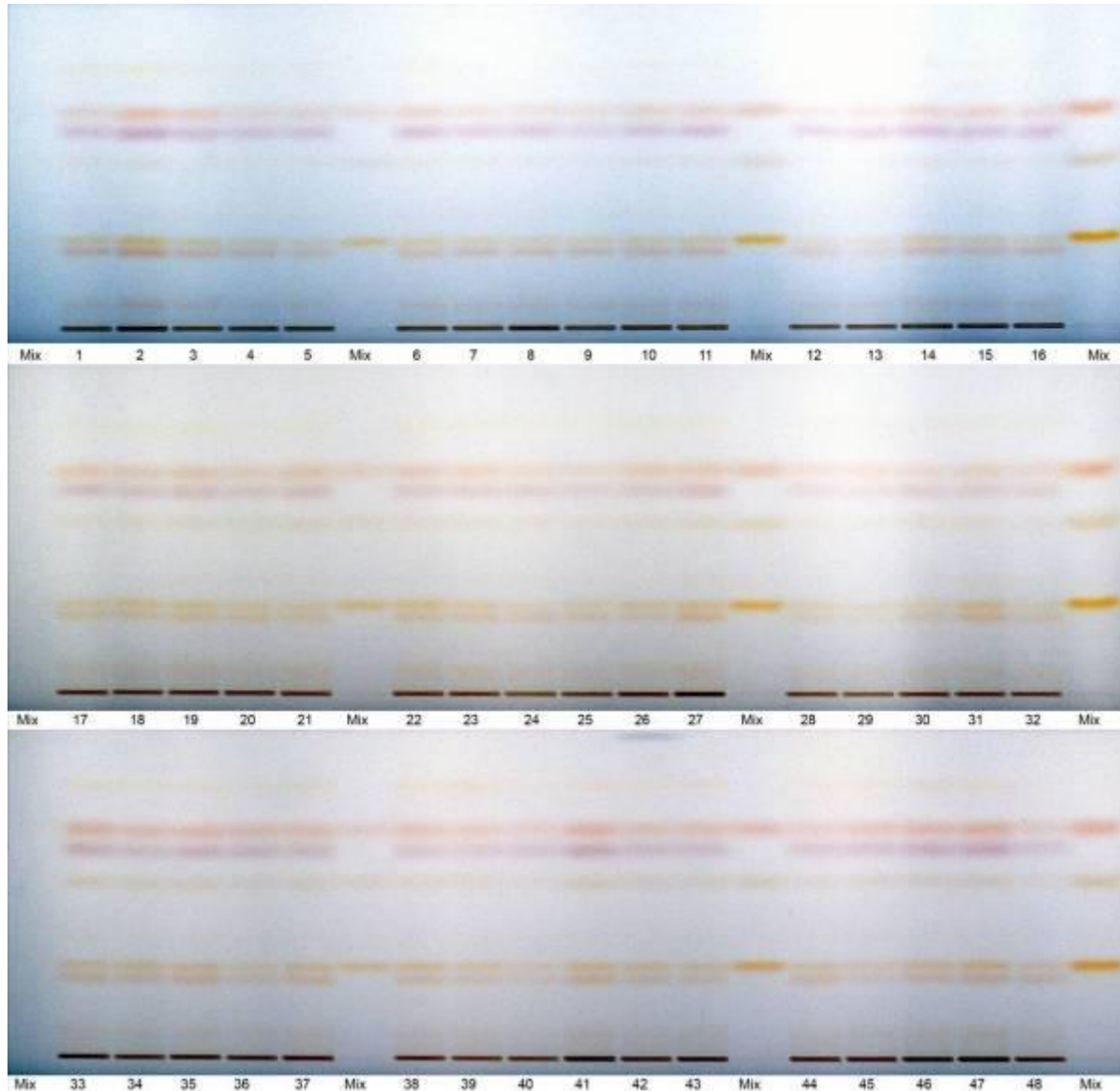
UV 366 nm



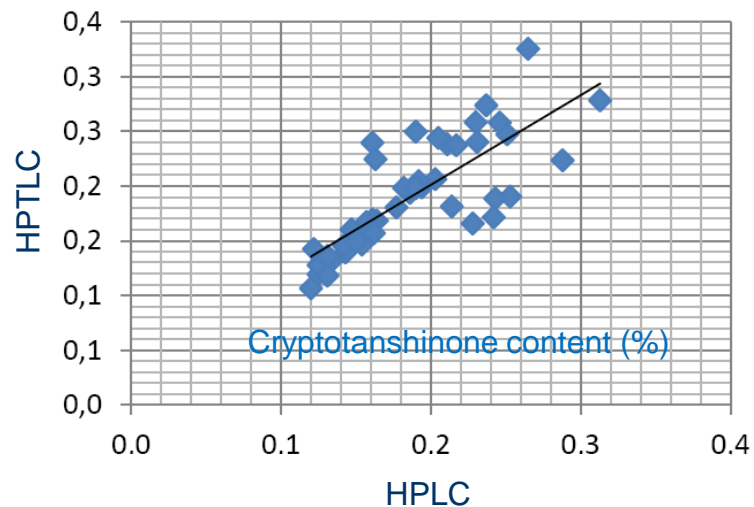
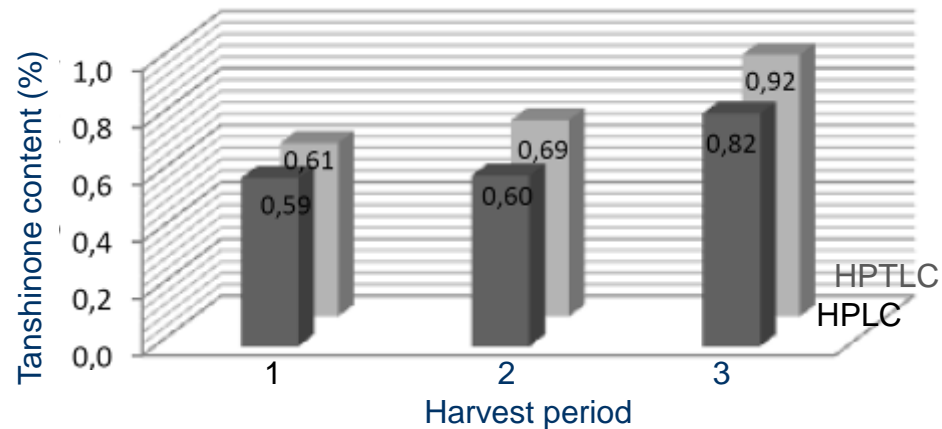
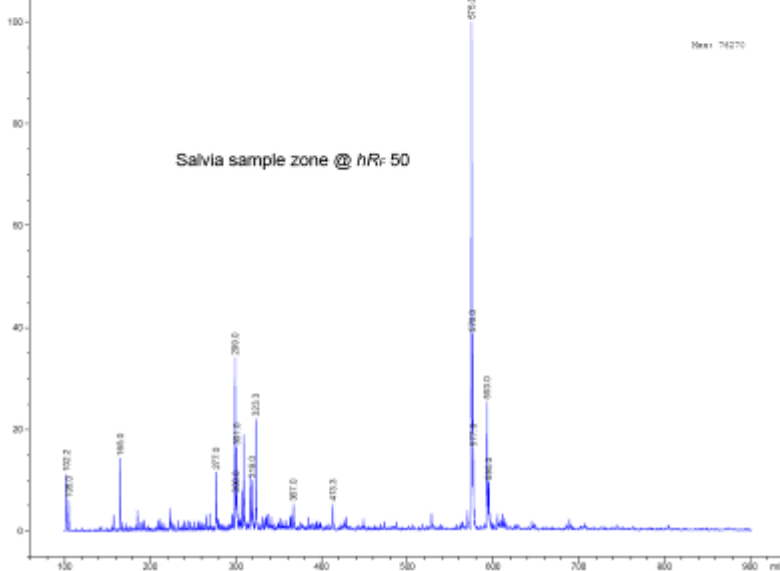
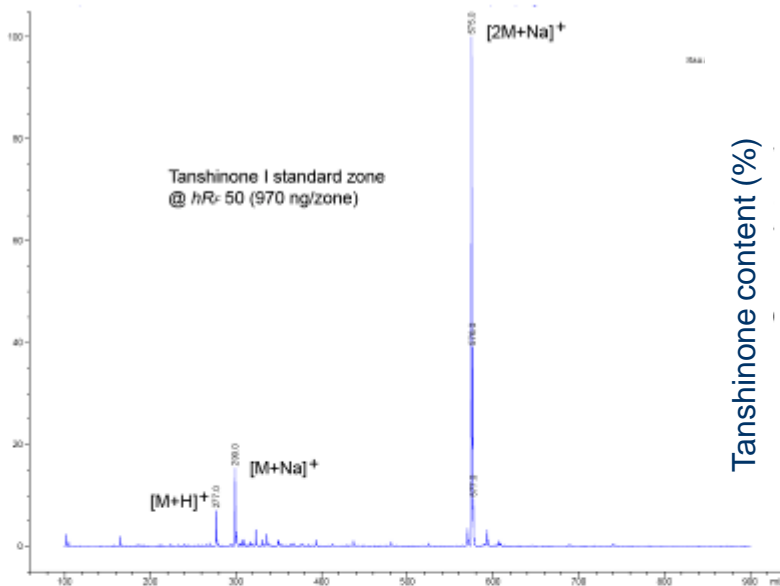
*Bacillus subtilis*



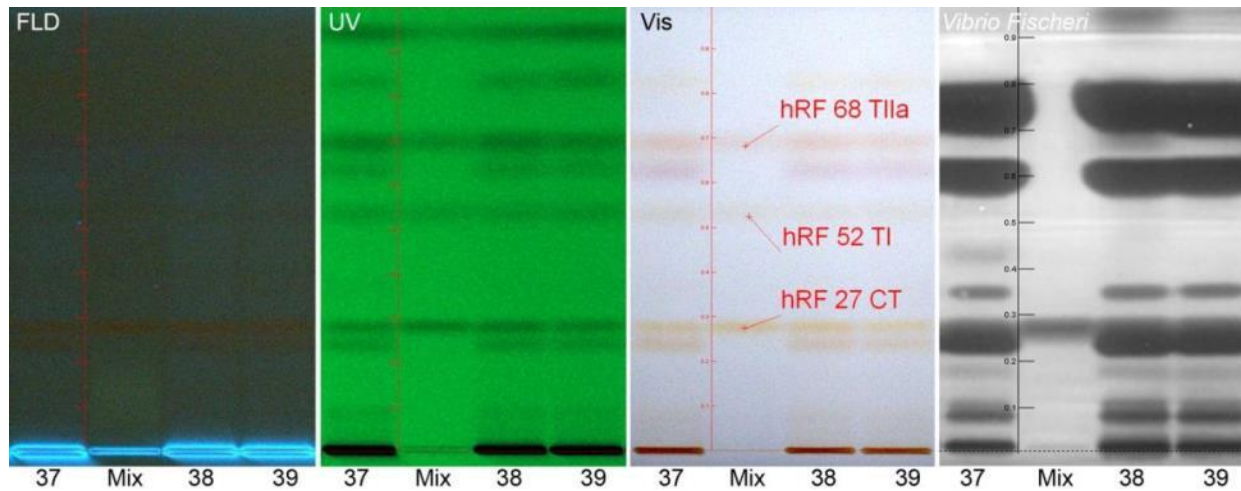
# Quantitation of tanshinons in Chinese salvia



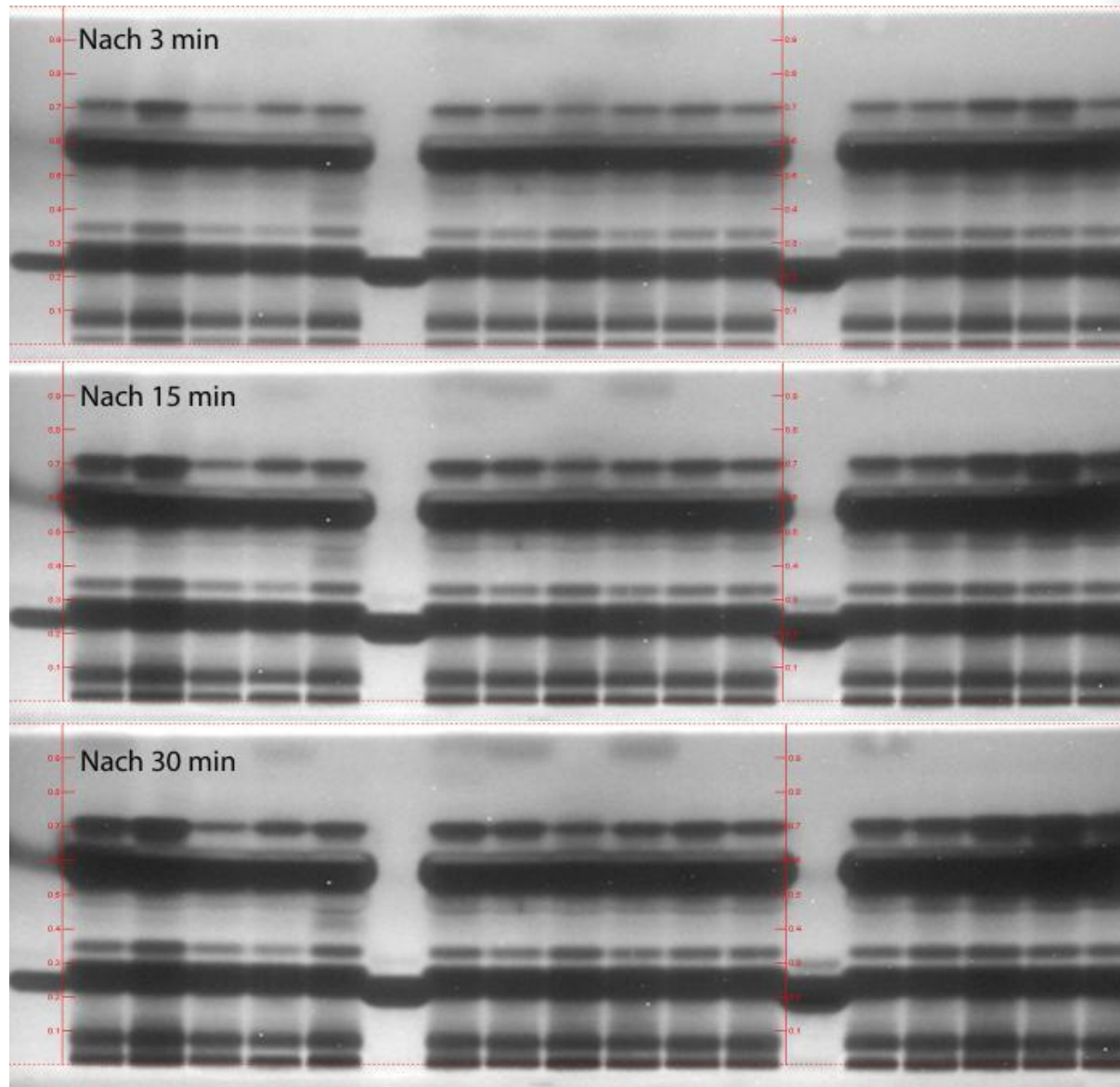
# Confirmation by MS and method comparison



# Detection of bioactive compounds

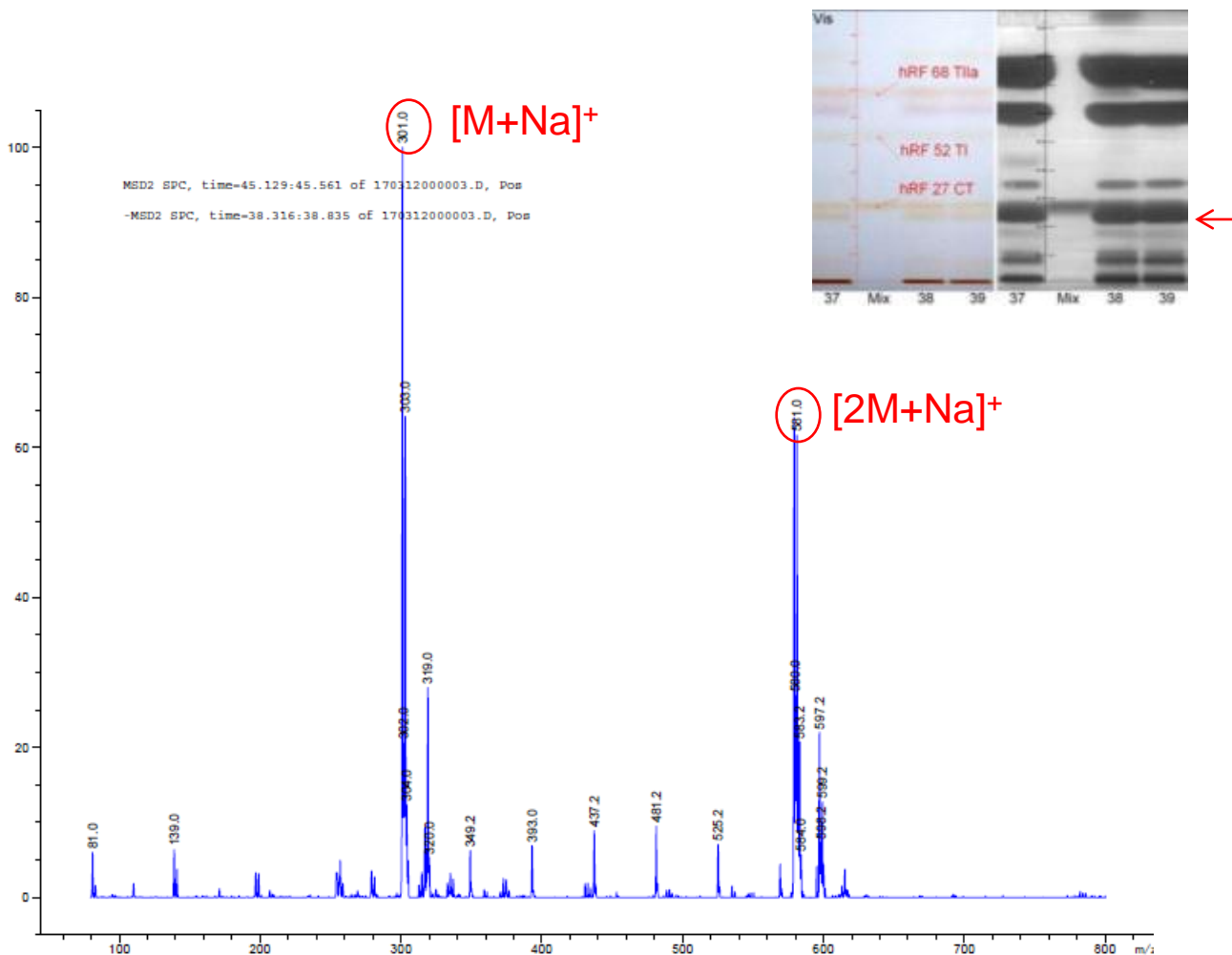


# Detection of bioactive compounds



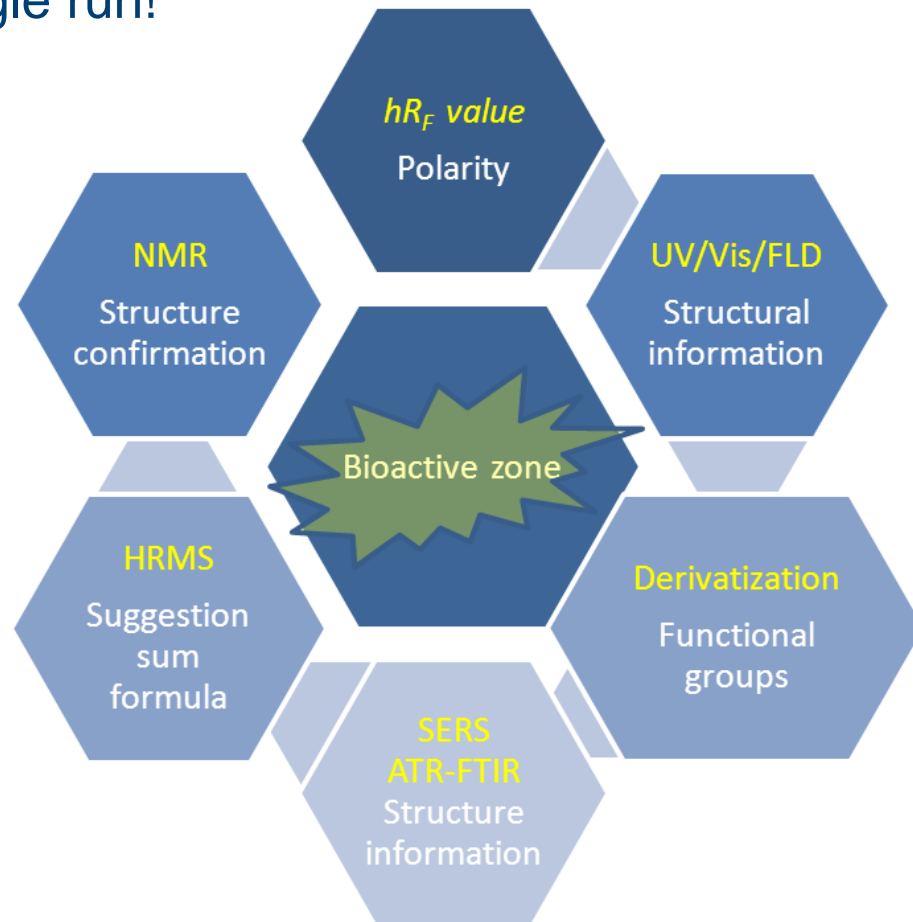
# Detection of bioactive compounds

Mass spectra recorded after detection with bioassay → salt adducts are pronounced!



# Goal: From bioactive zone to sum formula

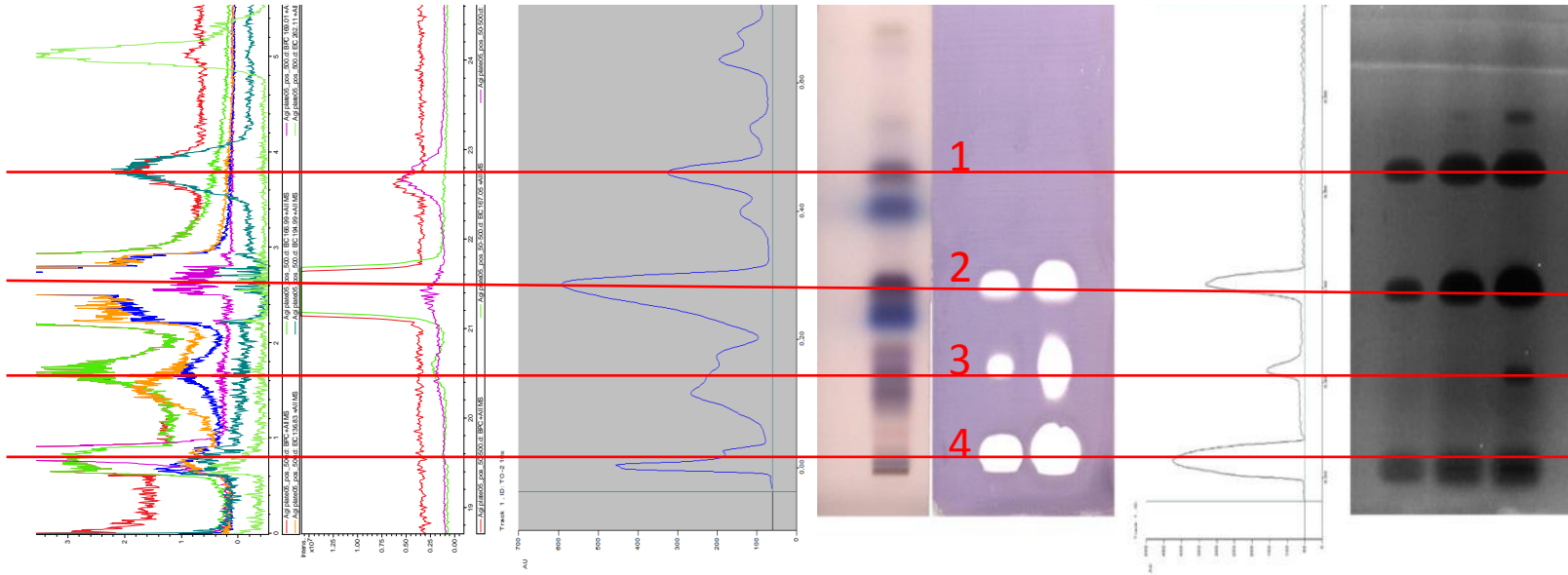
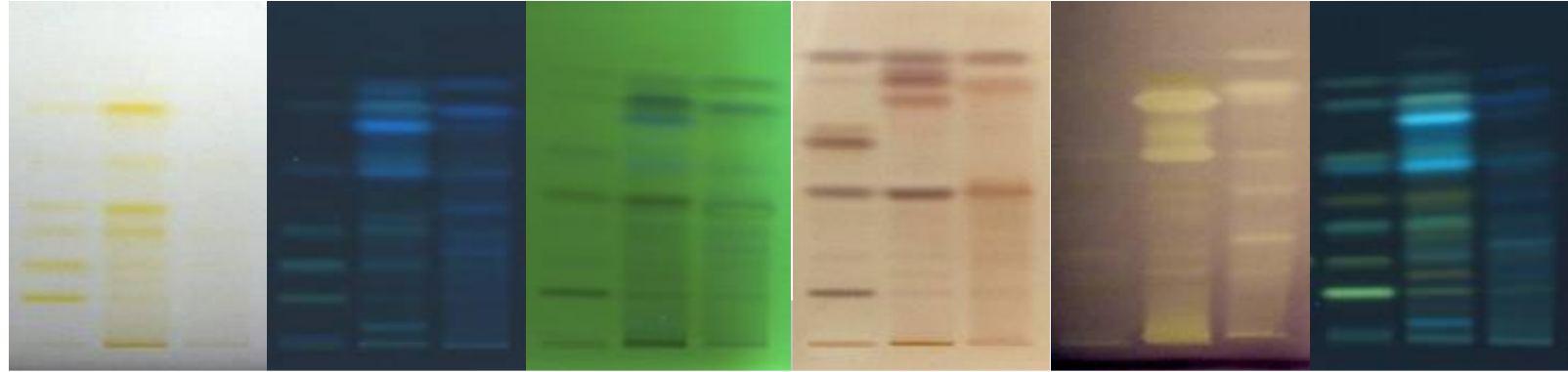
...in a single run!



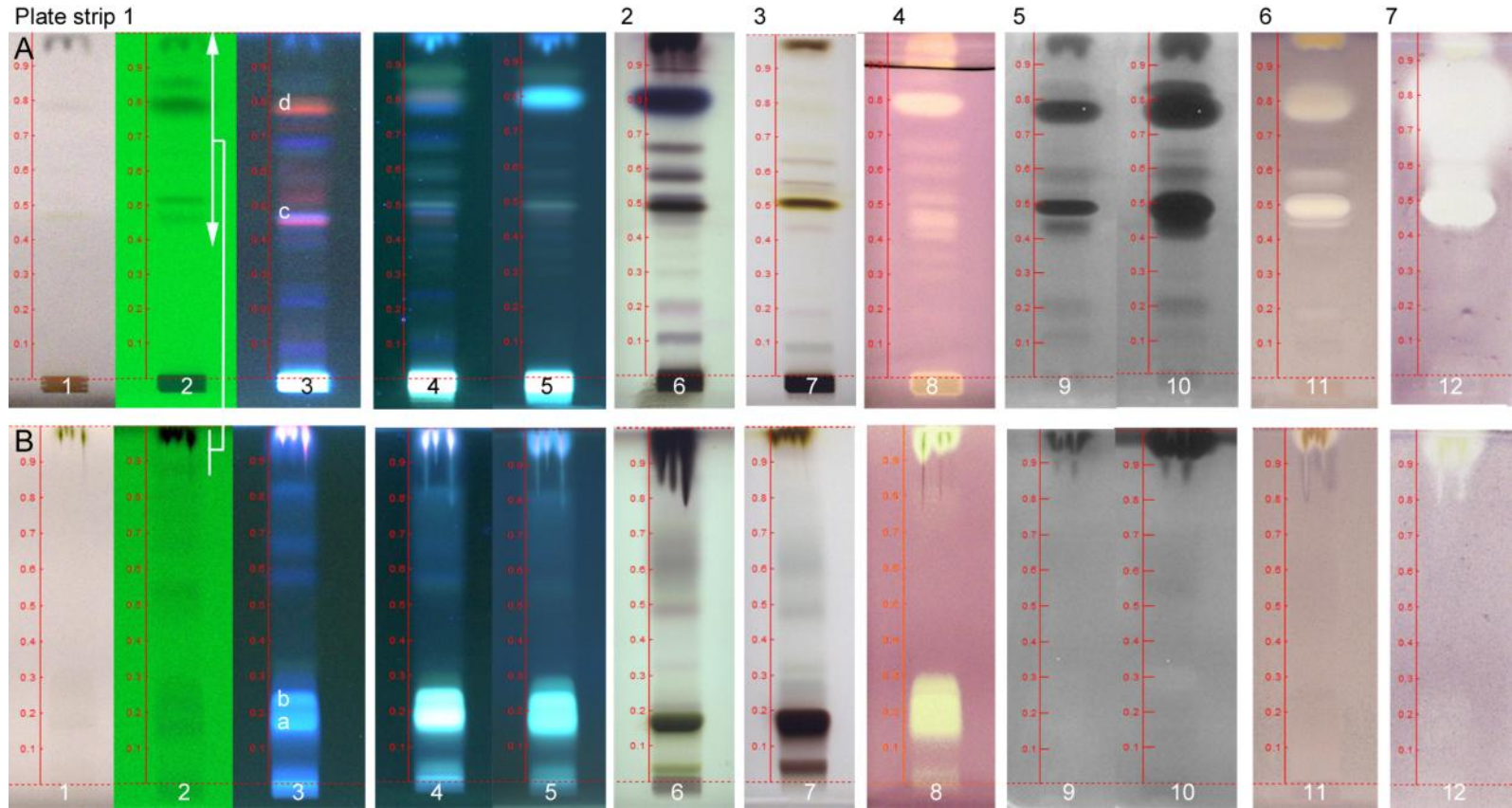




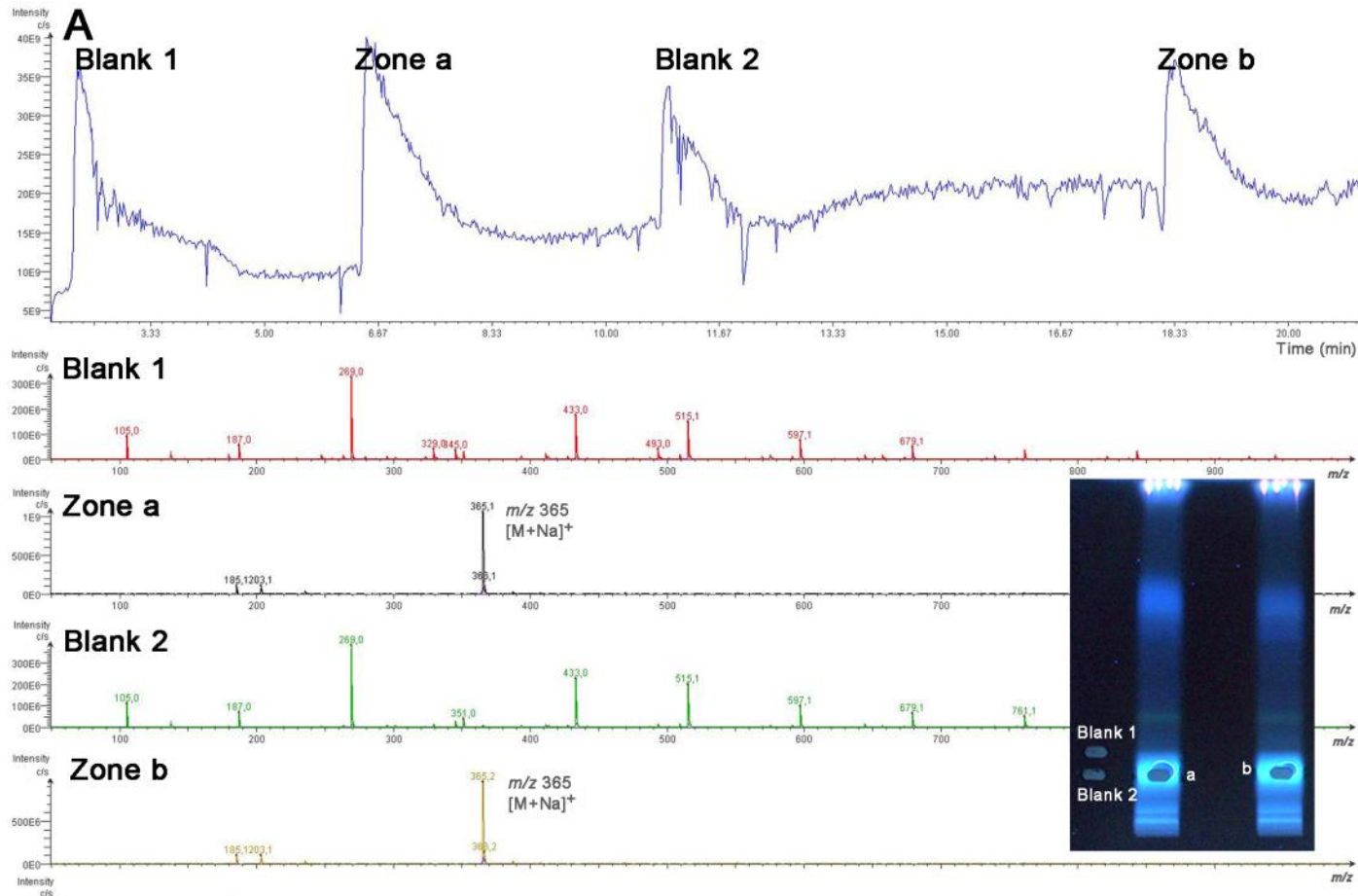
# Multi-detection



# EDA of *Pimpinella saxifraga* L. root extracts

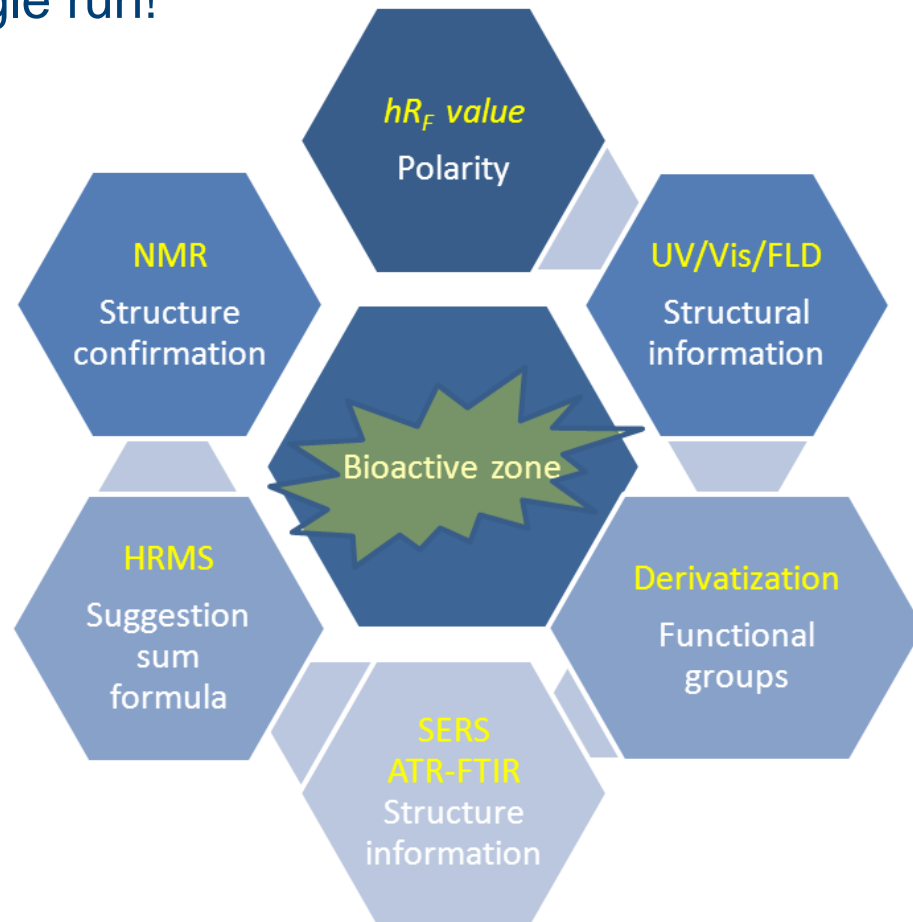


# EDA of *Pimpinella saxifraga* L. root extracts



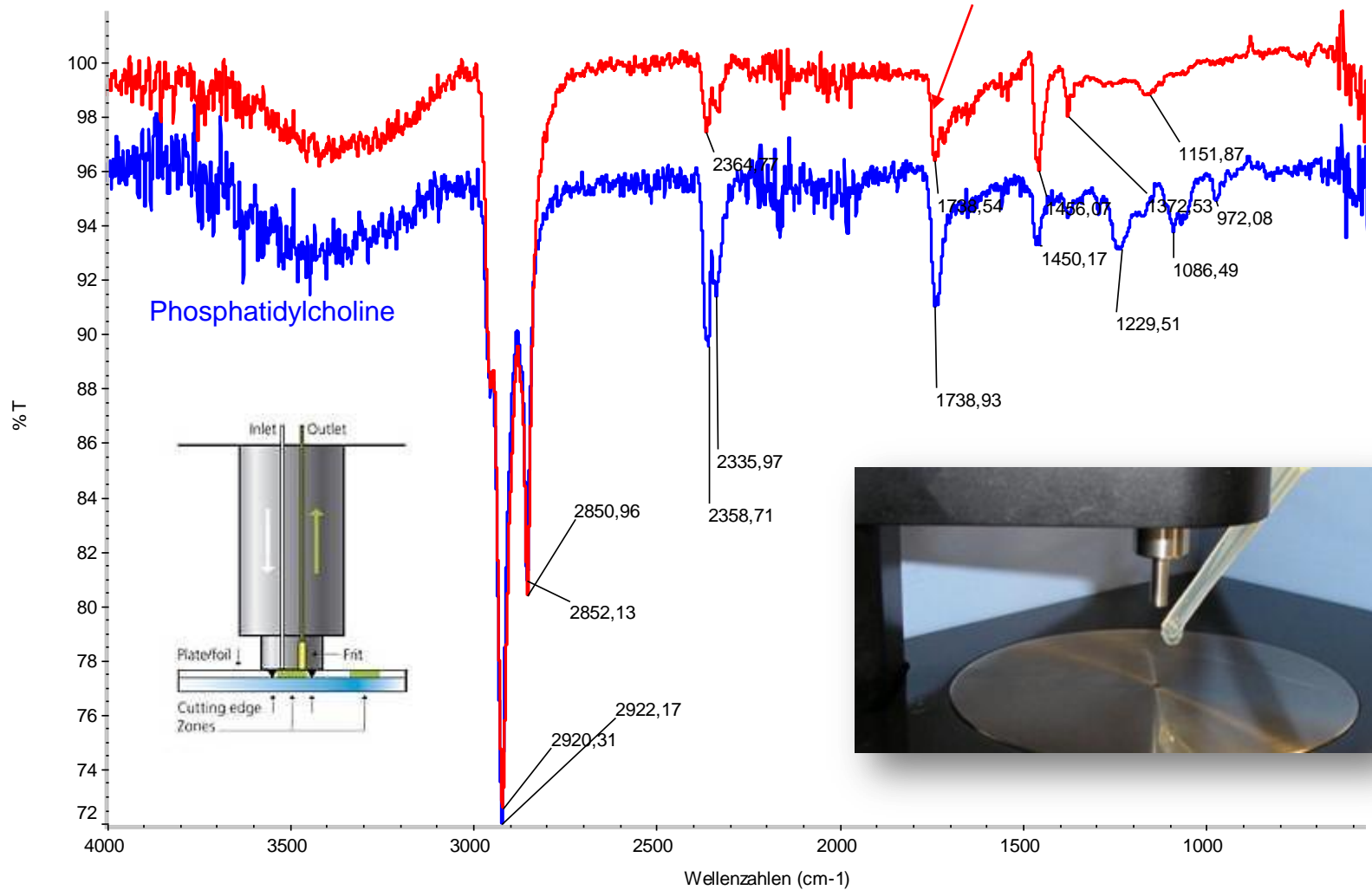
# Goal: From bioactive zone to sum formula

...in a single run!



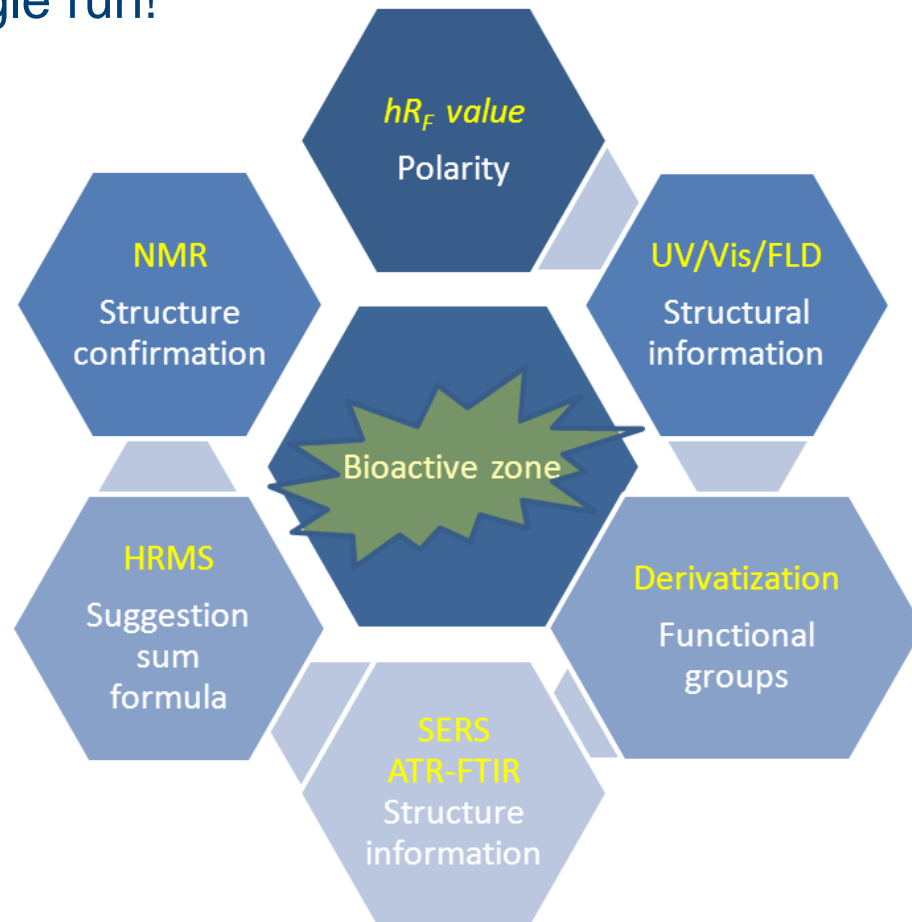
# HPTLC-ATR FTIR

Anti-inflammatory compound isolated from *Lactobacillus fermentum*



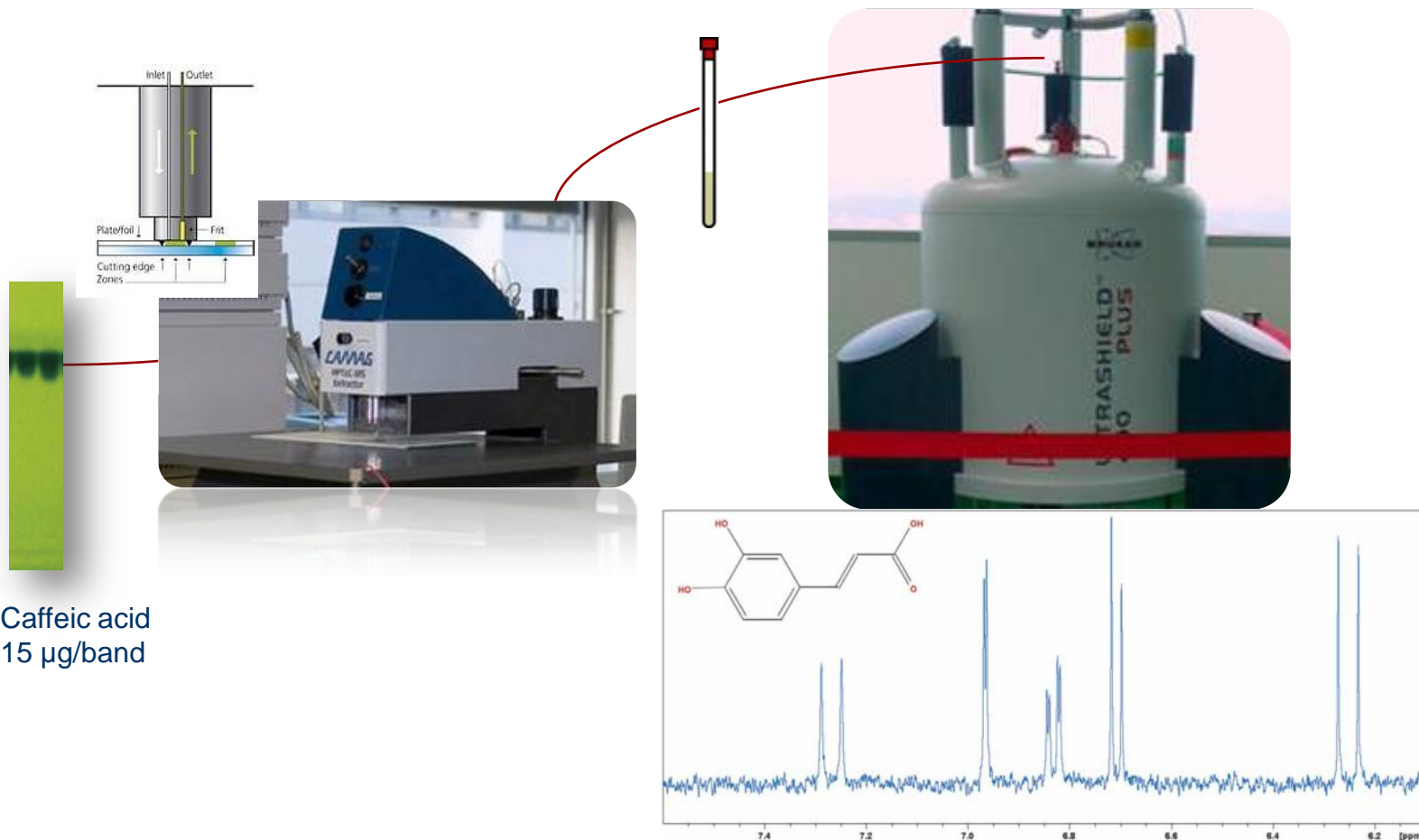
# Goal: From bioactive zone to sum formula

...in a single run!



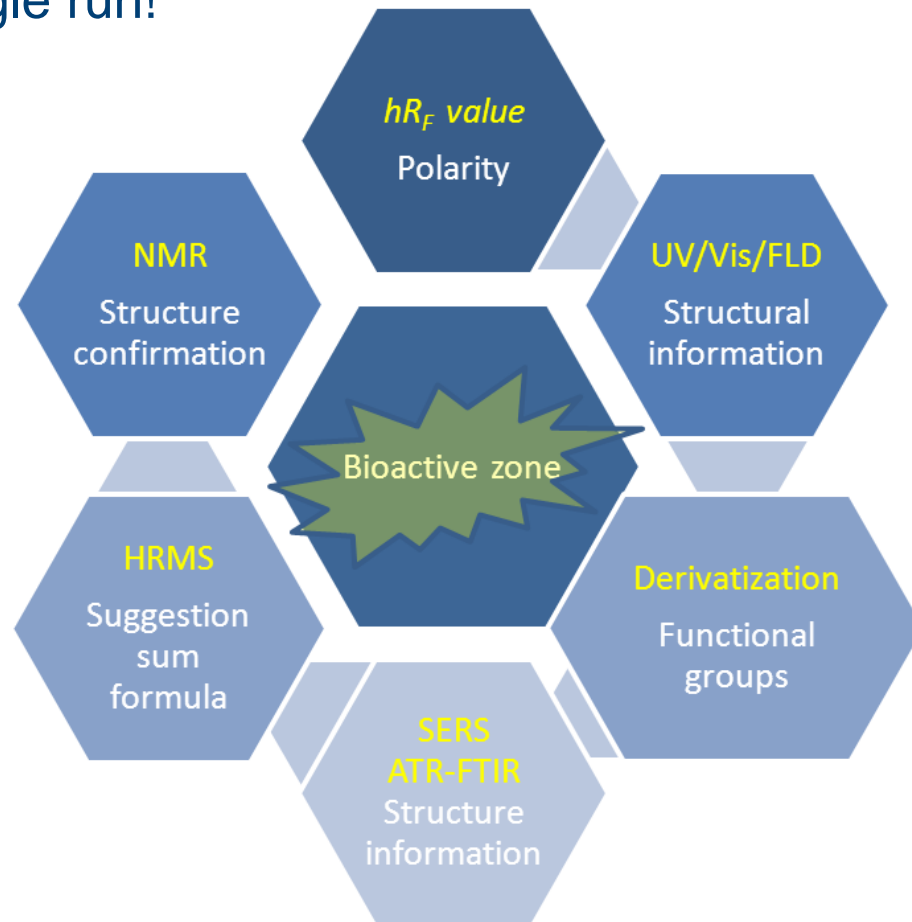
# HPTLC-NMR

→ Hyphenation of HPTLC with  $^1\text{H}$ -NMR via TLC-MS Interface



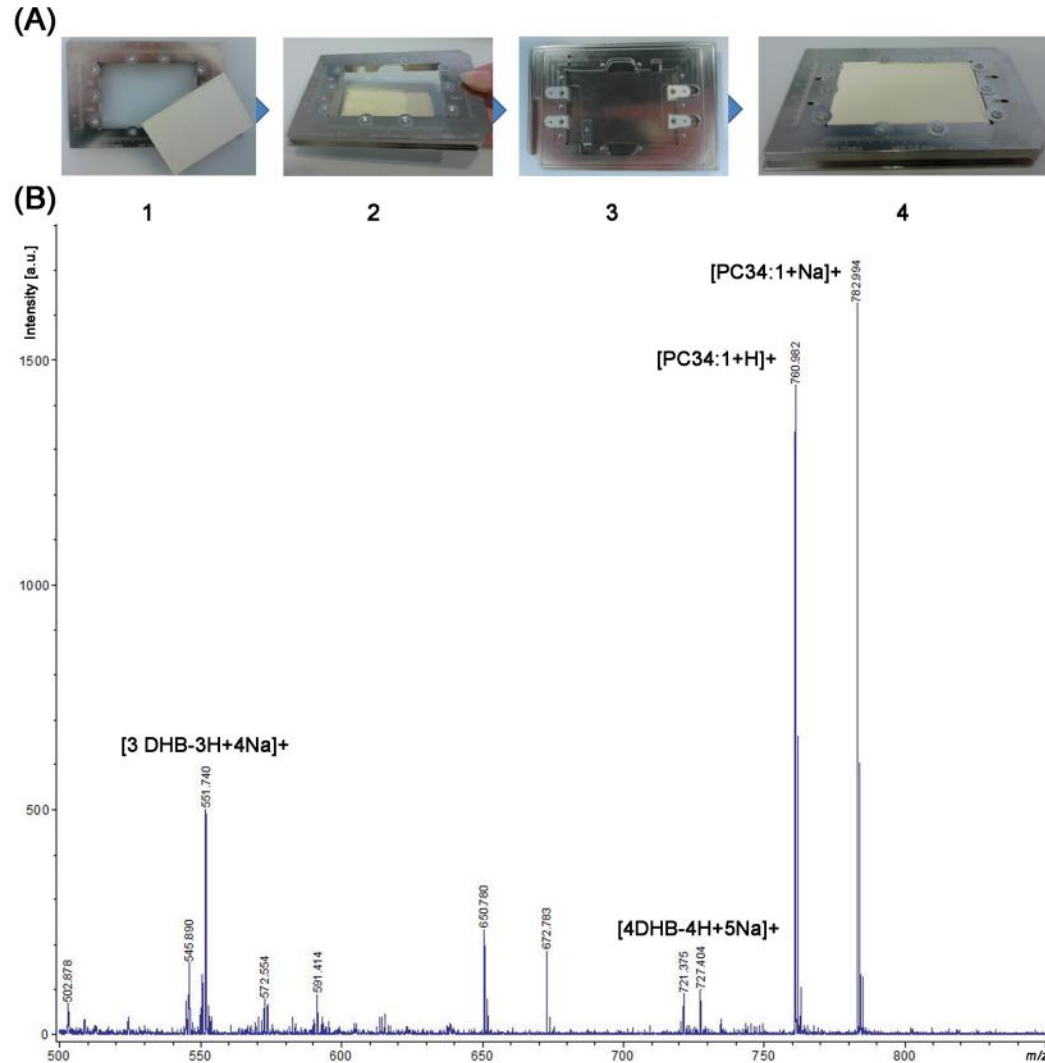
# Goal: From bioactive zone to sum formula

...in a single run!

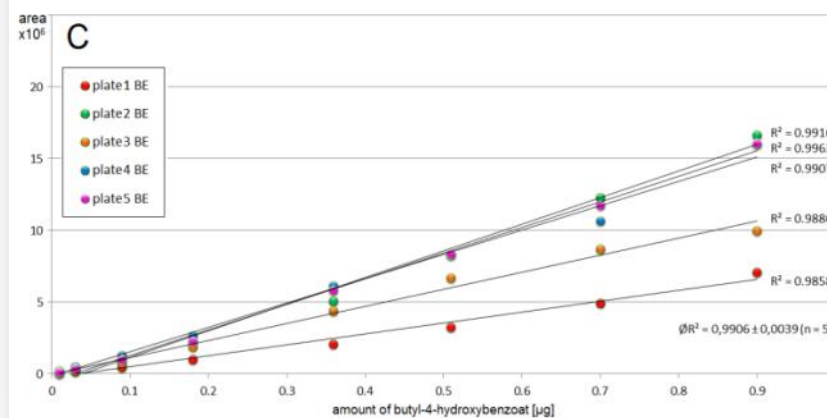
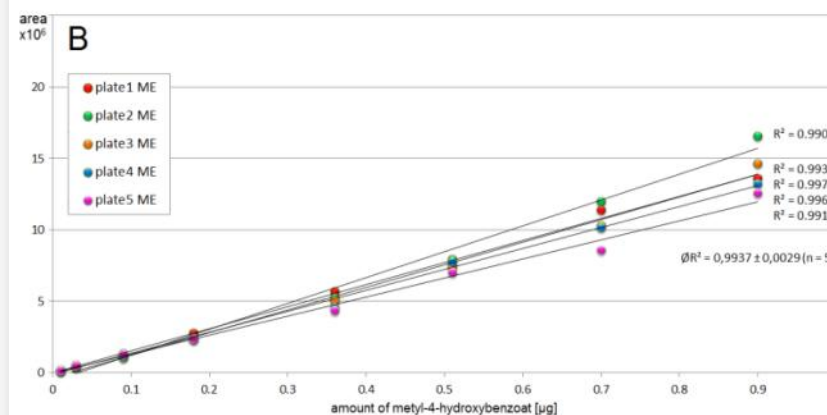
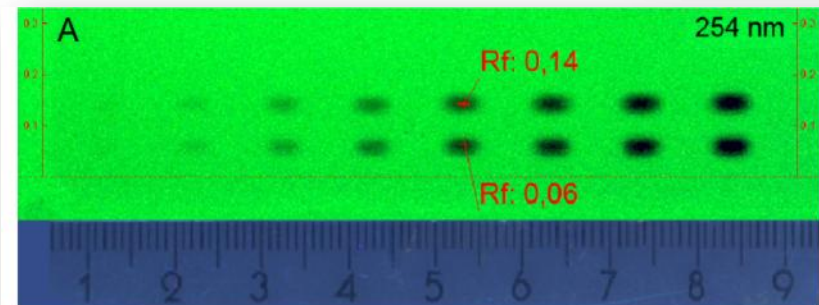
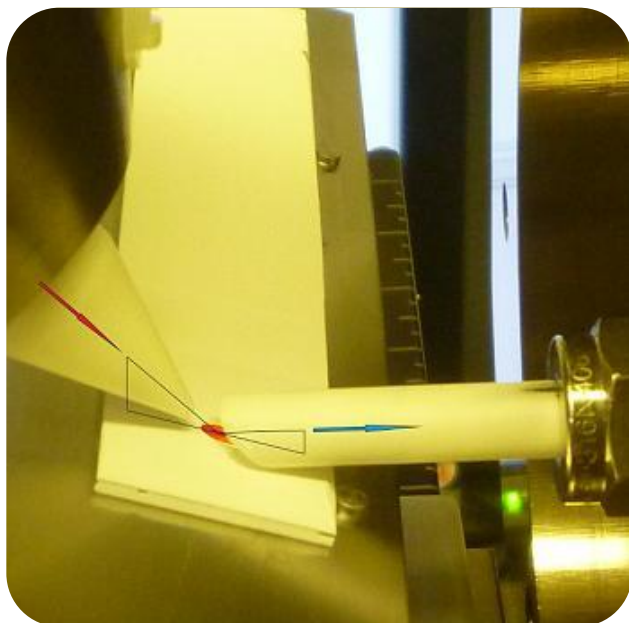
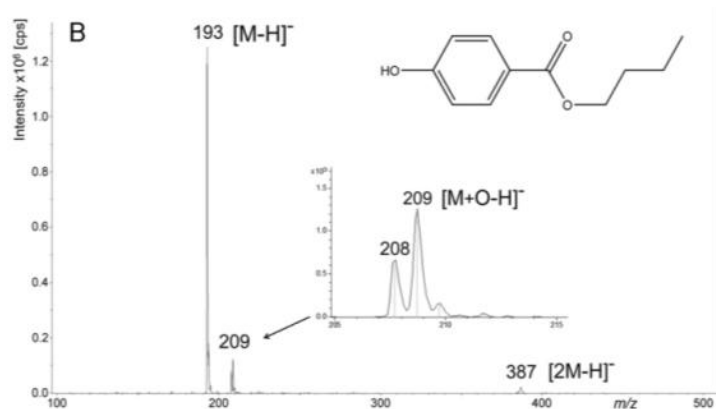




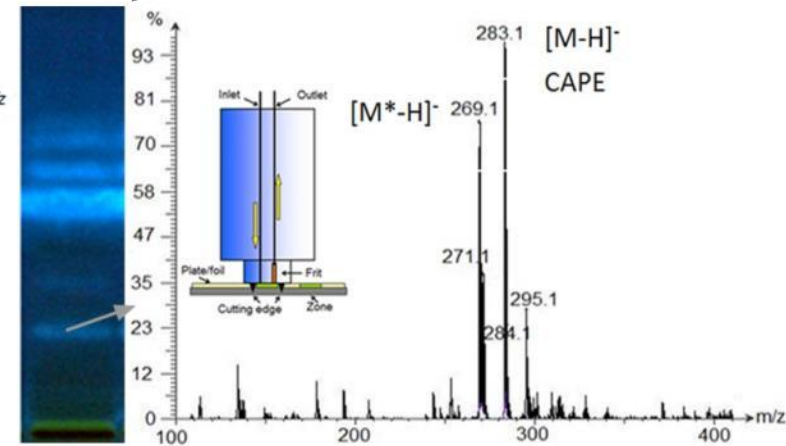
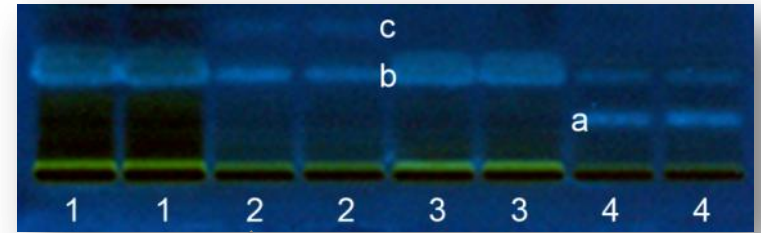
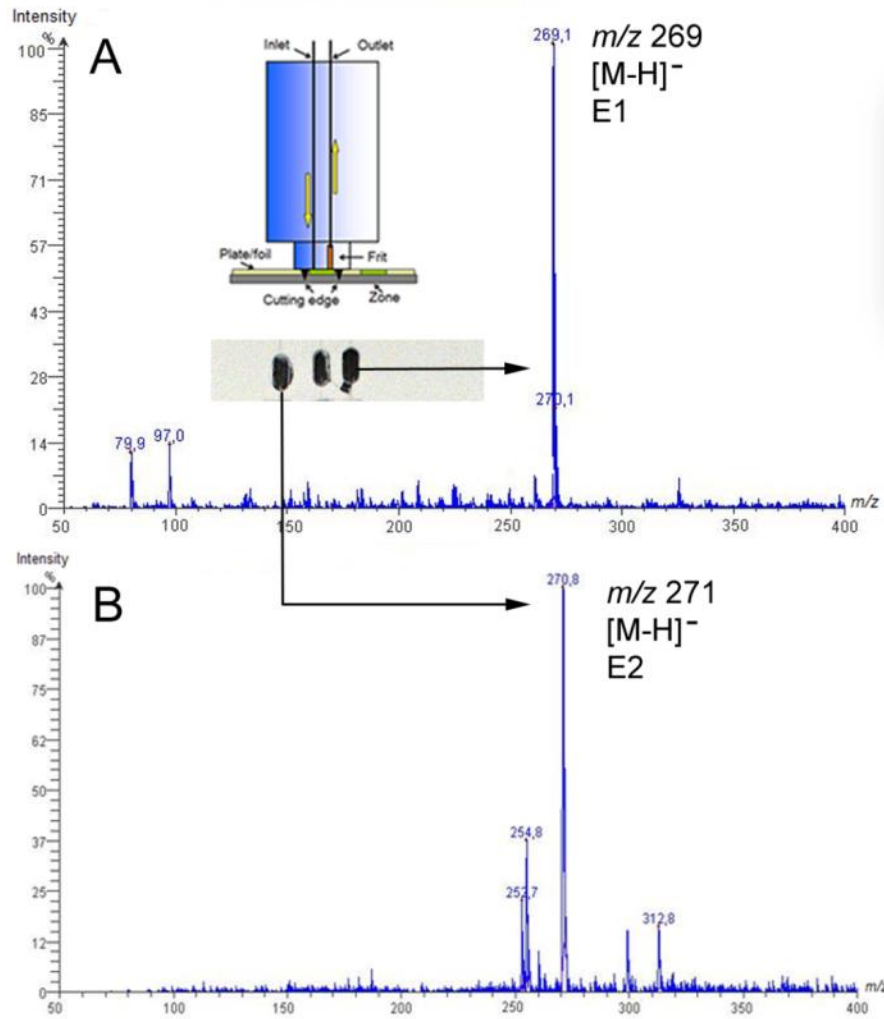
# HPTLC-MALDI-TOFMS of phospholipids



# HPTLC-DART-SVPA-MS of preservatives




# Confirmation by MS



I. Klingelhöfer, G. Morlock, *J. Chromatogr. A* 1360 (2014) 288-295

G. Morlock, I. Klingelhöfer, *Anal. Chem.* 86 (2014) 8289-8295

# GDCh course 335/15



GESELLSCHAFT DEUTSCHER CHEMIKER

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## Hyphenations in der HPTLC

HPTLC und Kopplungen  
(in Zusammenarbeit mit der JLU Gießen)

---

Prof. Dr. Gertrud Morlock

- Kopplungstechniken
- Massenspektrometrie (MS)
- Wirkungsbezogene Analytik (Bioassays)
- ATR-FTIR und NMR
- Effektive Analytik




335/15

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11. November 2015 · Gießen

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**Anerkannt mit 18 Punkten**  
(www.zfo.org)

Zentrum für  
Lebensmittelwissenschaften

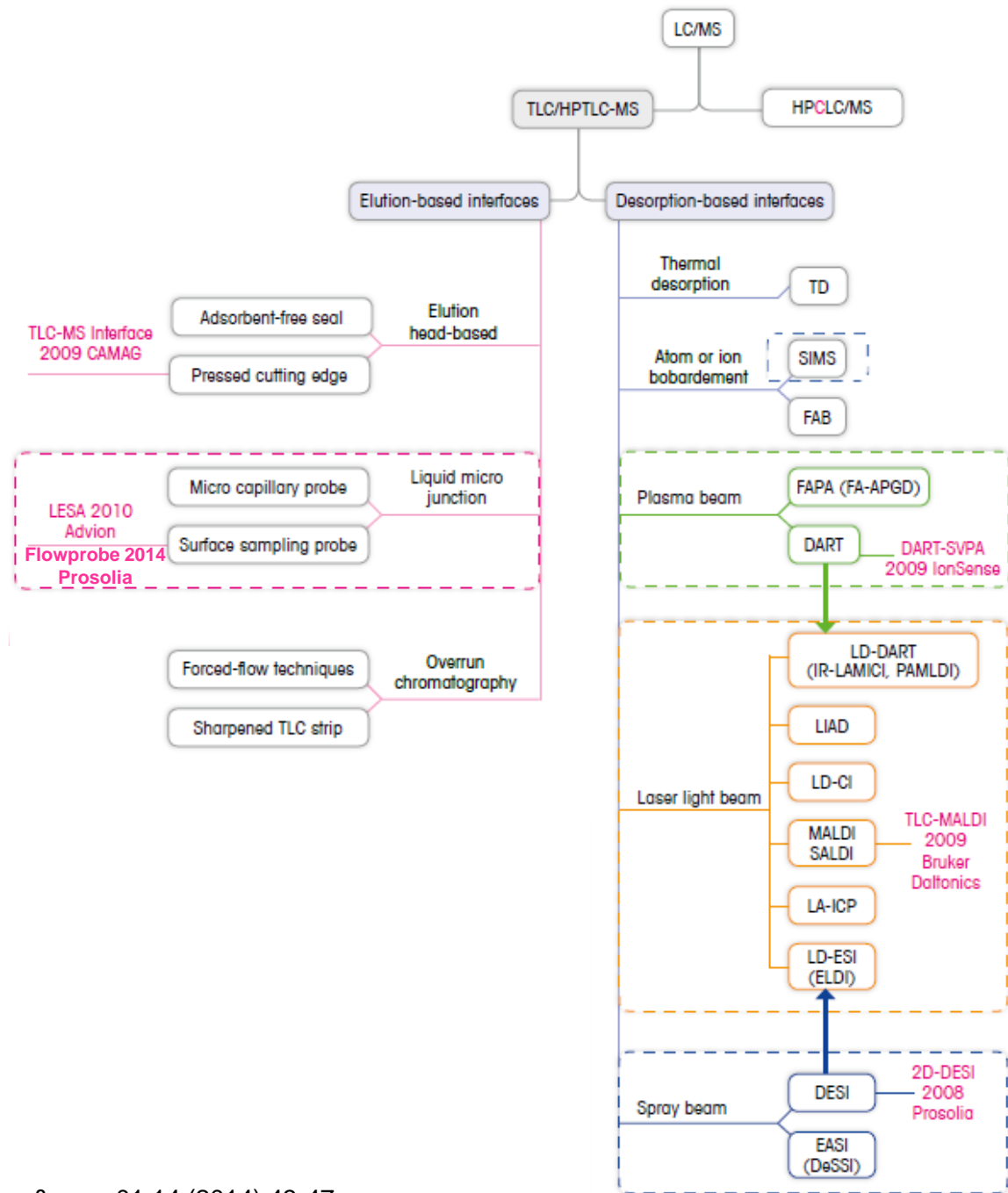
ANALYTISCHE CHEMIE

## PROGRAMM

Mittwoch, 11. November 2015

- 9.00 Begrüßung und Einführung in die HPTLC (Morlock)
- 9.45 HPTLC erfahren – Experimente (Häbe, Klingelhöfer)
- 10.45 Kaffeepause
- 11.00 Hyphenations in der Planar-Chromatographie – Teil 1 (Morlock, Schwack)
- 11.45 Gruppe 1: Experiment DC-HPLC/DAD-ESI MS (Oellig, Schwack)  
Gruppe 2: Experiment HPTLC-UV/Vis/FLD-MALDI-TOF MS/MS (Lochnit, Krüger)
- 12.30 Gruppe 1: Experiment HPTLC-UV/Vis/FLD-MALDI-TOF MS/MS (Lochnit, Krüger)  
Gruppe 2: Experiment DC-HPLC/DAD-ESI MS (Oellig, Schwack)
- 13.15 Mittagspause
- 13.45 Hyphenations in der Planar-Chromatographie – Teil 2 (Morlock)
- 14.00 Gruppe 1: Experiment HPTLC-UV/Vis/FLD-ATR FTIR (Klingelhöfer, Gerbig)  
Gruppe 2: Experiment HPTLC-UV/Vis/FLD-Bioassay-ESI MS (Krüger, Kirchert)
- 14.45 Gruppe 1: Experiment HPTLC-UV/Vis/FLD-Bioassay-ESI MS (Krüger, Kirchert)  
Gruppe 2: Experiment HPTLC-UV/Vis/FLD-ATR FTIR (Klingelhöfer, Gerbig)
- 15.30 Kaffeepause
- 15.45 Hyphenations in der Planar-Chromatographie – Teil 3 (Morlock)
- 16.00 Gruppe 1: Experiment HPTLC-UV/Vis/FLD-DART-MS (Häbe, Krüger)  
Gruppe 2: Experiment HPTLC-UV/Vis/FLD-DESI-MS (Kirchert, Stiefel)
- 16.15 Gruppe 1: Experiment HPTLC-UV/Vis/FLD-DESI-MS (Kirchert, Stiefel)  
Gruppe 2: Experiment HPTLC-UV/Vis/FLD-DART-MS (Häbe, Krüger)
- 16.30 Diskussion (Morlock)
- 17.00 Voraussichtliches Ende der Veranstaltung





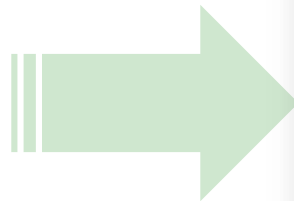


Bioactive compounds of interest in botanicals used as spices or food supplements?



# GDCh course 338/15

The day after



**GDCh**  
GESELLSCHAFT DEUTSCHER CHEMIKER

## EDA by HPTLC- (bio)assay-HRMS

Prof. Dr. Gertrud Morlock

**12. November 2015**

**ZFL** **ZFO**  
Anerkannt mit 18 Punkten  
(www.zfo.org)

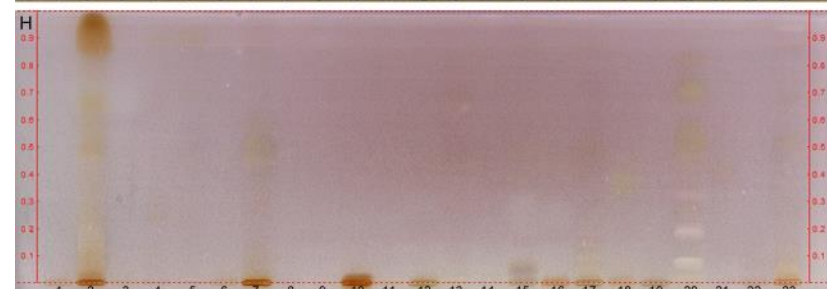
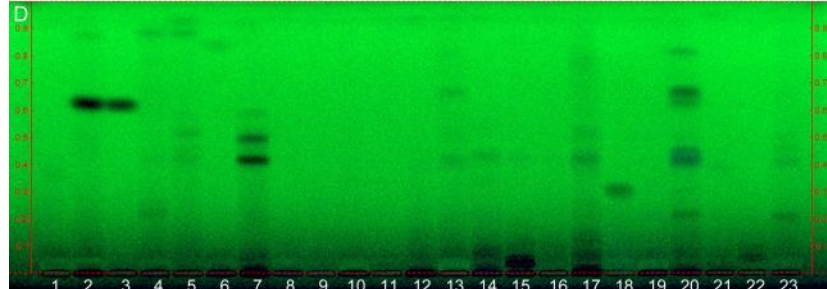
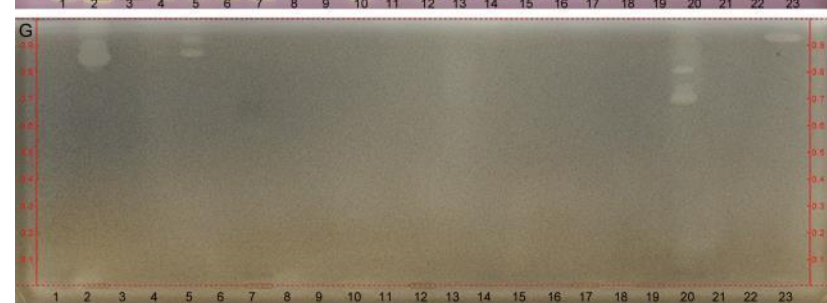
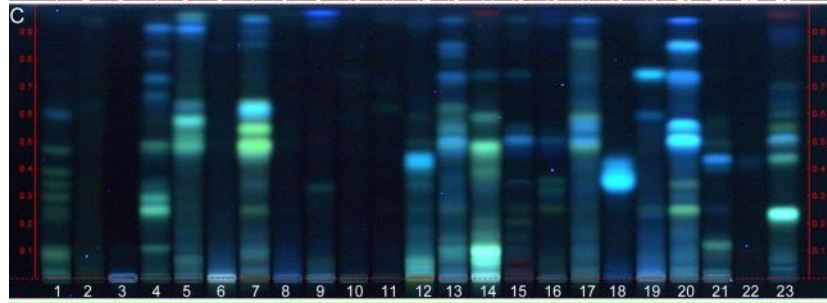
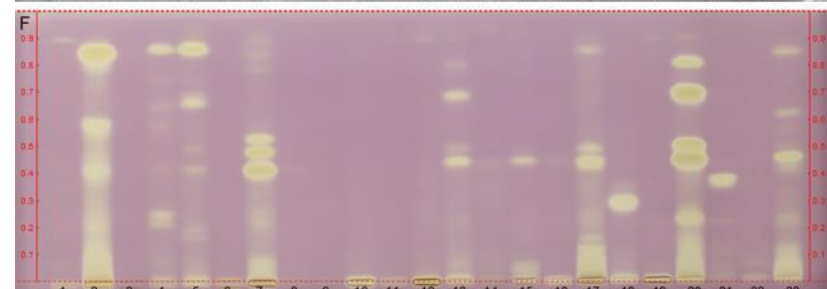
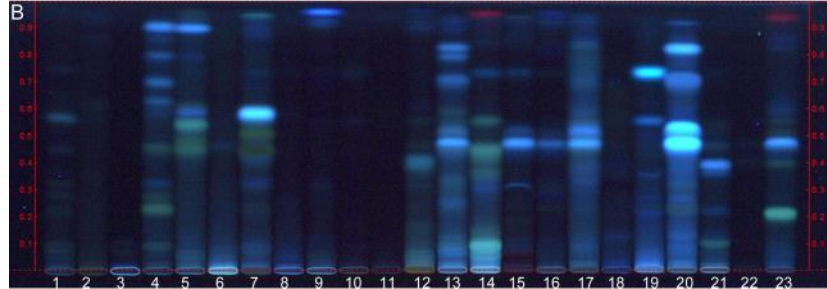
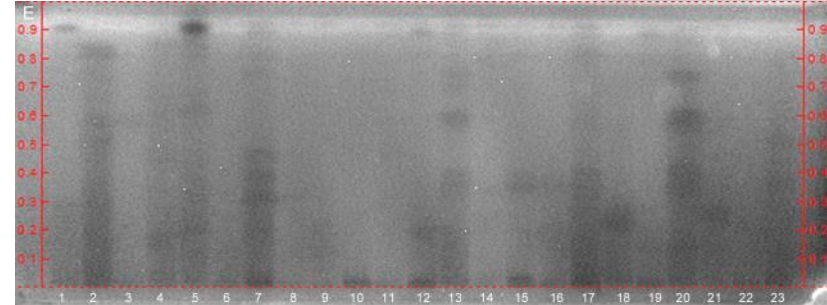
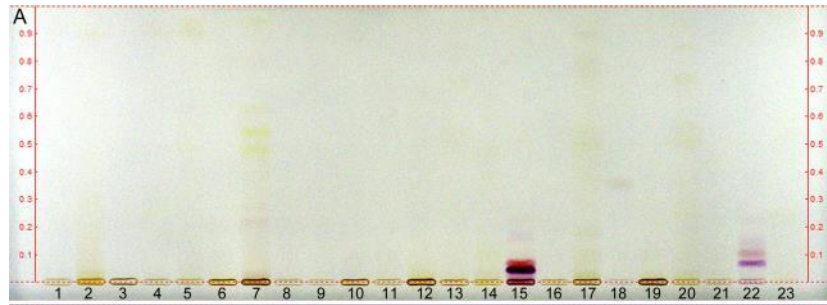
**A N A L Y T I S C H E C H E M I E**

# Effect-directed analysis (EDA) of 68 botanicals

JLU Gießen

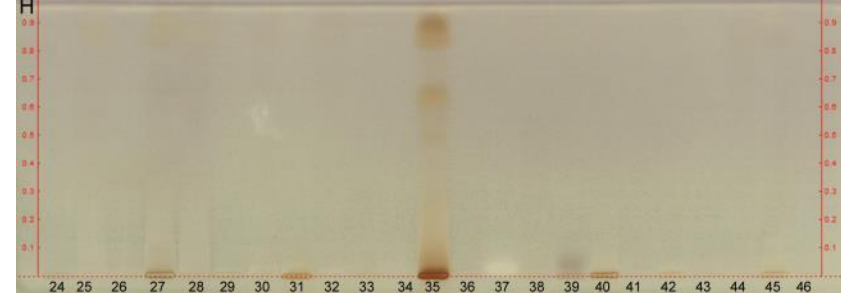
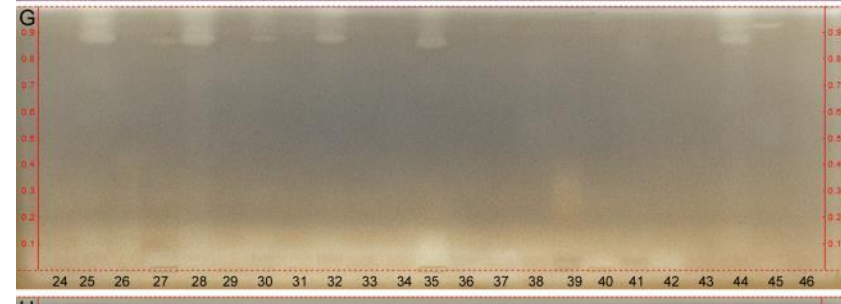
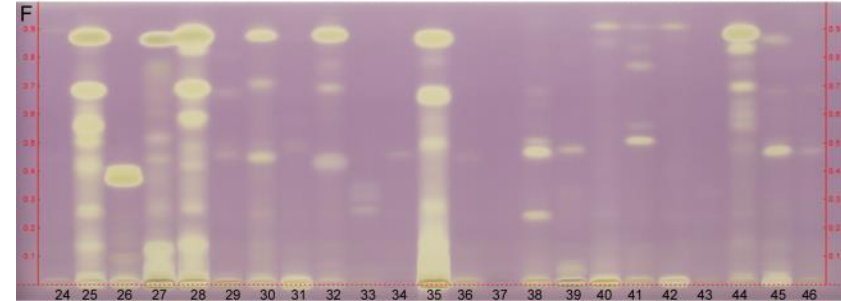
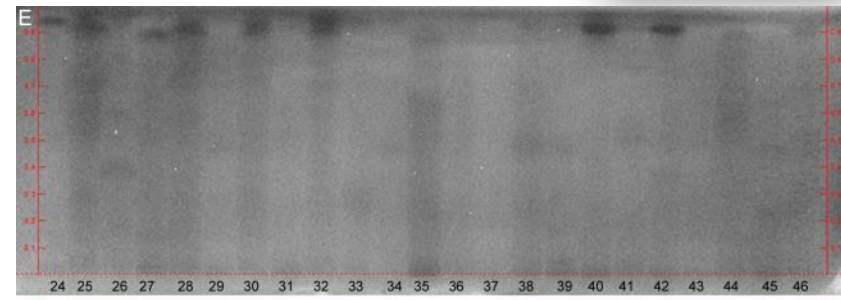
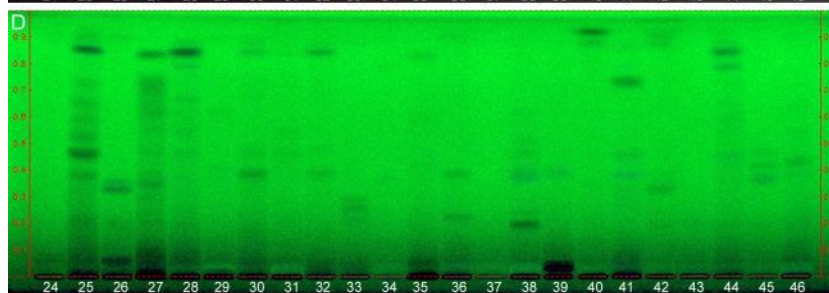
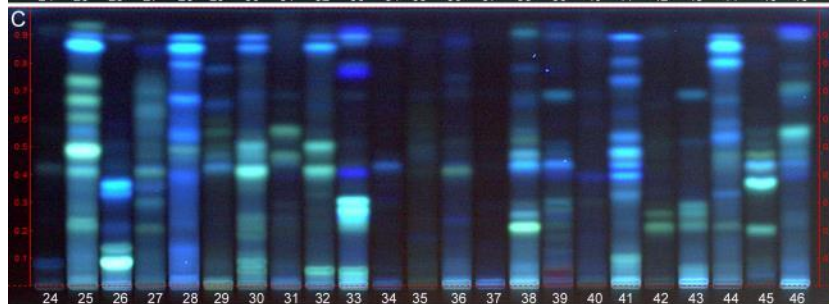
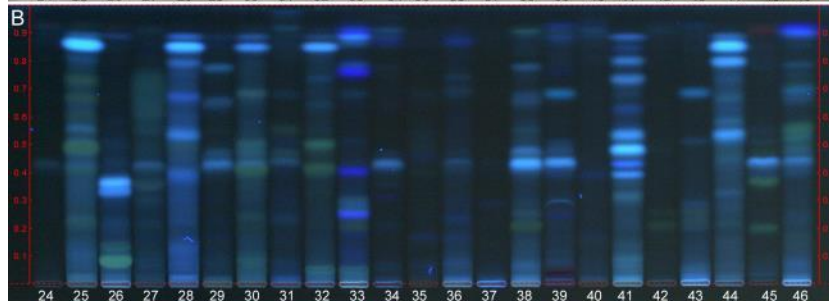
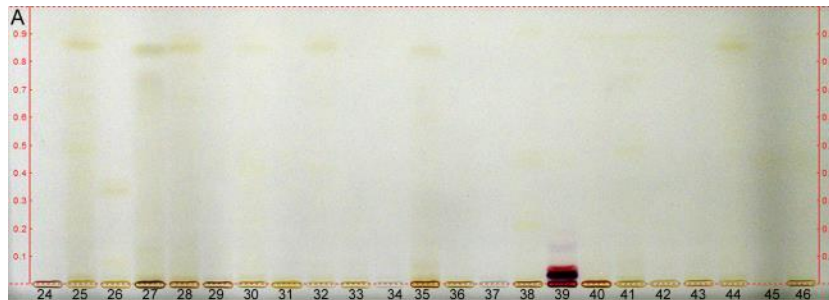
Food Science

G. Morlock

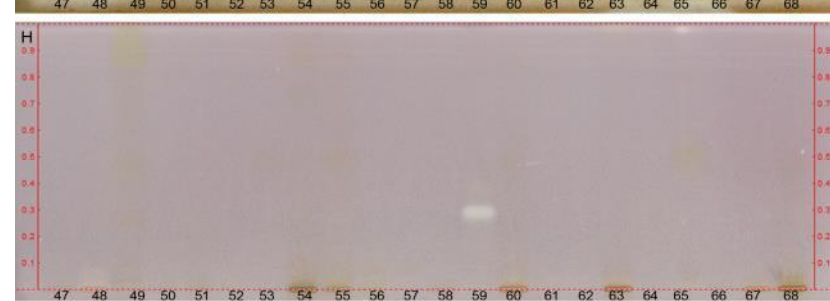
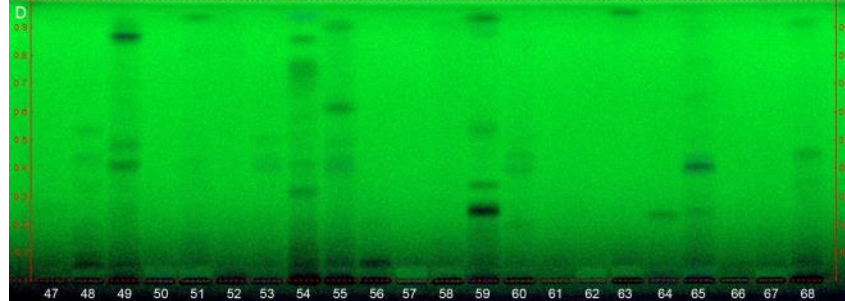
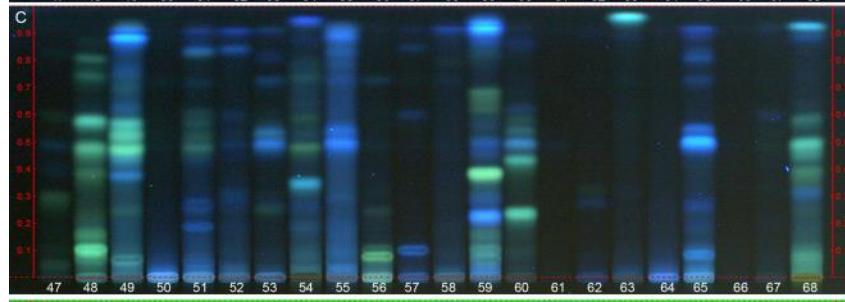
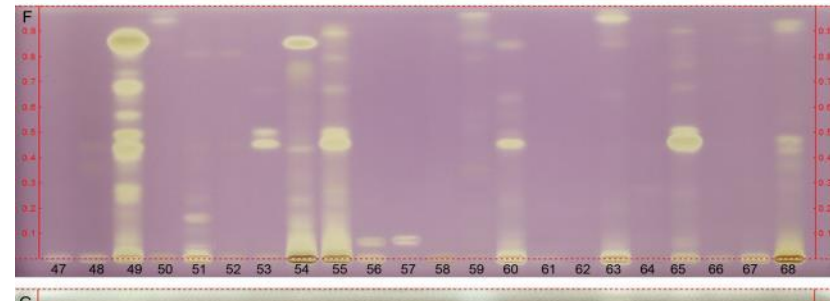
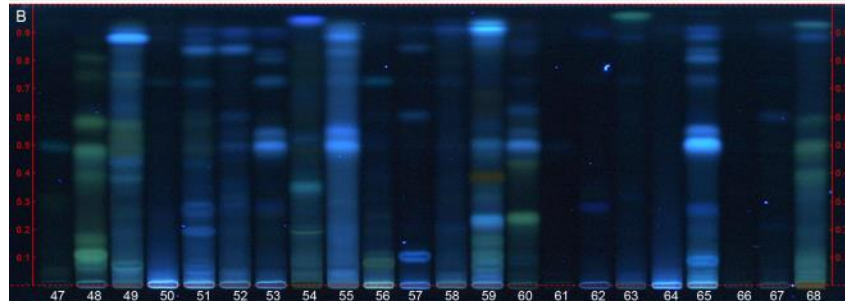
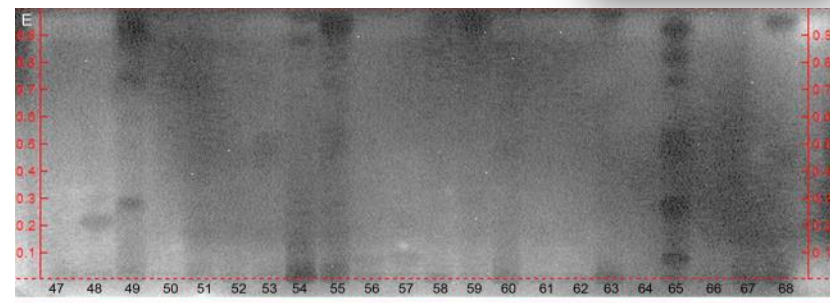
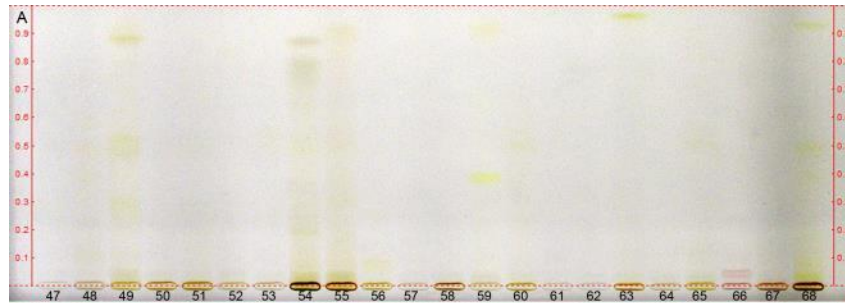




# EDA of 68 botanicals (#24-46)



# EDA of 68 botanicals (#47-68)



# EDA of 68 botanicals

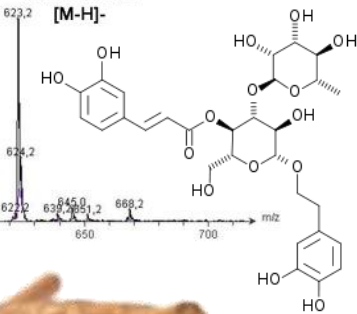
1. Ginkgo	24. Heidelbeere	47. Sanddorn
2. Guarana	25. Rosmarin	48. Bockshornklee
3. Kola	26. <b>Lemon verbena</b>	49. <b>Thyme</b>
4. Pfefferminze	27. Eukalyptus	50. Ingwer
5. Oregano	28. Melisse	51. Andorn
6. Zichorie	29. Kümmel	52. Brennessel
7. Weinblätter	30. Salbei	53. Artischocken
8. Kardamom	31. Hopfen	54. Nelken
9. Zitronenschale	32. Majoran	55. Gerösteter Matete
10. Hagebutte	33. Orangenschale	56. Jasmin
11. Wacholderbeere	34. Liebstöckelwurzel	57. Sonnenhut
12. Honigbusch	35. Traubenkerne	58. Koriander
13. Schafgarbe	36. Fenchel	59. <b>Licorice</b>
14. Passionsblume	37. Ginseng	60. Weißdornextrakt
15. Hibiskus	38. Holunderblüten	61. Apfelschalen
16. Gelber Früchtetee	39. Roter Früchtetee	62. Sellerieknolle
17. Brombeerblätter	40. Zimtrinde	63. Galgant
18. Acerola	41. Schachtelhalm	64. Knoblauch
19. Holunderbeere	42. Sternanis	65. Taigawurzel
20. Grüner Matete	43. Orangenblüten	66. Himbeersaftkonzentrat
21. Spitzwegerich	44. Basilikum	67. Traubenschalen
22. Schwarzes Johannisbeersaftkonzentrat	45. Weißdornblätter	68. Rooibos
23. Weißdornblätter #1	Charge 2	
	46. Kamille	

# Characterization of bioactive compounds

Lemon verbena (*Aloysia citrodora*)



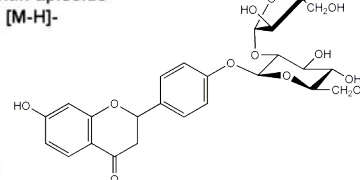
Verbascoside



Licorice (*Glycyrrhiza glabra*)



Liquiritin apioside

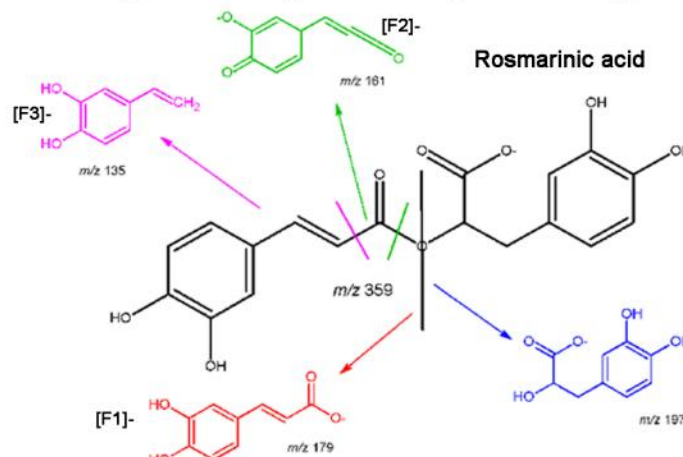


Thyme (*Thymus vulgaris*)



Rosmarinic acid

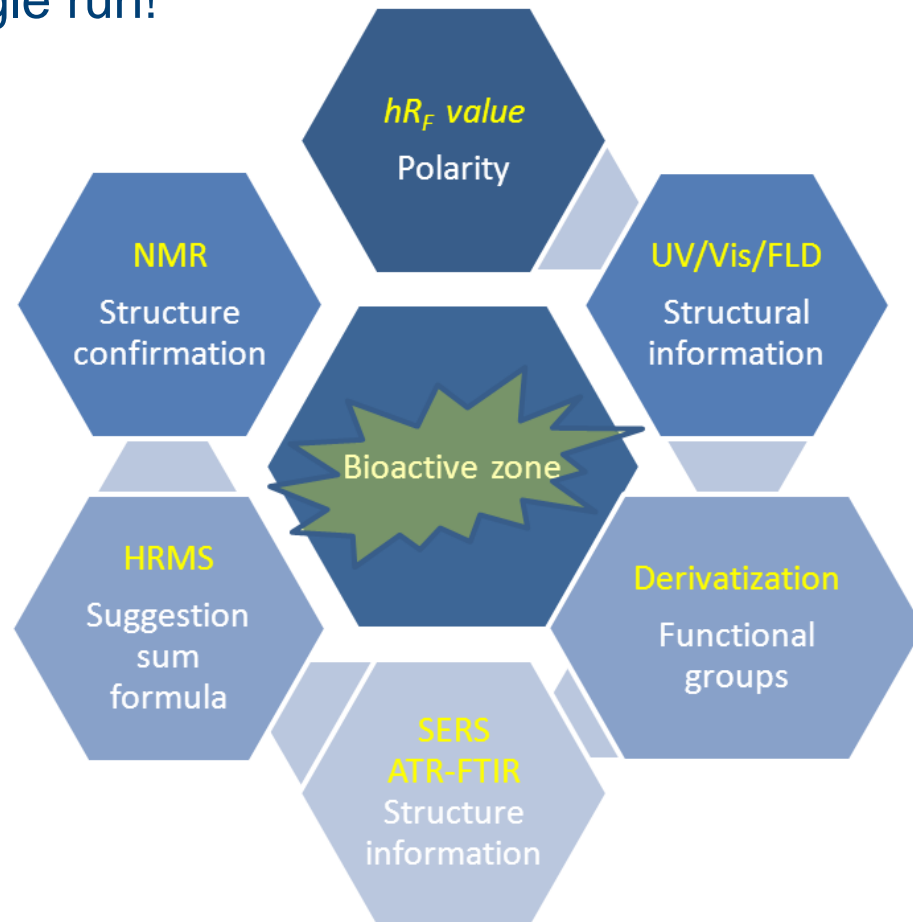
[M-3H+Na]<sup>-</sup>



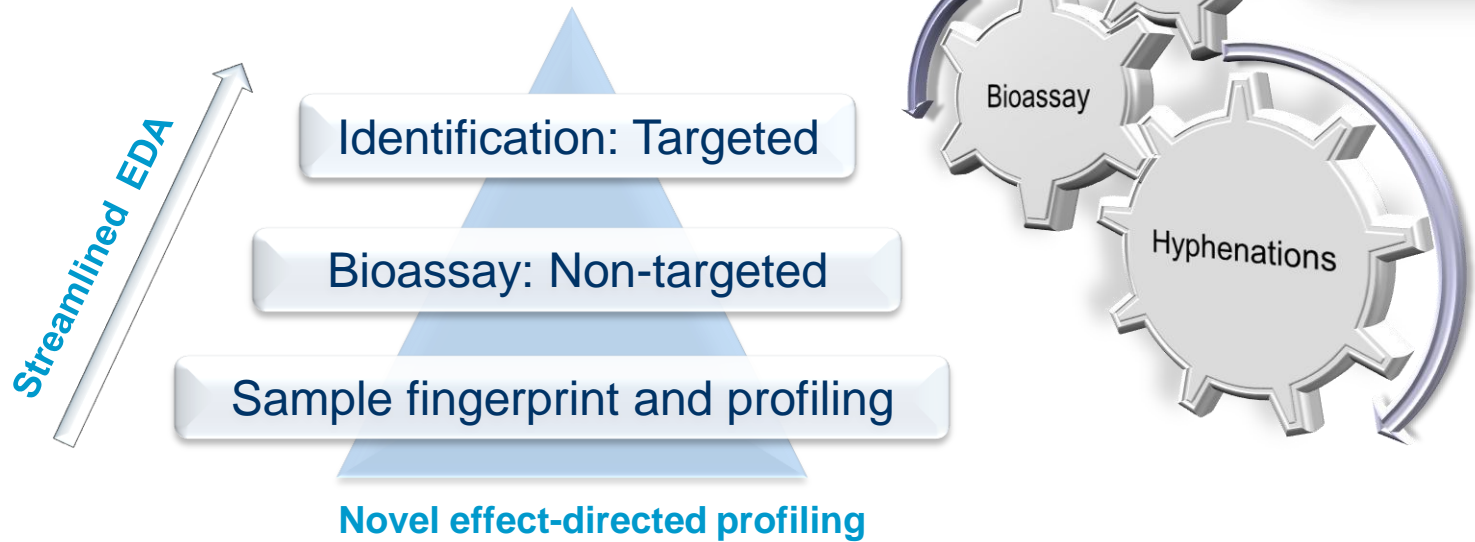
S. Krüger, R. Fornasari, I. Scainelli,  
L. Hüsken, G. Morlock, in preparation

# Goal: From bioactive zone to sum formula

...in a single run!



# Conclusion



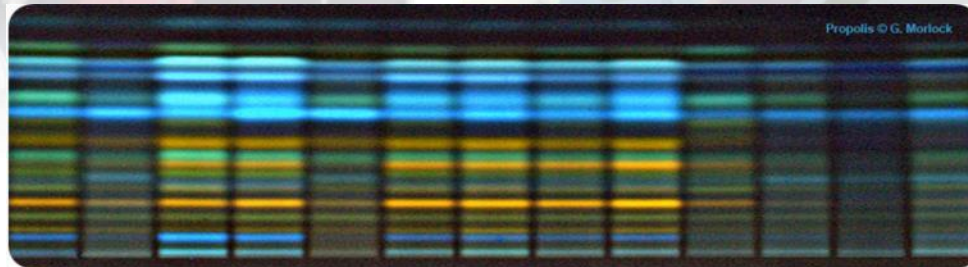
## Comprehensive information

- ➡ Analysing samples as raw as possible
- ➡ Identifying single compounds generating the effect
- ➡ Reduced matrix interference
- ➡ Improved capability of detection
- ➡ Support decisions in natural product search and health issues

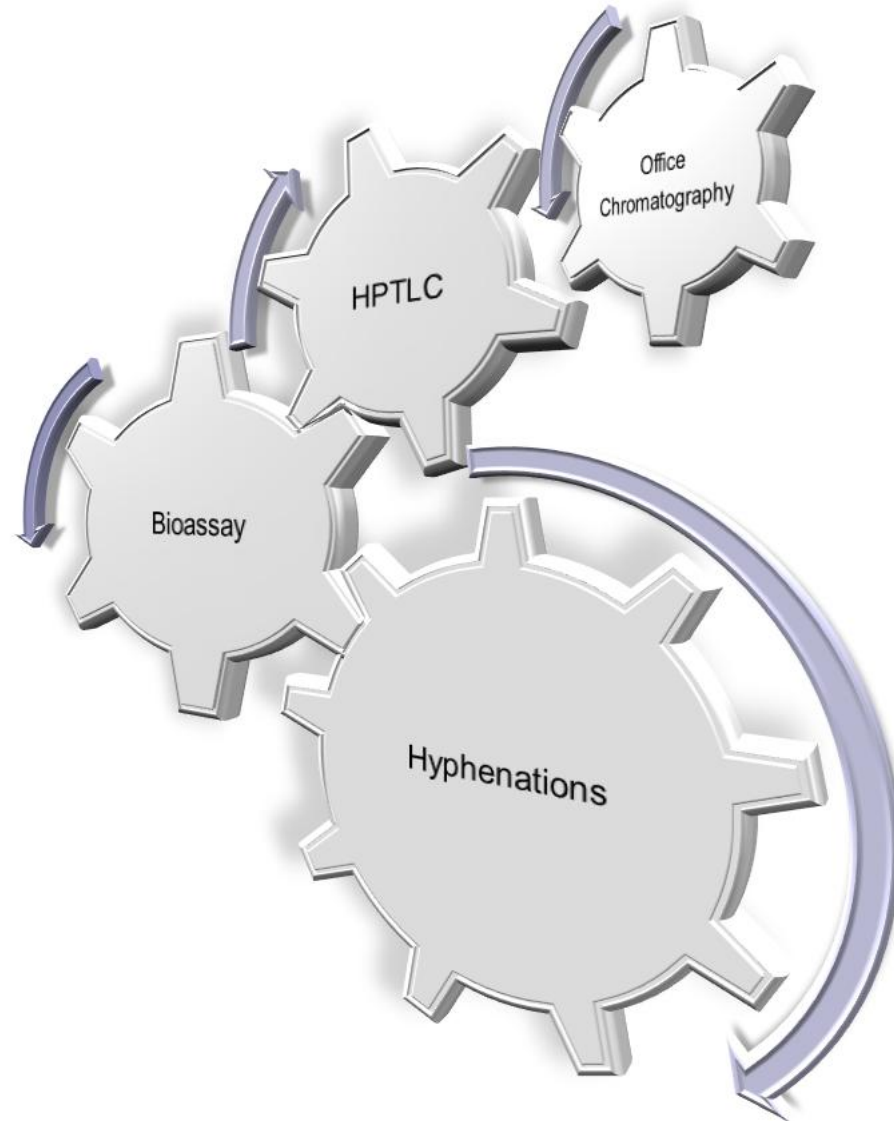


Thank you!

JUSTUS-LIEBIG-  
UNIVERSITÄT  
GIESSEN  
Food Science 

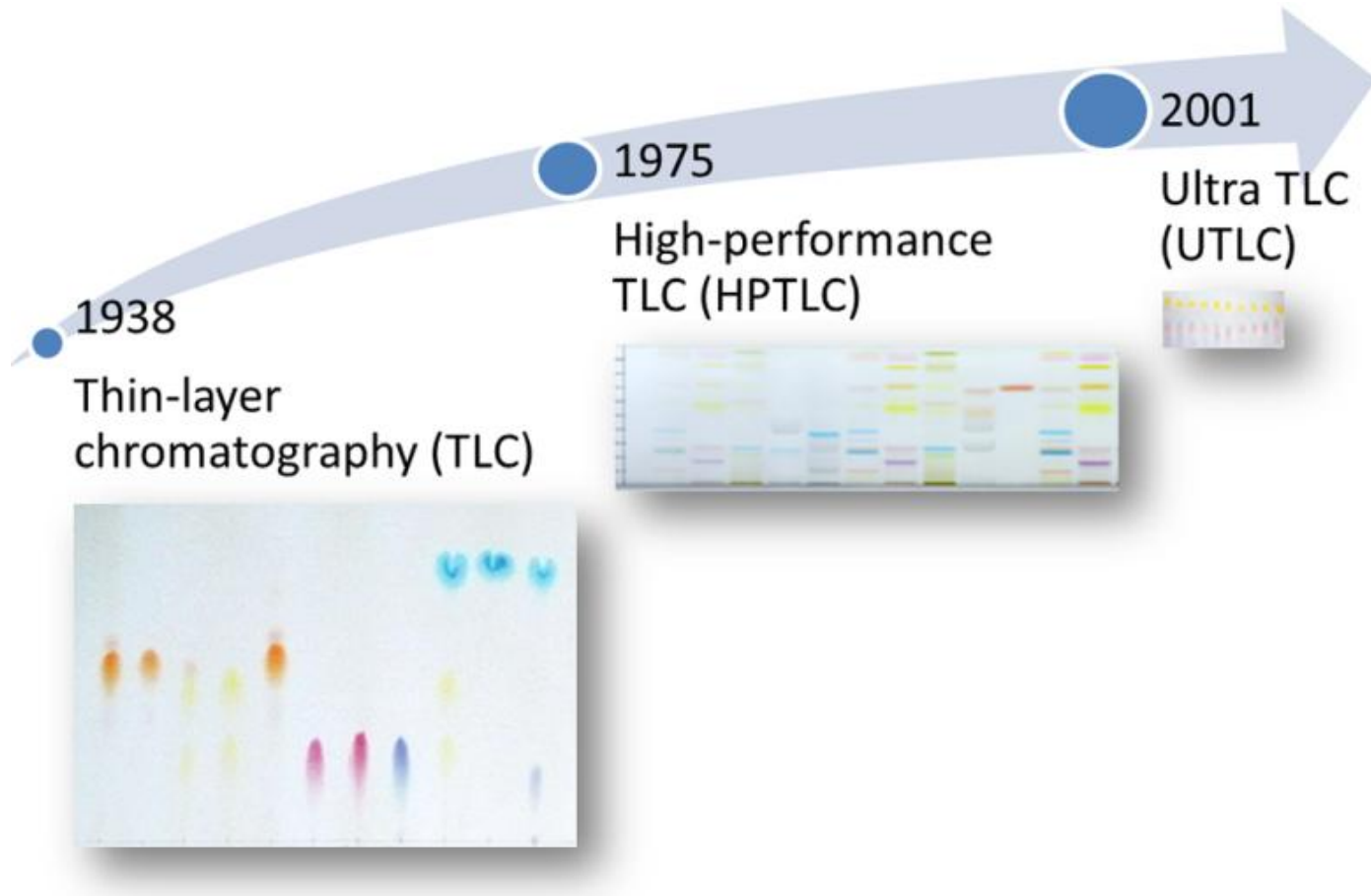


# Outlook

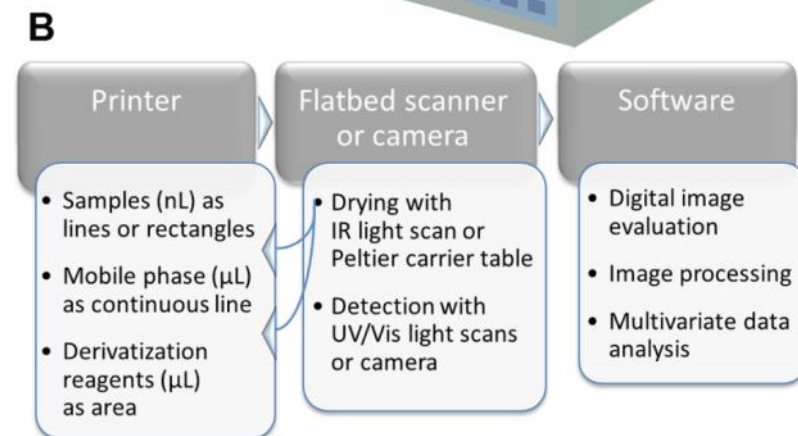
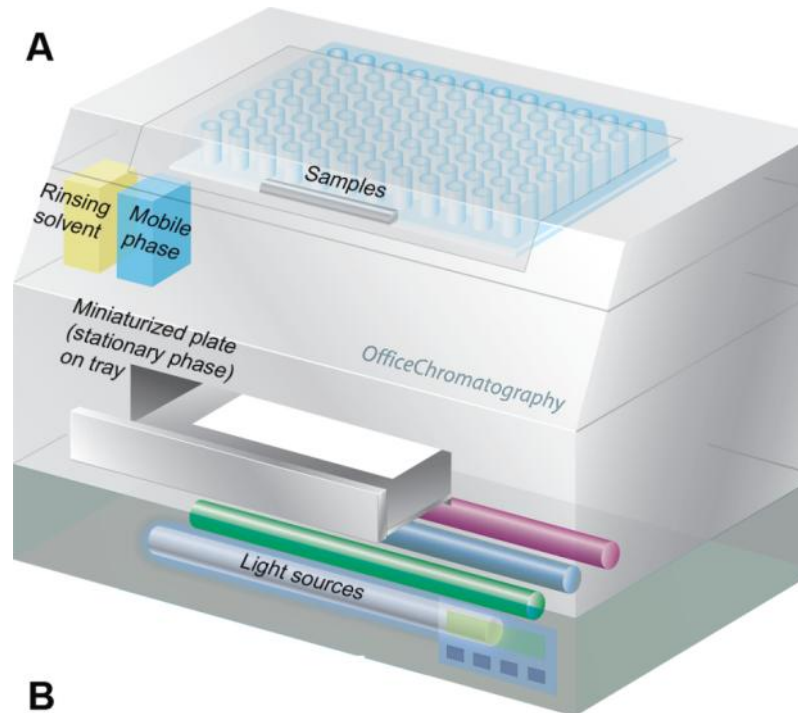




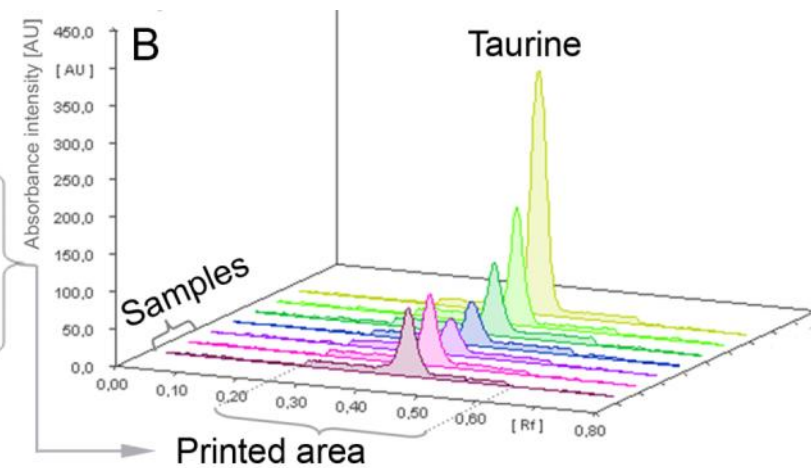
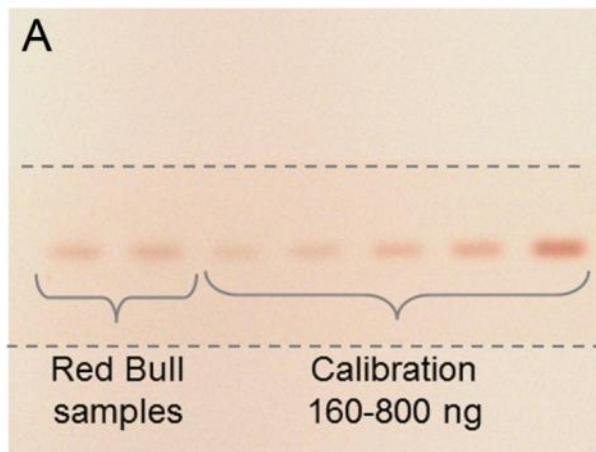
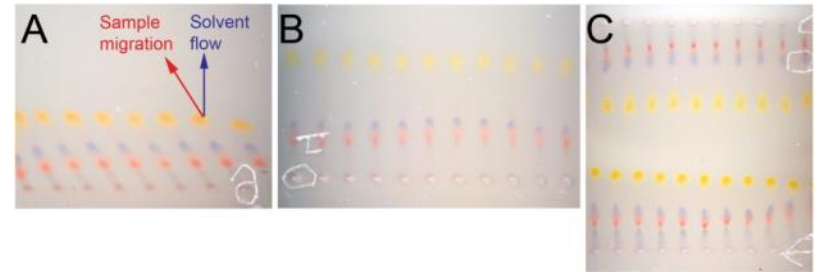
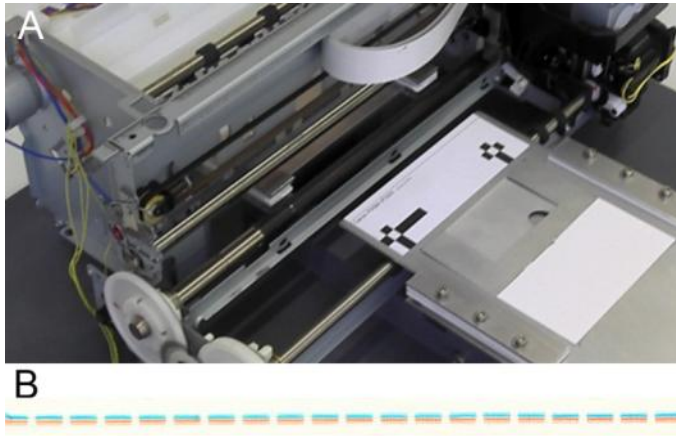
# Miniaturization



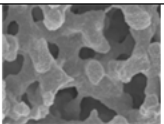
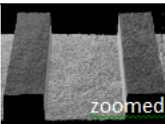
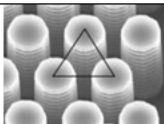
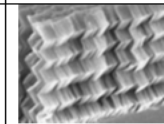
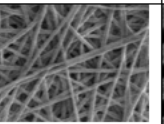

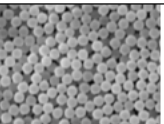
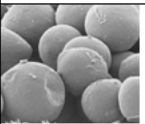
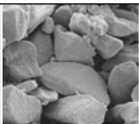
# Office chromatography



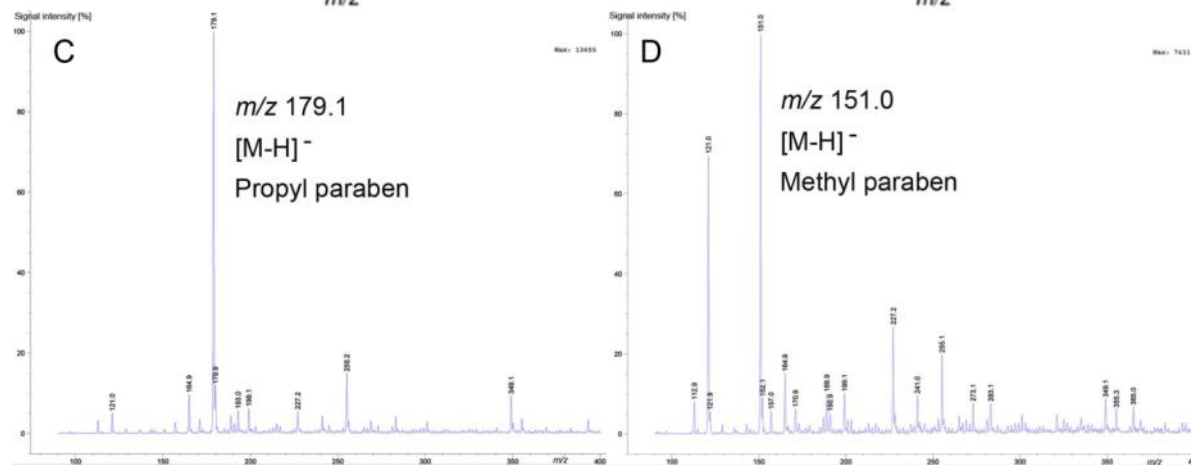
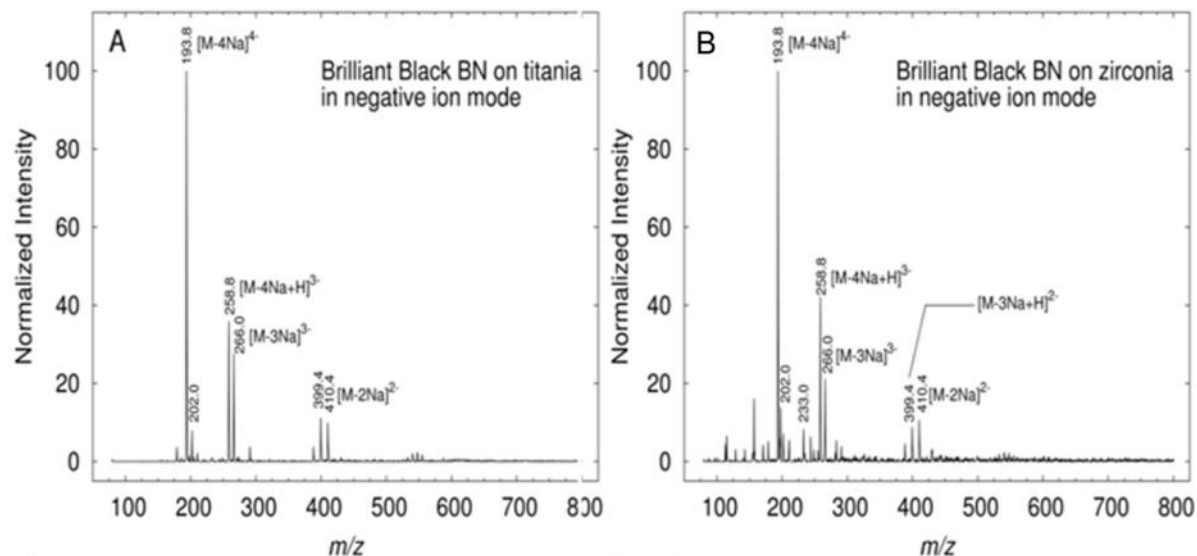
# UTLC & PMT



# Novel layers

	UTLC							HPTLC	
First report	2001 [5]	2001 [47]	2007 [49]	2008 [28]	2009 [56]	2011 [60]	2011 [63]	1975 [3]	
Layer type	Monolithic layer	Monolayer on channel bottom	Ordered (non)porous pillar arrays	Nanostructured layer	Electrospun mat	Carbon-nanotube-templated microfabrication (CNT-M)	Submicrometer particulate layer with cross linked polymer brushes	Particulate layer	
Technique of fabrication	Polymerization on glass plate (sol-gel process) and opt. photografting	DRIE of Si-wafer surface and coating	Mid/Deep-UV lithography, DRIE of Si-wafer surface and coating	GLAD of inorganic oxides on glass plates	Electrospinning of (composite) polymer solutions on aluminium foil	Coating CNTs with silica by (pseudo) atomic layer deposition plus second coating	Slurry overlay on Si-wafer and brush coating by polymerization	Slurry overlay on various carriers (glass plate, aluminium or polymer foil) and coatings	
Layer icon									
Layer structure	Monolithic texture with 1-2 $\mu\text{m}$ macropores	Monolayered porous silicon bottom of a nanochannel (0.7 $\mu\text{m}$ wide, 0.3 $\mu\text{m}$ deep)	Monolayer coating or monolithic silica shell of cylindrical pillars ( $\varnothing$ 4 $\mu\text{m}$ , 10 $\mu\text{m}$ high, spaced 0.3 - 1.7 $\mu\text{m}$ ) in a 70 $\mu\text{m}$ nanochannel	Column array of verticals, posts, helices, zig-zags or blades (spaced 2 - 50 nm) with (an)isotropic structure	Spun mat of nanofibers ( $\varnothing$ 200-400 nm, cm to m long) forming cylindrical channels	Silica coated (20-60 nm) herring-bone hedge array (3-4 $\mu\text{m}$ wide, spaced 4-7 $\mu\text{m}$ ) forming channels (50-100 $\mu\text{m}$ long)	Non-porous particles coated with a polymer brush layer	Spherical	Irregular
Layer thicknesses ( $\mu\text{m}$ )	10 - 50	Monolayer 0.05 - 0.3	Monolayer or 0.5 $\mu\text{m}$ porous shell	1.3 - 7	15 - 25	50	15	50 - 200	
Adsorbent types	Silica gel, poly-(4-methylstyrene-co-chloromethylstyrene-co-divinylbenzene)	C8, C18	C8; C18 [50]	Silica, zirconia alumina, titania, C18	Glassy carbon, polyvinyl alcohol, polyacrylonitrile	Silica, amino	Polyacrylamide; poly(GMA-co-DEGDMA-NH <sub>2</sub> ) [64]	Silica, amino, cyano, diol, C2, C8, C18, cellulose, etc.	
Layer geometry (mm)	60 x 36 or 30 x 33	0.7 x 20	0.14 x 40; 10 x 30 [50]	25 x 25 or 100 x 20	30 x 60, individually sliceable	12 x 60	25 x 25	200 x 100, individually sliceable	

# Mass spectra



# Bioassay printing

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