



University of Hohenheim  
Institute of Food Chemistry

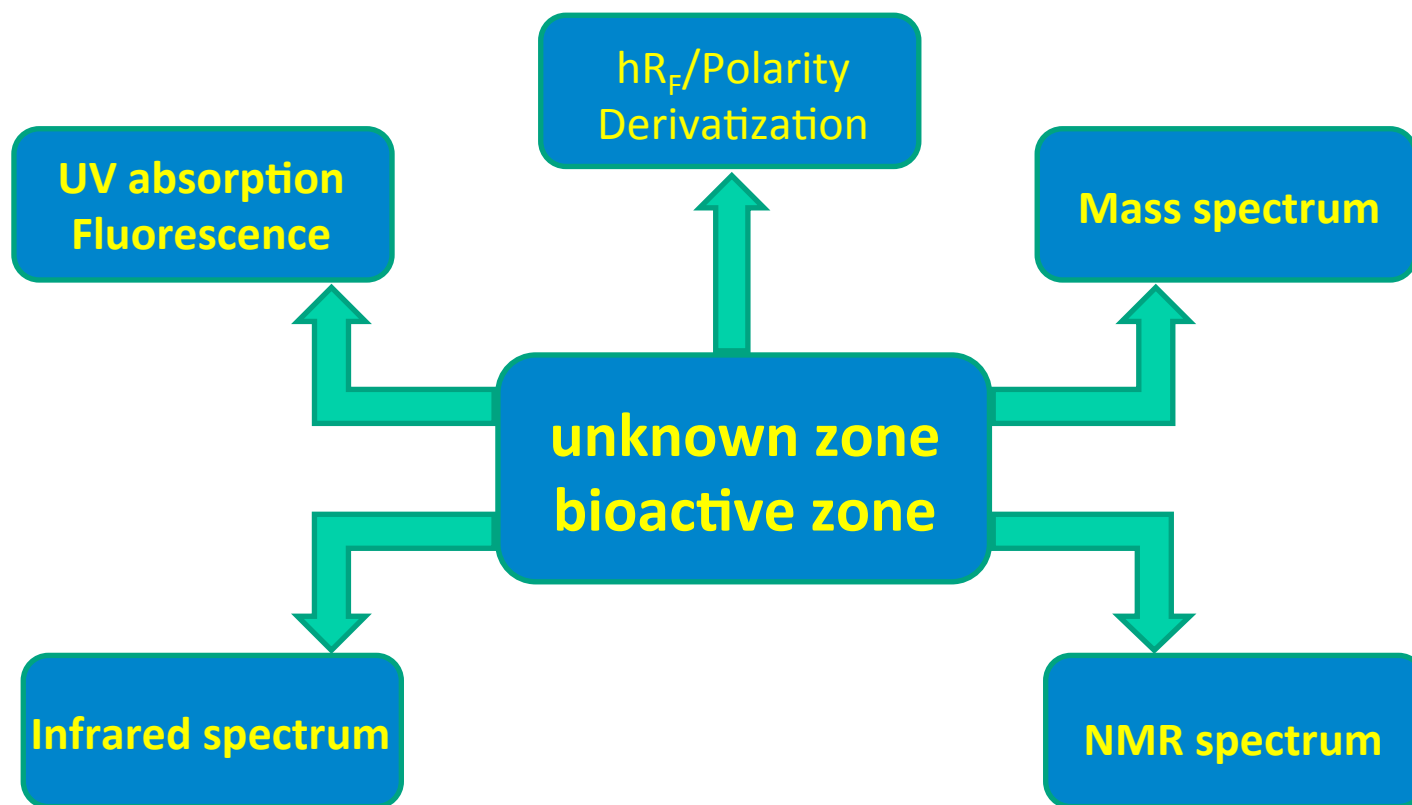
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# HPTLC hyphenations – potential for structure elucidation

Wolfgang Schwack  
University of Hohenheim



# From a detected zone to the chemical structure





# Derivatization/Staining

**Iodine:** olefins, aromatics

**Fluram:** amines, amino acids

**Ninhydrine:**  $\alpha$ -amino acids, peptides

**2,4-Dinitrophenylhydrazine:** aldehydes, ketones

**Primuline:** lipids

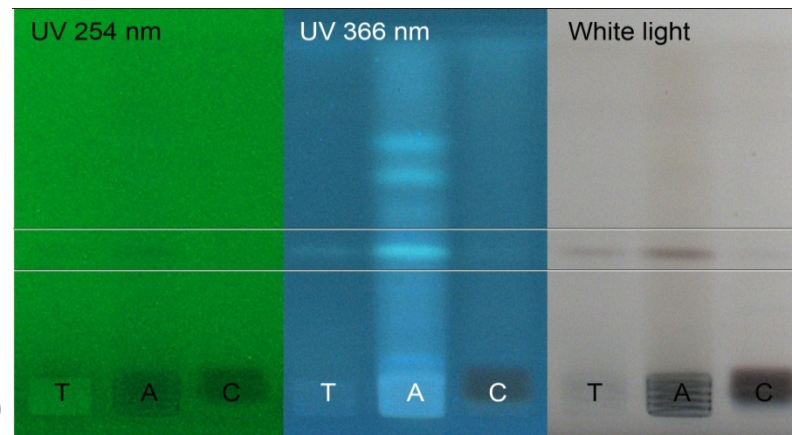
**$\beta$ -Naphthol:** glycosides

**p-Aminobenzoic acid:** glycosides

...

Sucralose  $\rightarrow$

G. Morlock et al. (2011)

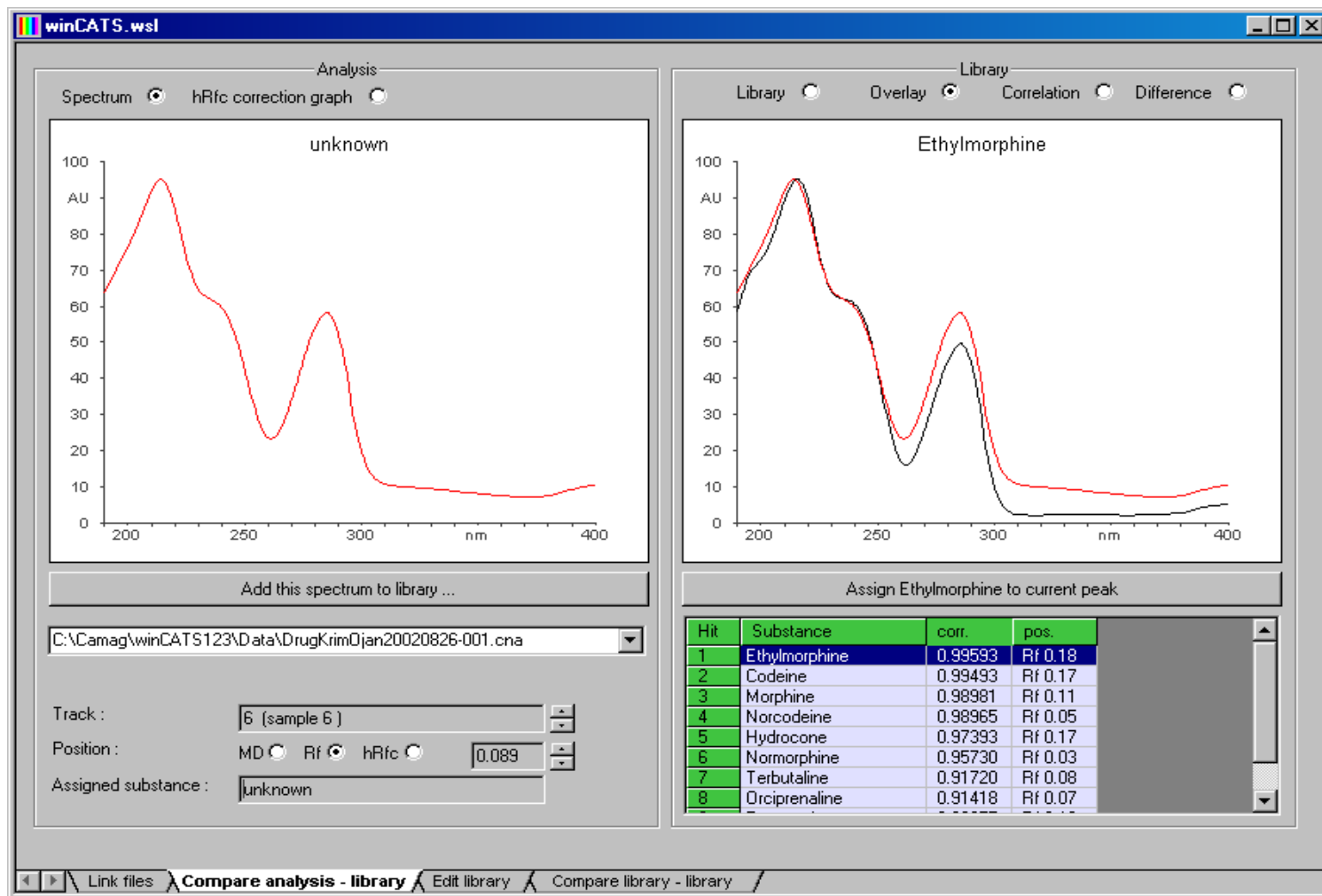


Fluram



# UV/Vis spectrum

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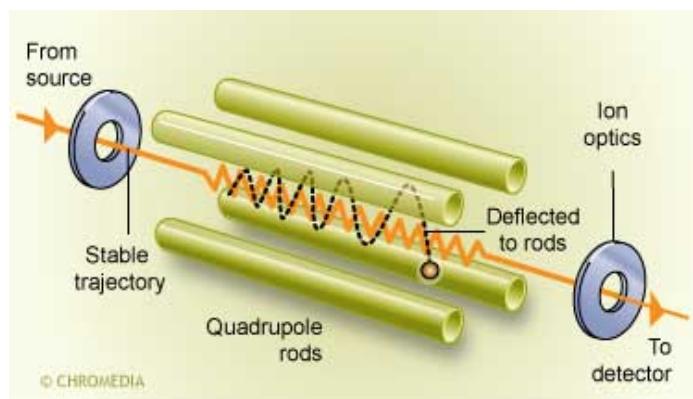
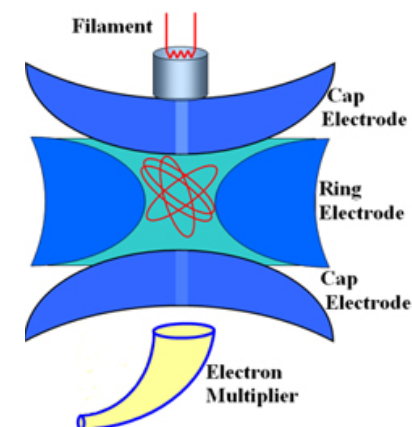




# Mass spectrum

## Low resolution mass spectrometer (nominal mass)

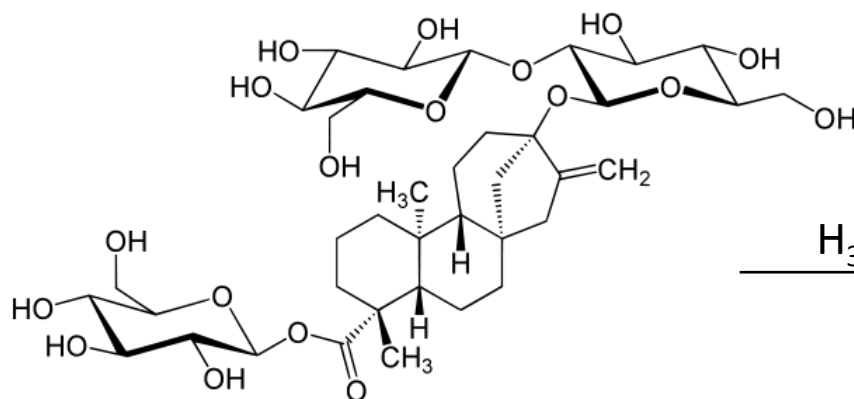
- ☞ Some structural information is available:
- Degradation/stability studies (parent is known)
  - Organic synthesis or derivatization (parents are known)
  - Identification of a bioactive compound in plant X (natural composition of X is well known)





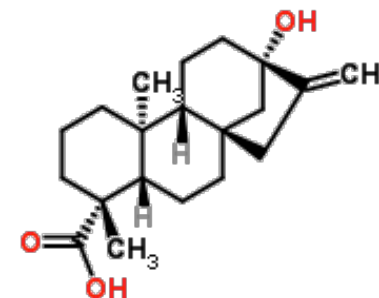
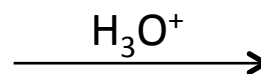
# Mass spectrum

Example: *Stevia rebaudiana*



Stevioside:  $[M+H]^+ = m/z$  805

☞ Sugar reagents: positive



Steviol:  $[M+H]^+ = m/z$  319

$[M-H]^- = m/z$  317

☞ Less polar

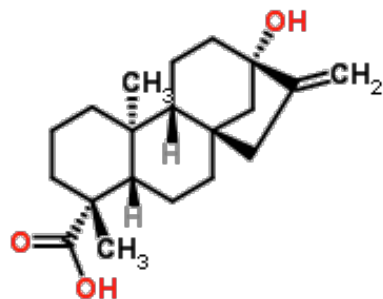
☞ Sugar reagents: negative

☞ Primuline: positive



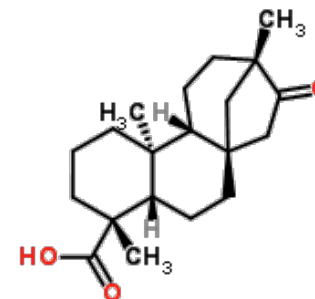
# Mass spectrum

However:



Steviol

or



iso-Steviol

☞ IR spectrum,  $^1\text{H-NMR}$  spectrum ?



# Mass spectrum

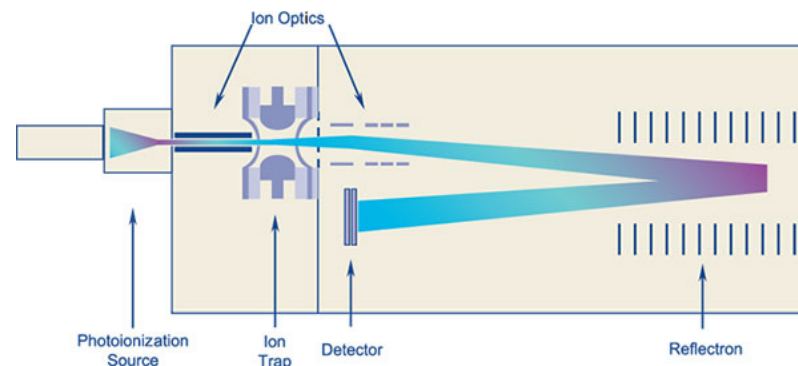
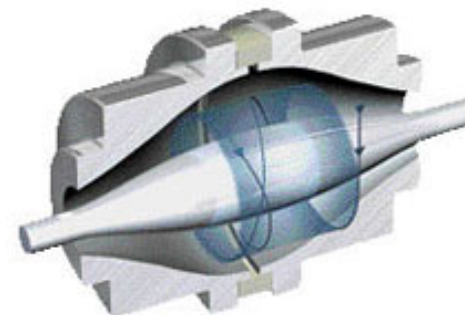
## High resolution mass spectrometer (exact mass)

☞ No structural information is available:

$$[M+H]^+ = m/z \ 319.22677$$

☞ ChemSpider search ([www.chemspider.com](http://www.chemspider.com)): compound known?

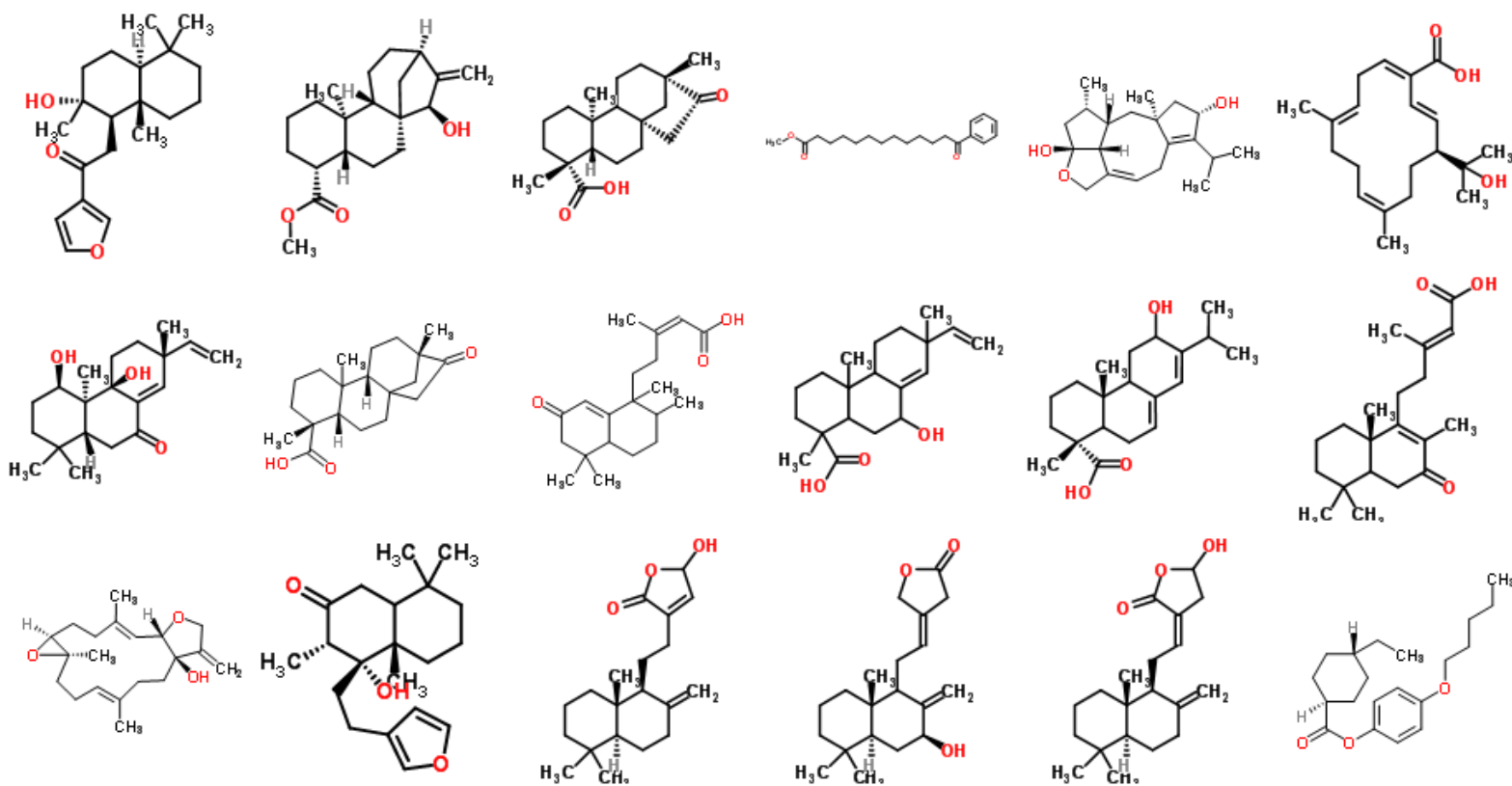
- Monoisotopic mass  $\pm 0.00032$  (1 ppm)  
=> 1060 hits (all with  $C_{20}H_{30}O_3$ )
- Monoisotopic mass  $\pm 0.0032$  (10 ppm)  
=> 1800 hits





# Mass spectrum

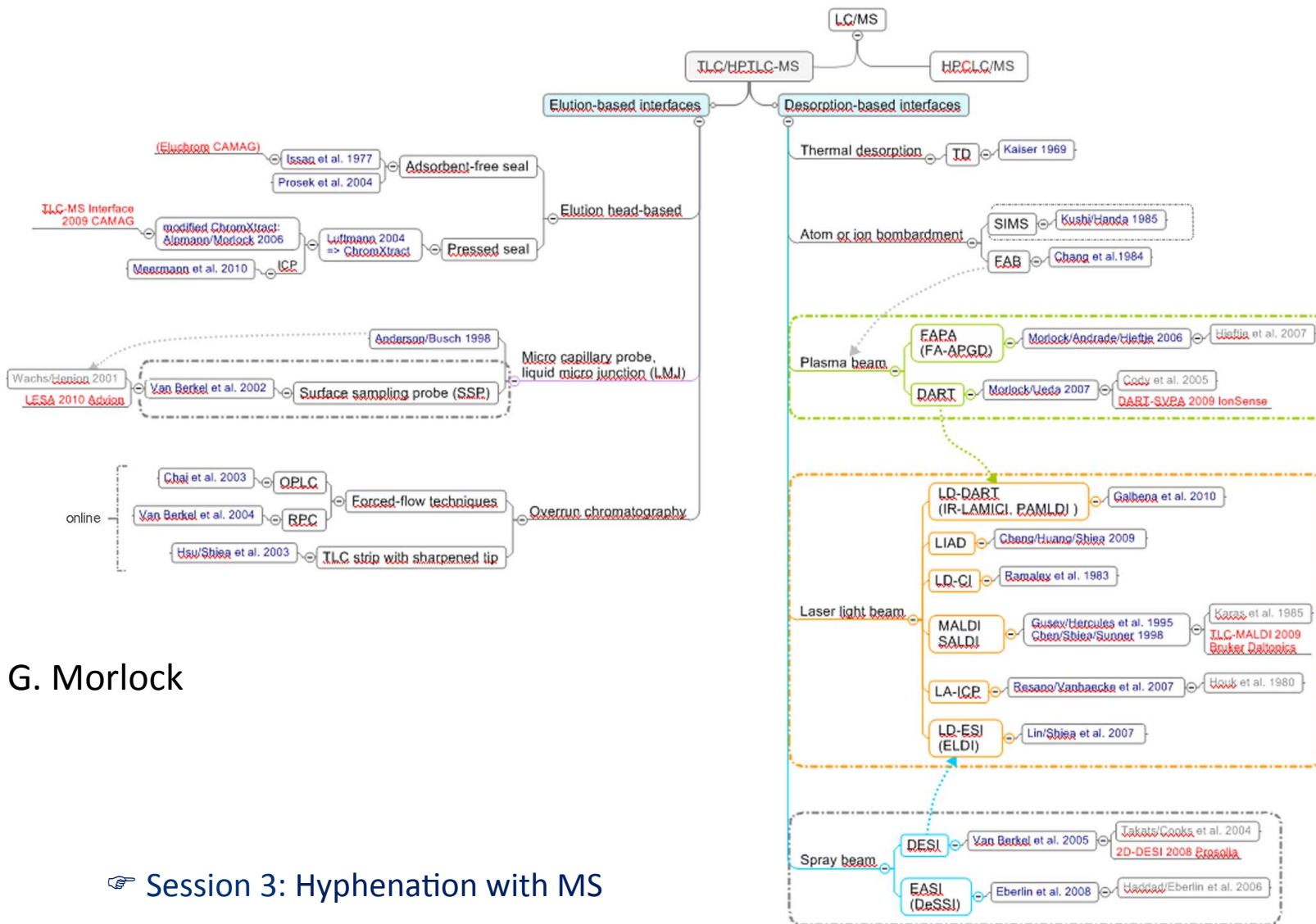
Just a selection ...



 **Additional structural information (filter)?**



# Interfacing HPTLC → MS



G. Morlock



# Interfacing HPTLC → MS

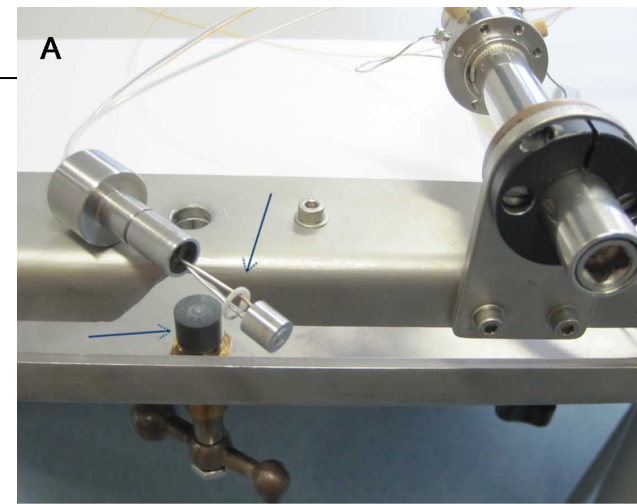
## 1. Elution-based TLC-MS

### Pressed seal

- ☞ TLC-MS Interface (CAMAG, 2009)
- ☞ ChromXtract (Luftmann, 2004)
- ☞ ChromXtract modified (Alpmann & Morlock, 2006)

### Liquid microjunction (LMJ)

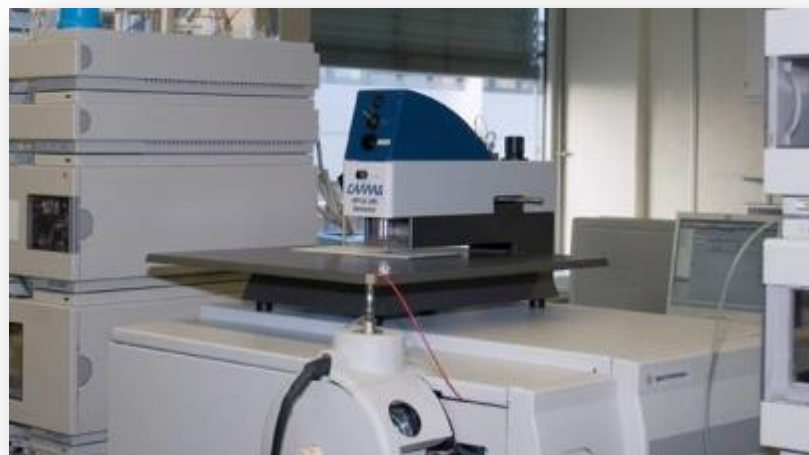
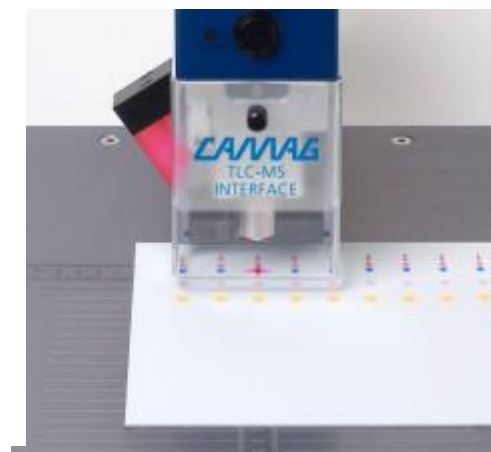
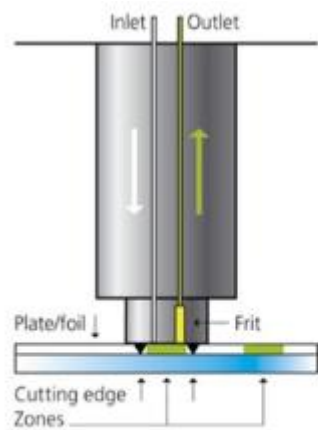
- ☞ In situ microextraction (flowprobe™): Prosofia (2008)
- ☞ Liquid Extraction Surface Analysis (LESA®): Advion (2010)
- ☞ TLC-surface sampling probe (SSP)(Van Berkel et al., 2002)







# TLC-MS Interface

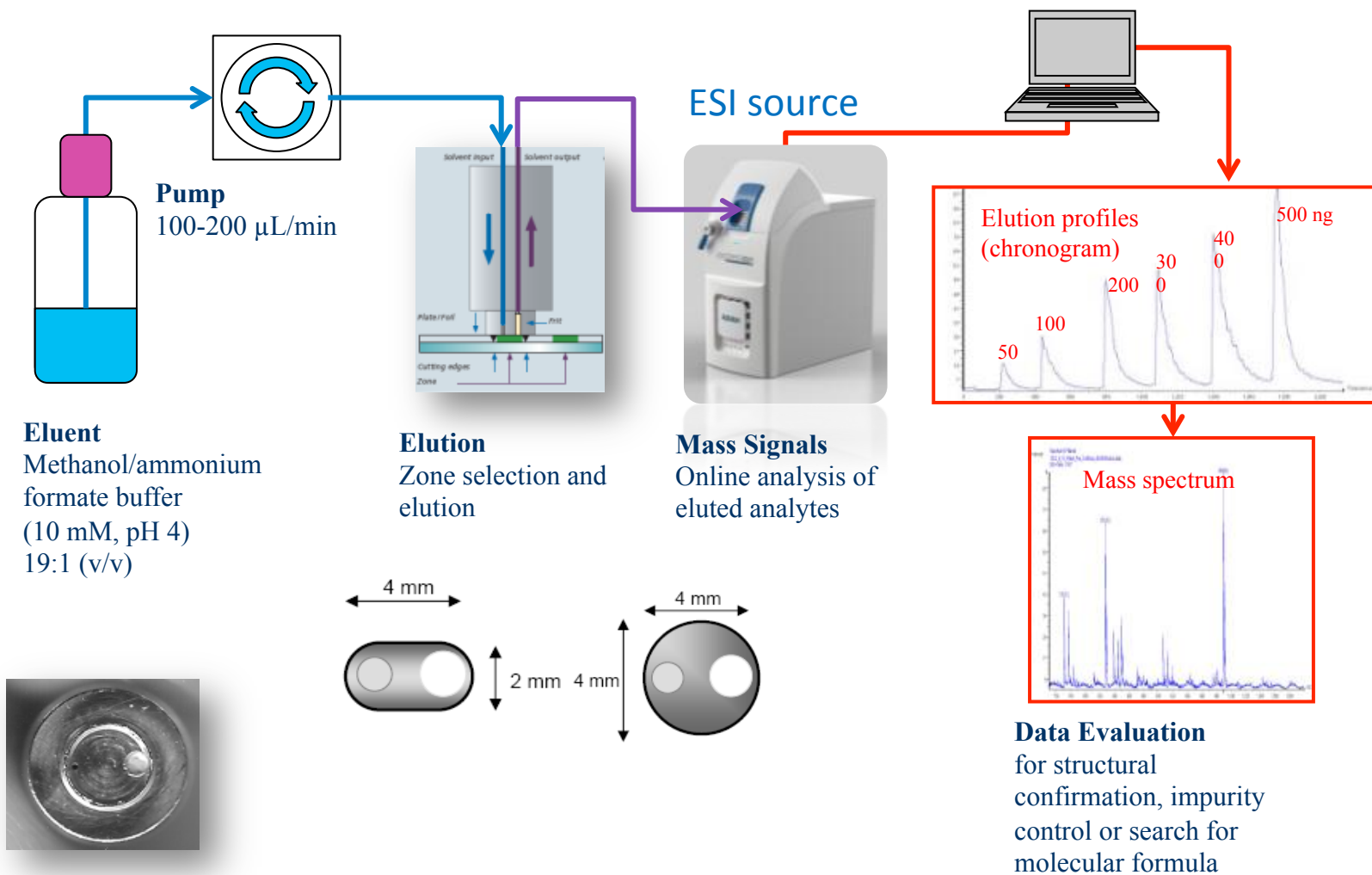






# TLC-MS Interface

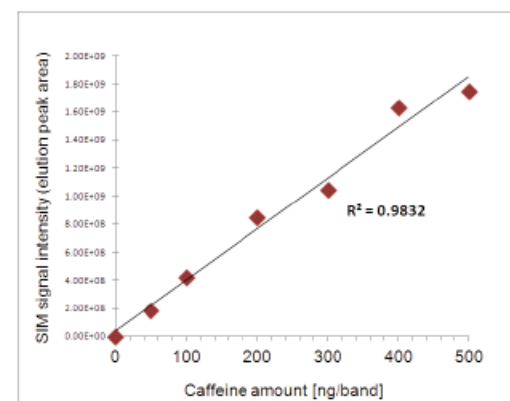
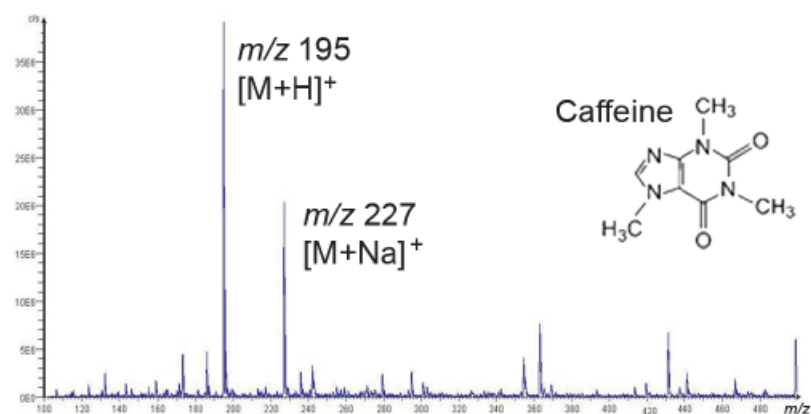
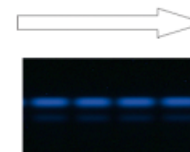
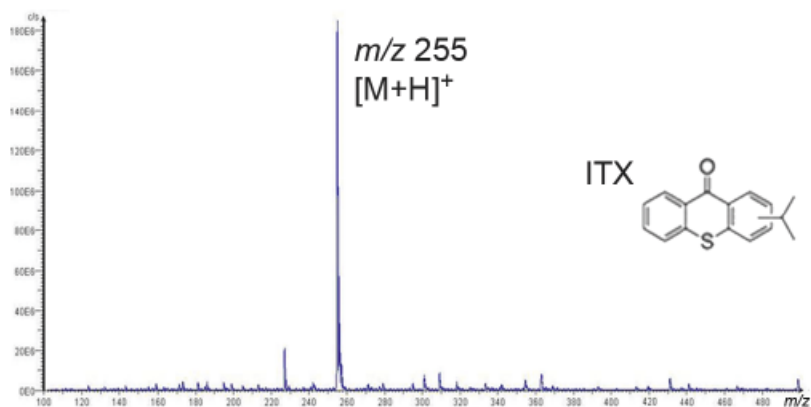
## How it works





# TLC-MS Interface

## Performance



Up: Mass spectrum of ITX and its repeatability in the SIM mode (%RSD = 4 %)

Down: Mass spectrum of caffeine and its analytical response in the SIM mode ( $R^2 = 0.9832$ )



# Mass spectrometers/interfaces

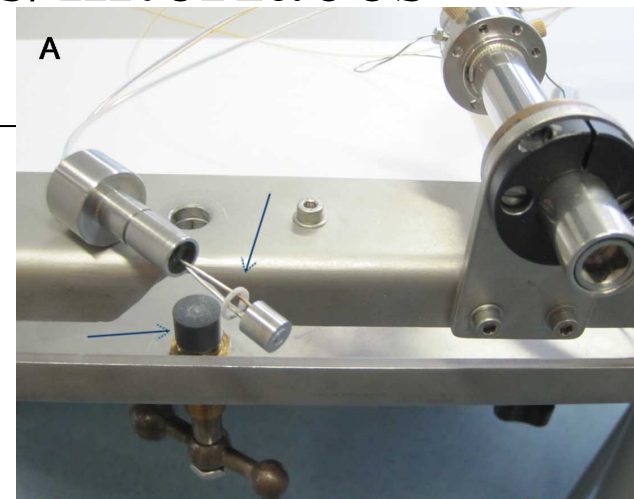
## 1. Elution-based TLC-MS

### Pressed seal

- ☞ ChromXtract (Luftmann, 2004)
- ☞ ChromXtract modified (Alpmann & Morlock, 2006)
- ☞ TLC-MS Interface (CAMAG, 2009)

### Liquid microjunction (LMJ)

- ☞ In situ microextraction (flowprobe™): Prosofia (2008)
- ☞ Liquid Extraction Surface Analysis (LESA®): Advion (2010)
- ☞ TLC-surface sampling probe (SSP)(Van Berkel et al., 2002)

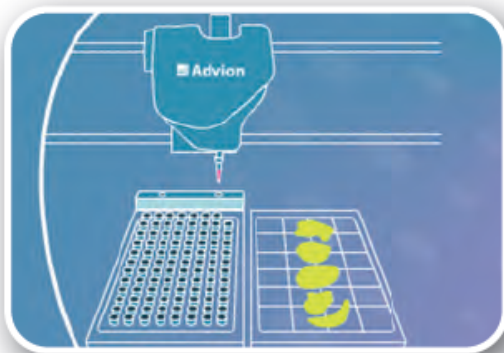




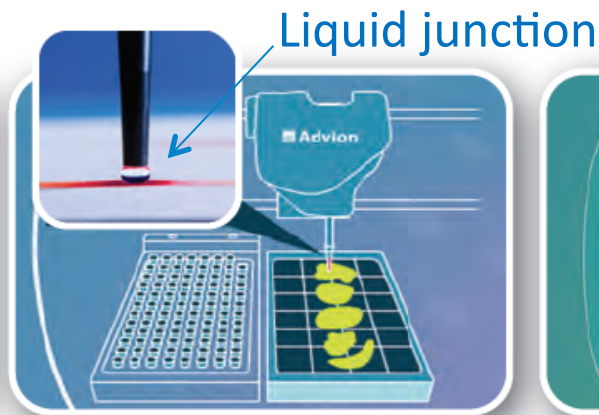
# Liquid microjunction

## Liquid extraction surface analysis: LESA-TriVersa Nanomate (Advion)

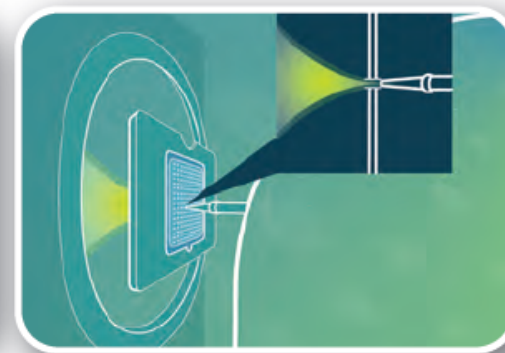
### How It Works



The TriVersa NanoMate picks up a pipette tip from the tip rack, then aspirates extraction solvent from the reservoir.



The robot brings the extraction solvent into contact with the surface of the sample. The analyte is extracted from the surface.



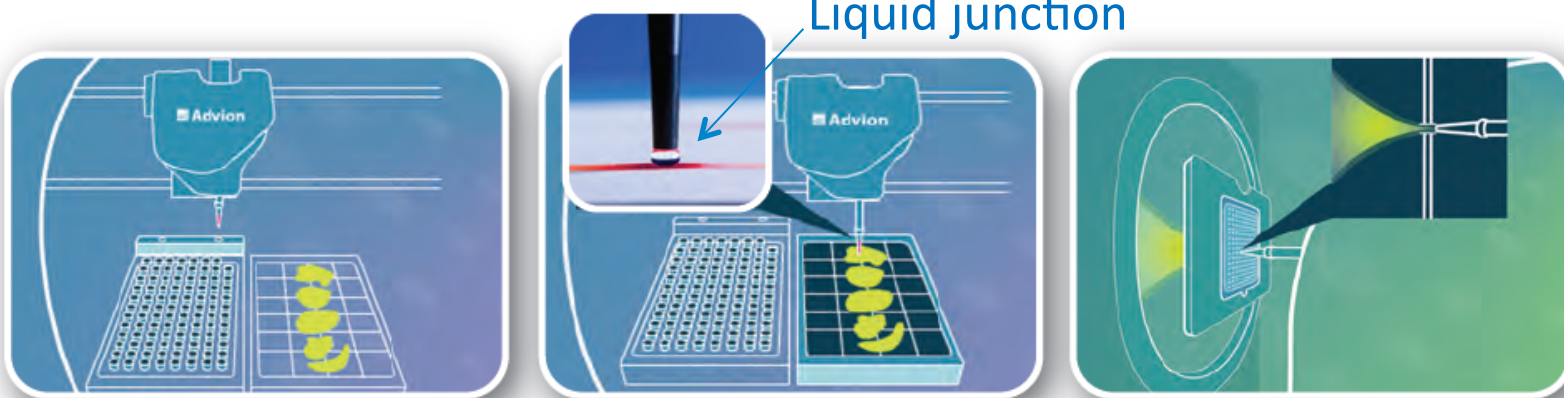
The solvent is retracted into the pipette tip and is analyzed by chip-based infusion.



# Liquid microjunction

## Liquid extraction surface analysis: LESA-TriVersa Nanomate (Advion)

### How It Works



The TriVersa NanoMate picks up a pipette tip from the tip rack, then aspirates extraction solvent from the reservoir.

The robot brings the extraction solvent into contact with the surface of the sample. The analyte is extracted from the surface.

The solvent is retracted into the pipette tip and is analyzed by chip-based infusion.

 **RP plates only!**

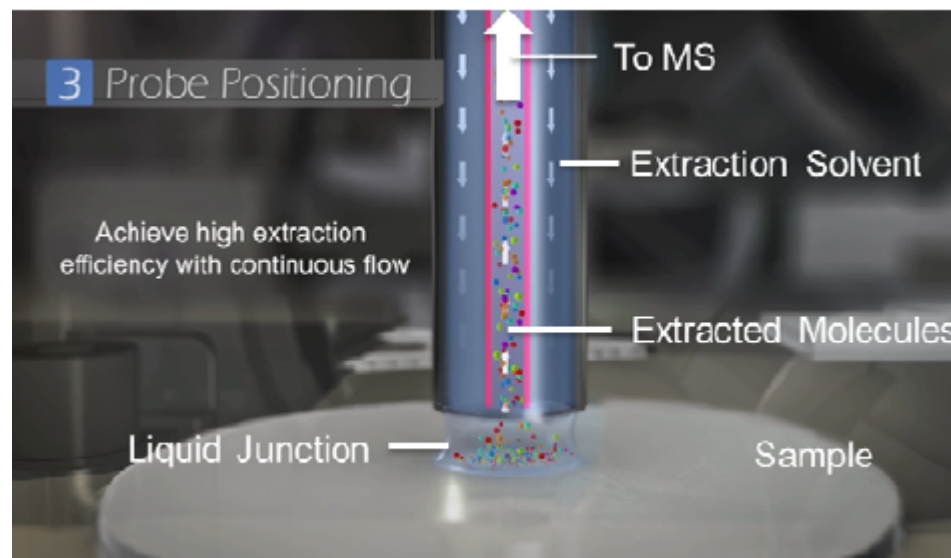


# Liquid microjunction

## In situ microextraction (flowprobe™): Prosolia

### How it works:

The flowprobe uses a two step process: 1) dissolution of the material at the surface by a **continuous flow** of solvent; and 2) ionization of the analyte(s) via electrospray (ESI) ionization.

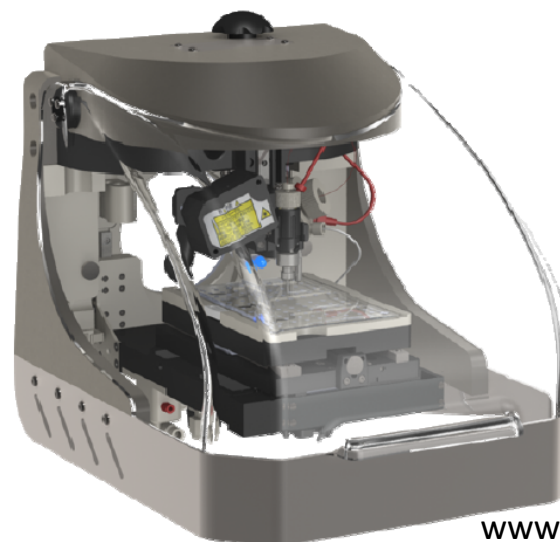






# Liquid microjunction

In situ microextraction (flowprobe™): Prosolia



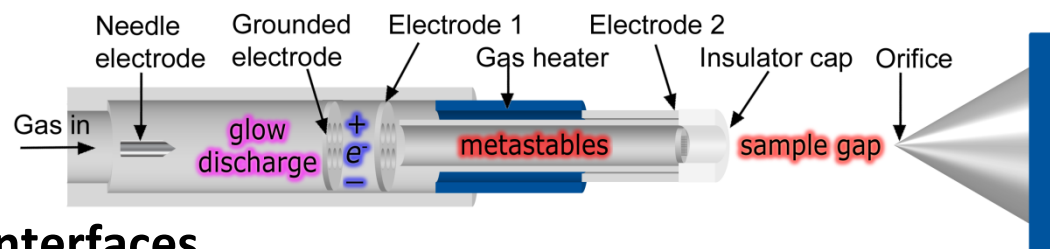
[www.prosolia.com](http://www.prosolia.com)

- ➡ RP plates directly (extraction with methanol/water)
- ➡ NP plates sprayed with silicone oil after development <sup>1)</sup>
- ➡ Line scanning option!

1) M. J. Walworth et al., Anal. Chem. (2011)



# Interfacing HPTLC → MS



## 2. Desorption-based interfaces

Direct Analysis in Real Time (DART®): Ion Sense (2005)

☞ DART-TLC (Morlock & Ueda, 2007)

☞ DART SVP 45A – 3+D Scanner (Ion Sense, 2009)

Matrix Assisted Laser Desorption and Ionization (MALDI):

☞ TLC-MALDI (Bruker Daltonics, 2009)

Desorption Electrospray Ionization (DESI):

☞ TLC-DESI (Van Berkel et al., 2005; Pasilis et al., 2007)

☞ Omnispray® ion sources (2D-DESI): Prosolia (2008)



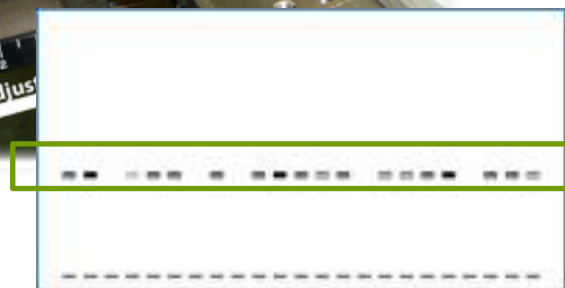
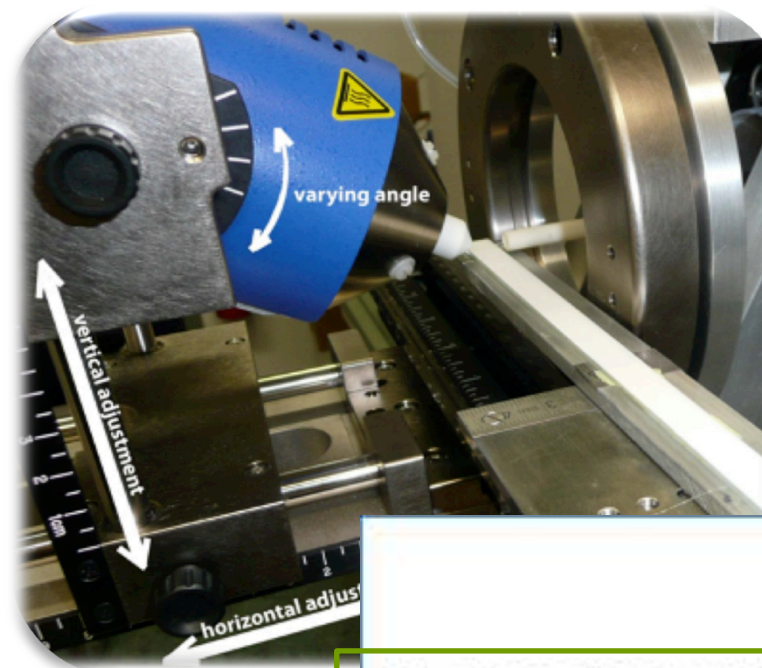
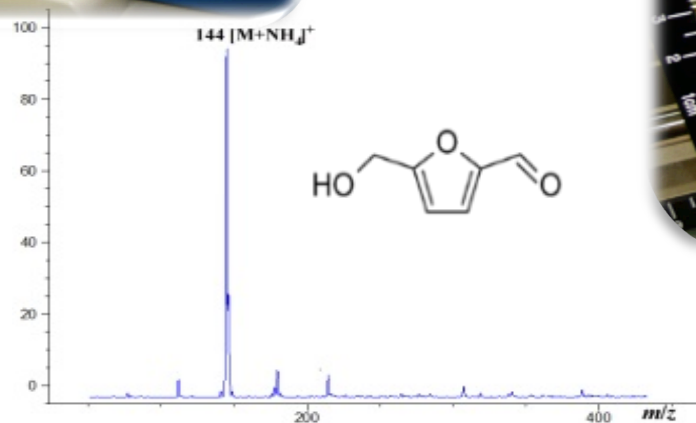


# Desorption-based interfaces

DART 100 → DART SVP 45A – 3+D Scanner



2006 ↔ 2011

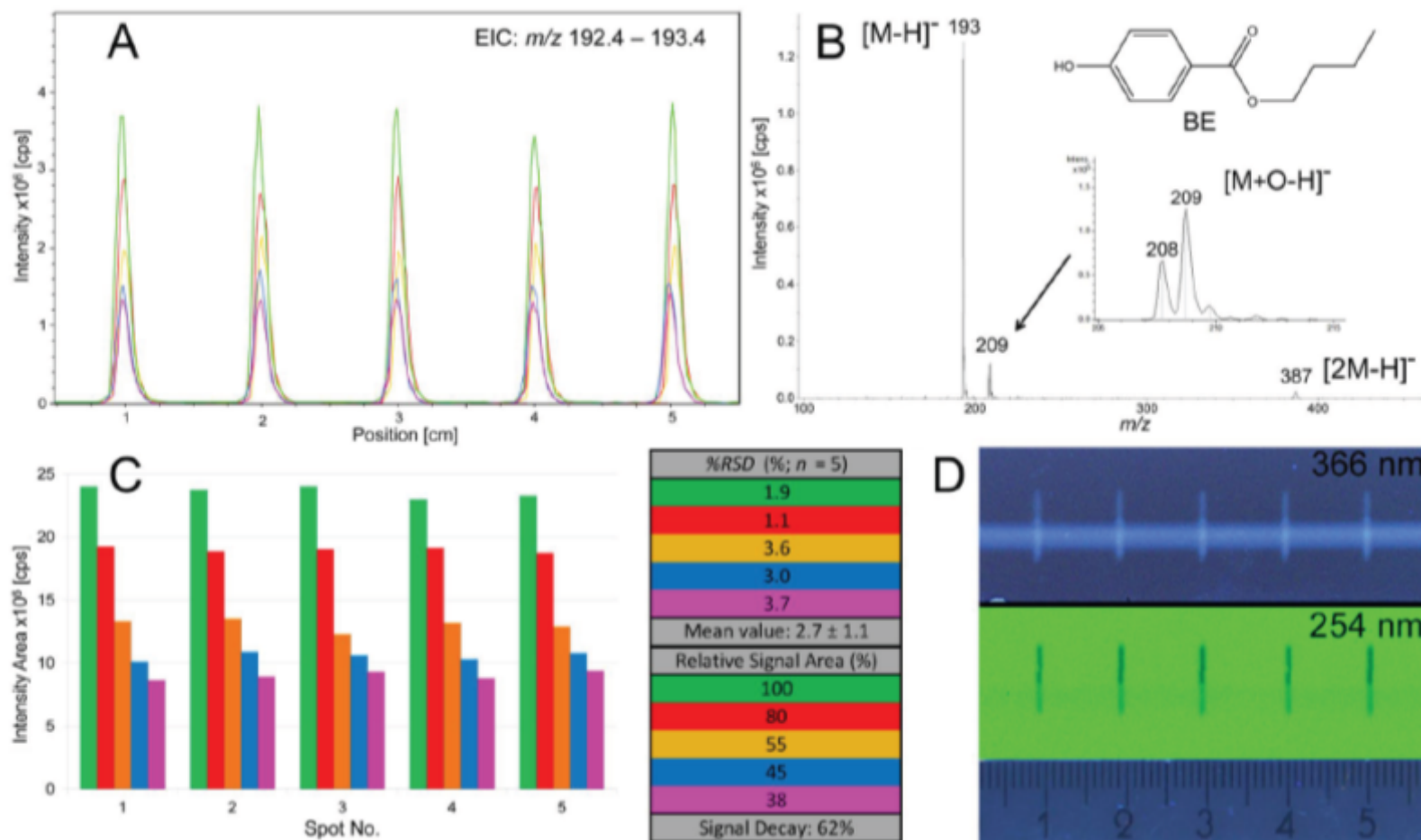


G. Morlock, Y. Ueda, *J Chromatogr* (2007)  
G. Morlock, E. Chernetsova, *Cent Eur J Chem* (2012)

👉 E. Crawford (0-9)



# HPTLC-DART SVPA-MS





# Interfacing HPTLC → MS

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## 2. Desorption-based interfaces

Direct Analysis in Real Time (DART®): Ion Sense (2005)

- ☞ DART-TLC (Morlock & Ueda, 2007)
- ☞ DART SVP 45A – 3+D Scanner (Ion Sense, 2009)

Matrix Assisted Laser Desorption and Ionization (MALDI):

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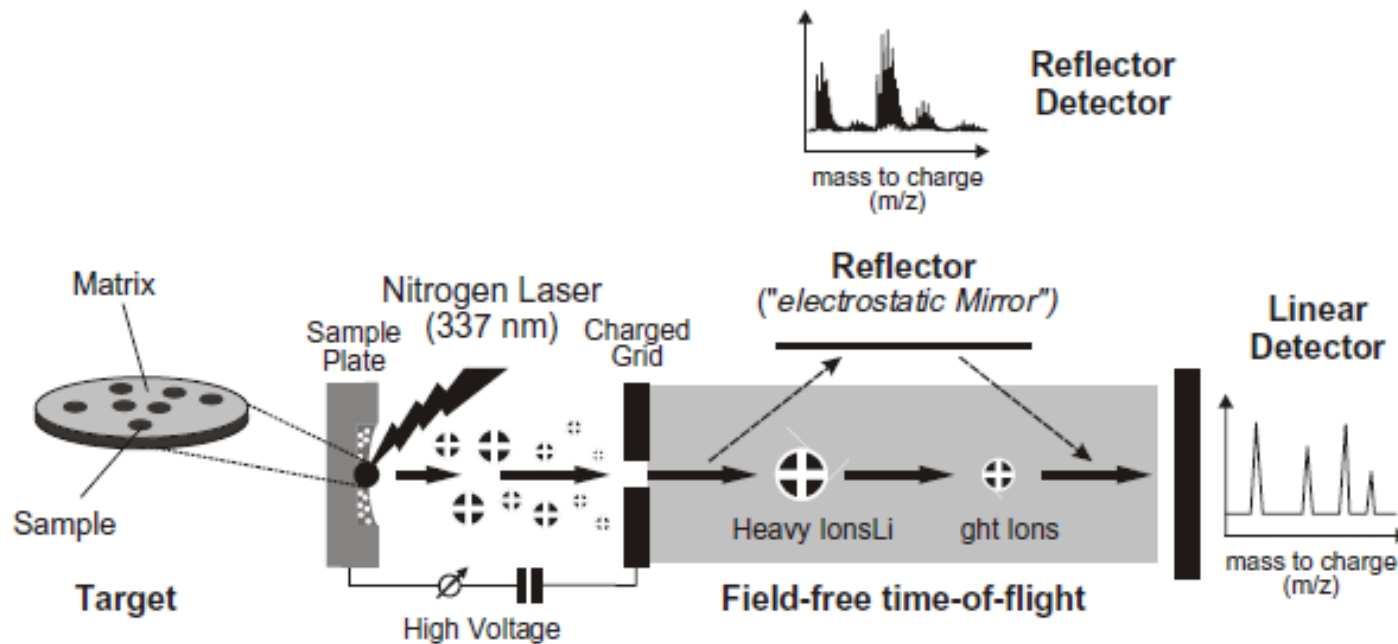
Desorption Electrospray Ionization (DESI):

- ☞ TLC-DESI (Van Berkel et al., 2005; Pasilis et al., 2007)
- ☞ Omnispray® ion sources (2D-DESI): Prosolia (2008)



# MALDI-TOFMS

MALDI: Matrix Assisted Laser Desorption and Ionization  
(Tanaka et al., 1988)





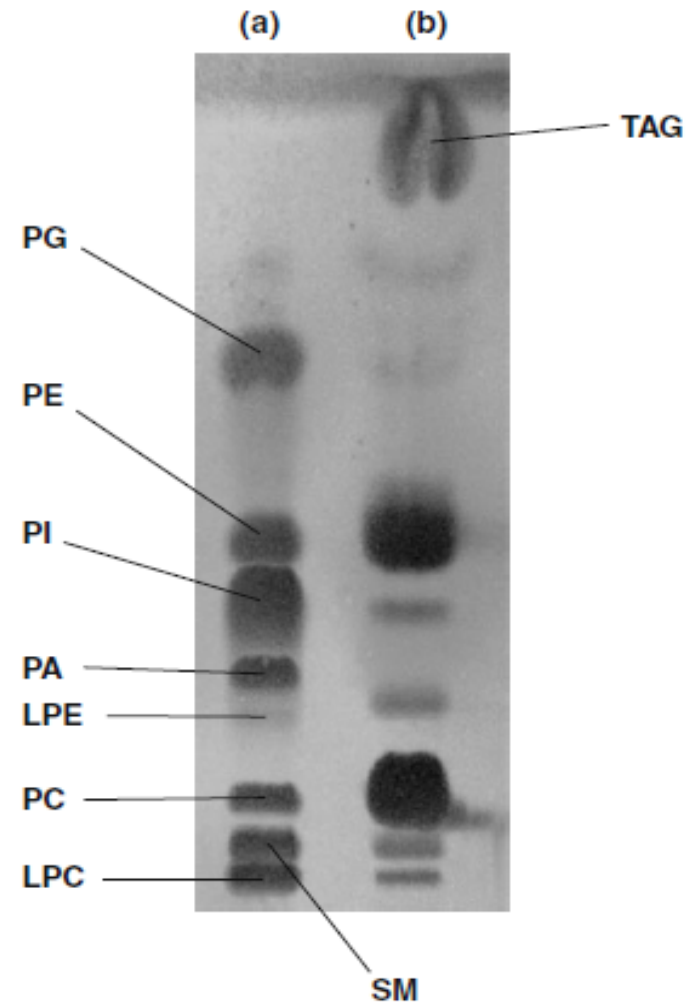
# TLC-MALDI-TOFMS

## Offline TLC-MALDI-TOFMS:

1. HPTLC and primuline staining
2. Scrape off and extract
3. Mix with matrix
4. MALDI-TOFMS

Phospholipids:

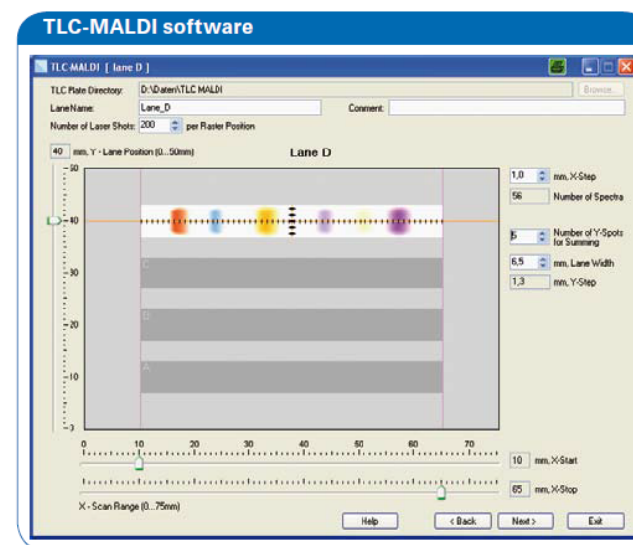
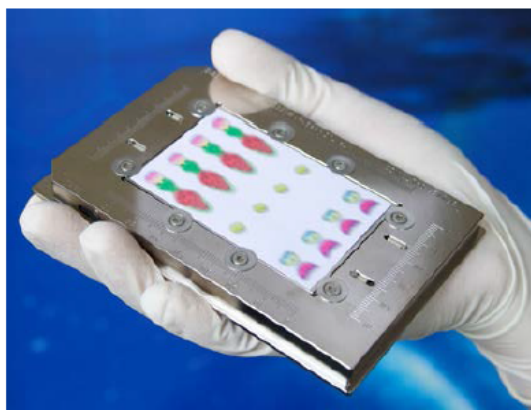
- a) standards
- b) hen yolk extract





# TLC-MALDI-TOFMS

## Online TLC-scanning MALDI-TOFMS: Bruker Daltonics (2009)

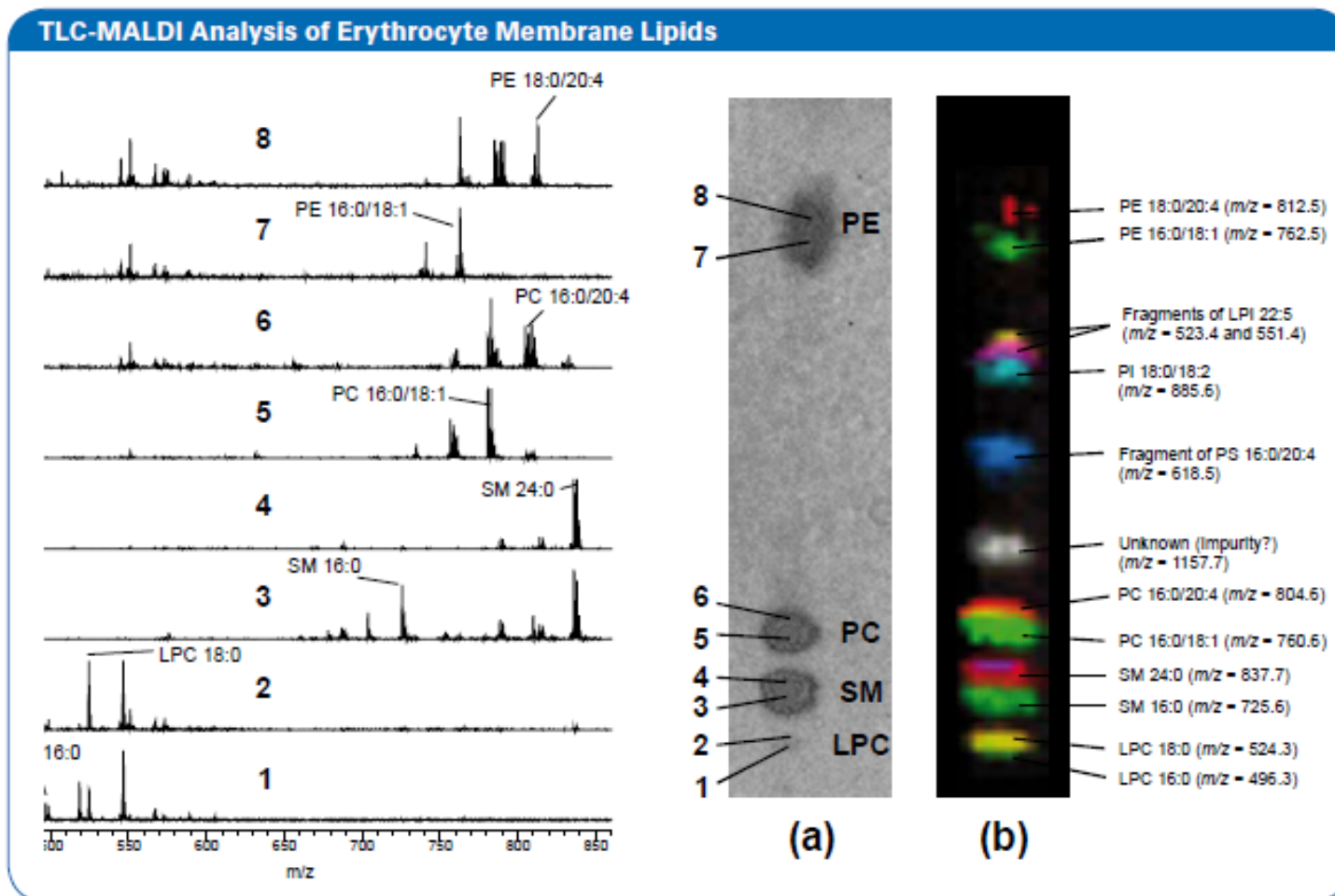


- Aluminium (electrically conductive ) backed plates (7.5 x 5 cm<sup>2</sup>)
- Dipping the plate into matrix solution (20 % DHB)
- Scanning up to 4 tracks on a plate
- Scanning time 5 min per 5-cm track
- Imaging time several hours





# TLC-MALDI-TOFMS





# Interfacing HPTLC → MS

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## 2. Desorption-based interfaces

Direct Analysis in Real Time (DART®): Ion Sense (2005)

☞ DART-TLC (Morlock & Ueda, 2007)

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☞ Omnispray® ion sources (2D-DESI): Prosolia (2008)





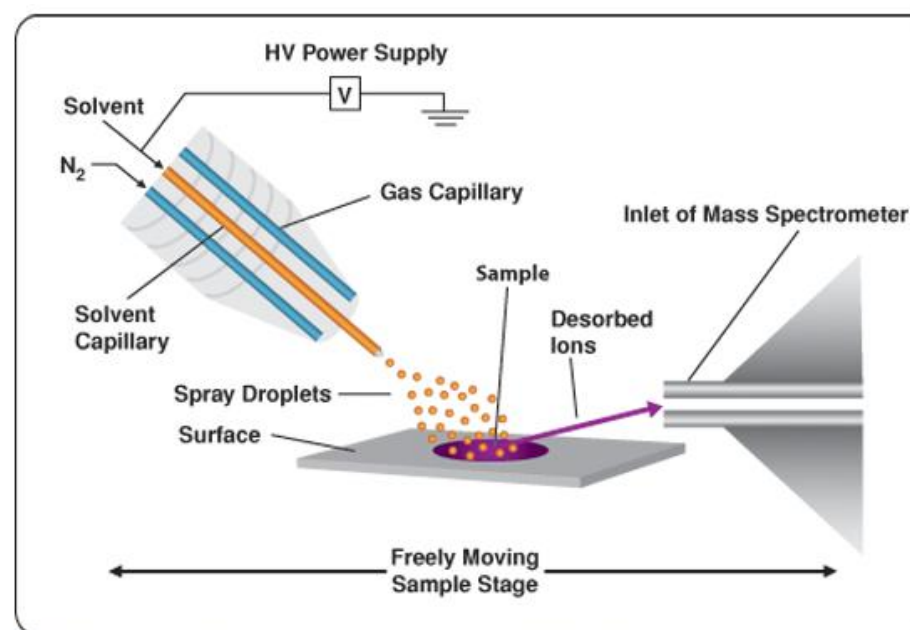
# DESI ion sources

## Principle of Operation

Desorption Electrospray Ionization (DESI) is carried out by directing high velocity charged droplets produced from a pneumatically-assisted electro spray onto a surface to be analyzed at atmospheric conditions. Ions of chemical species present on the surface are produced through the interaction of the charged droplets and the sample. The resulting mass spectra are similar to ESI mass spectra.



2-D DESI





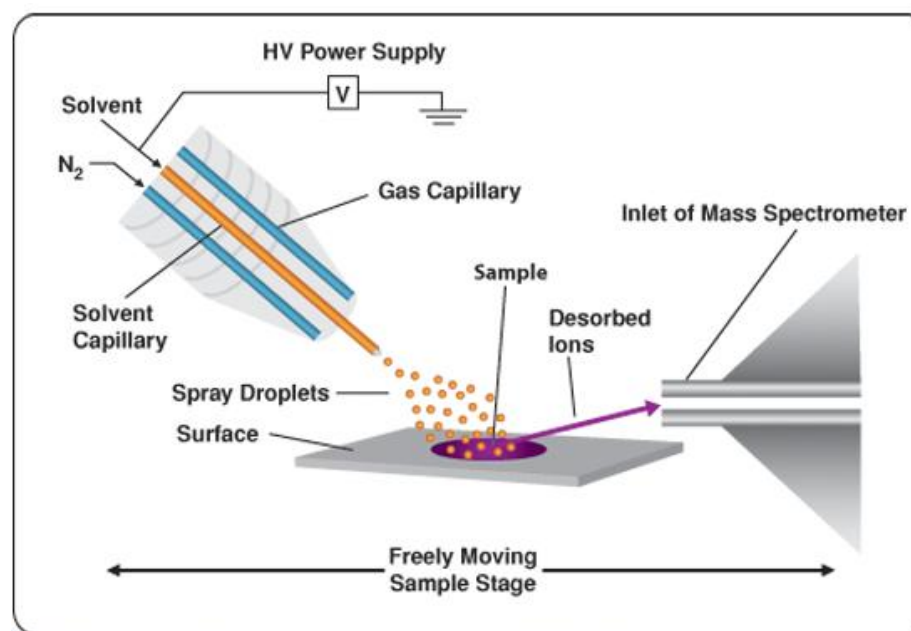
# DESI ion sources

## 2-D DESI options:

- Scanning
- Imaging
- 2 plates (5 x 5 cm<sup>2</sup>)
- NP and RP
- Sensitivity RP >> NP



Omnispray 2-D DESI





# TLC-MS

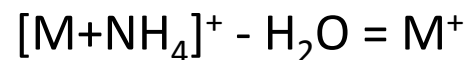
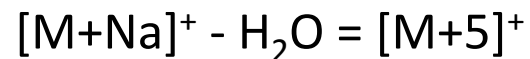
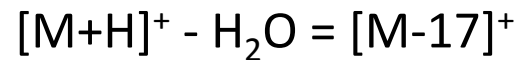
## Generally soft ionization processes (ESI, APCI, DART, MALDI)

Protonated/deprotonated molecules

Sodium adducts

Ammonium adducts

Low (in-source) fragmentations: neutral losses (even/odd rule)



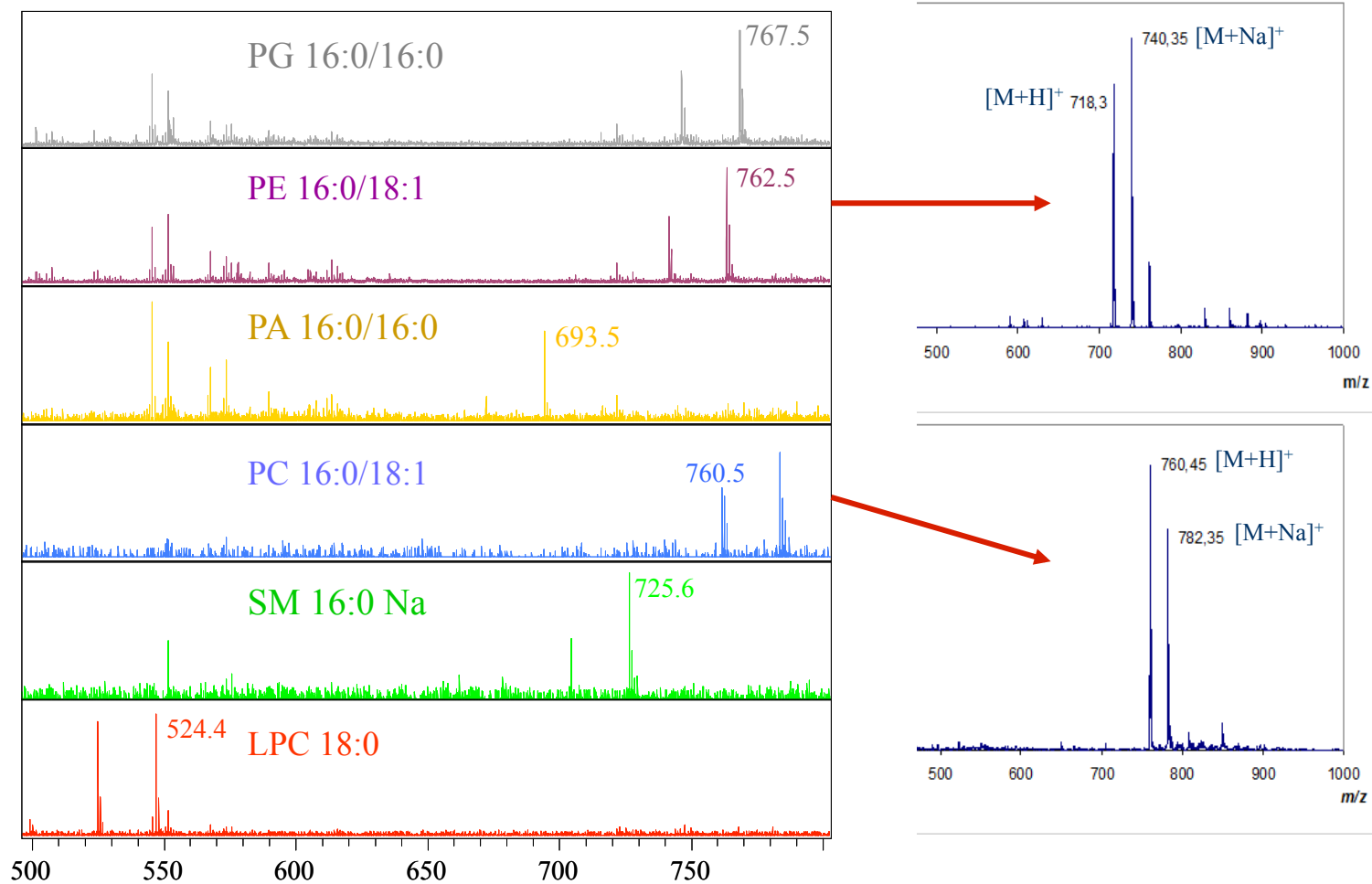
Enhanced fragmentations (CID): MS/MS, QTRAP, QTOF

☞ Substance identification, structural information



# MALDI $\leftrightarrow$ ESI

University of Hohenheim  
Institute of Food Chemistry





# FTIR spectrum

Mid-infrared → functional groups, carbon skeleton

Diffuse Reflectance Infrared Fourier Transform (DRIFT)

- TLC-DRIFT (scanning device) <sup>1)</sup>

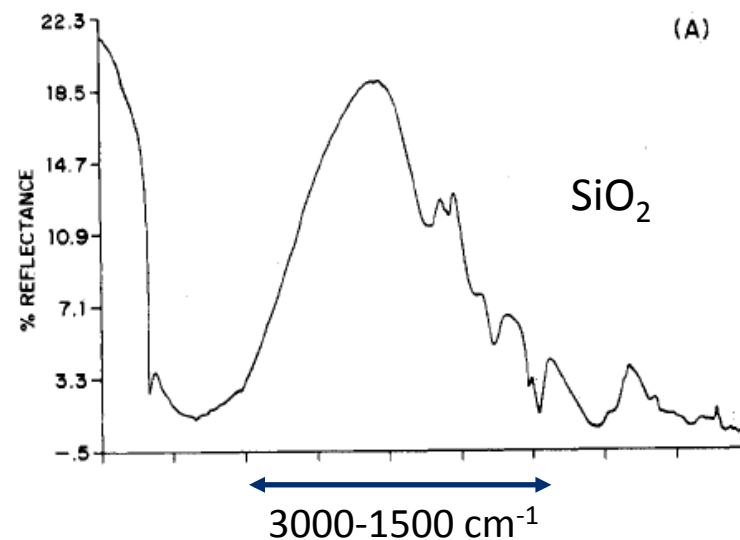
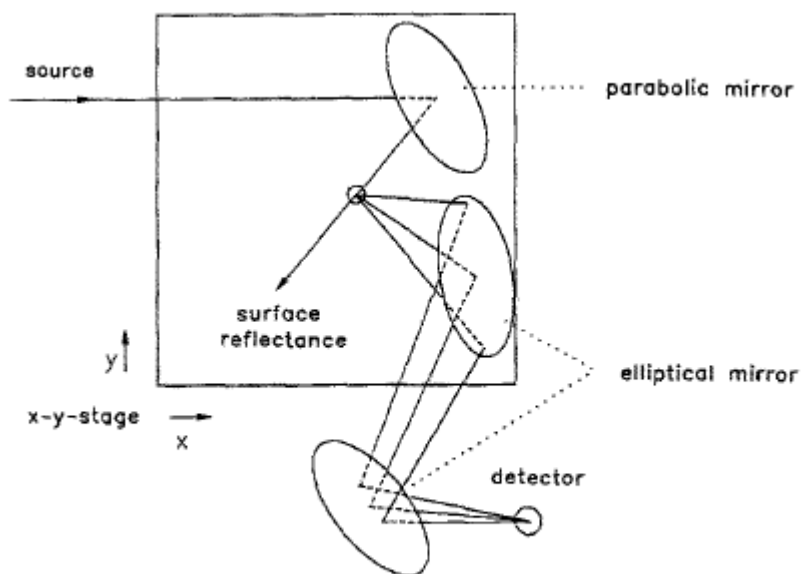
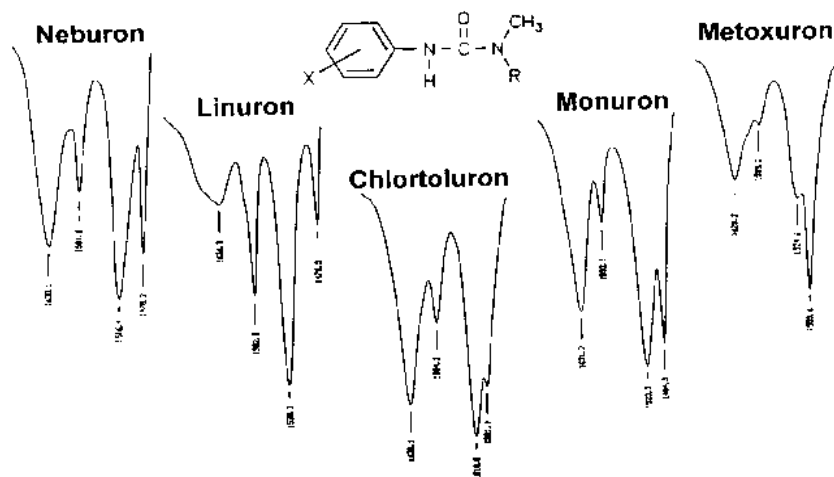


Fig. 1. Scheme of the constructed DRIFT unit for on-line measurements

1) E. Glauning et al., Fres. J. Anal. Chem. (1990)

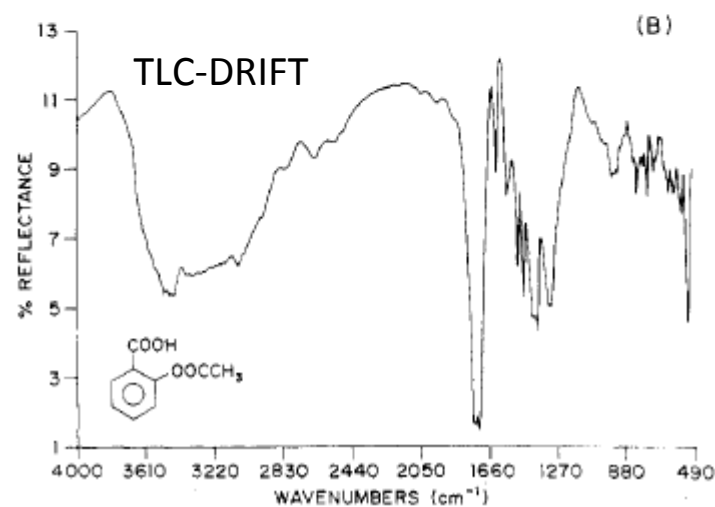
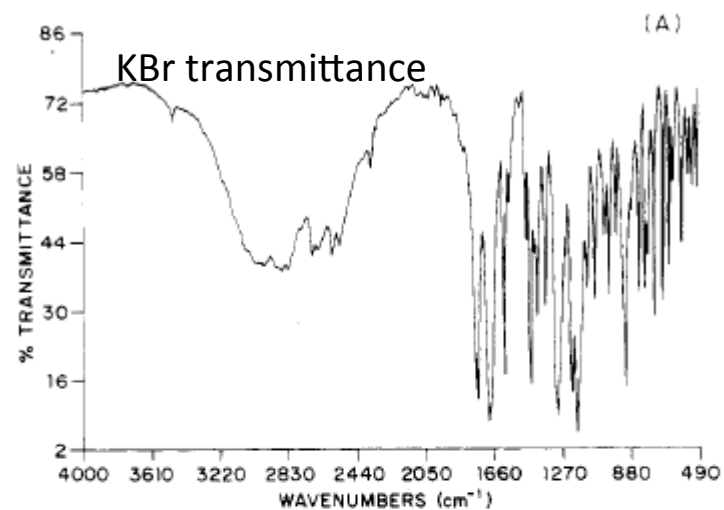


# TLC-DRIFT



G. Morlock, PhD thesis (1995)

☞ Reference libraries <sup>1)</sup>



1) E. Glauning et al., Fres. J. Anal. Chem. (1990)

G. E Zuber et al., Anal. Chem. (1984)



# FTIR spectrum

Diffuse Reflectance Infrared Fourier Transform (DRIFT)

- Actual DRIFT units

☞ Small samples (powders, pills, ...)



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



# FTIR spectrum

## Elution-based FTIR techniques

- ☞ Transfer to FTIR-microscopy slides

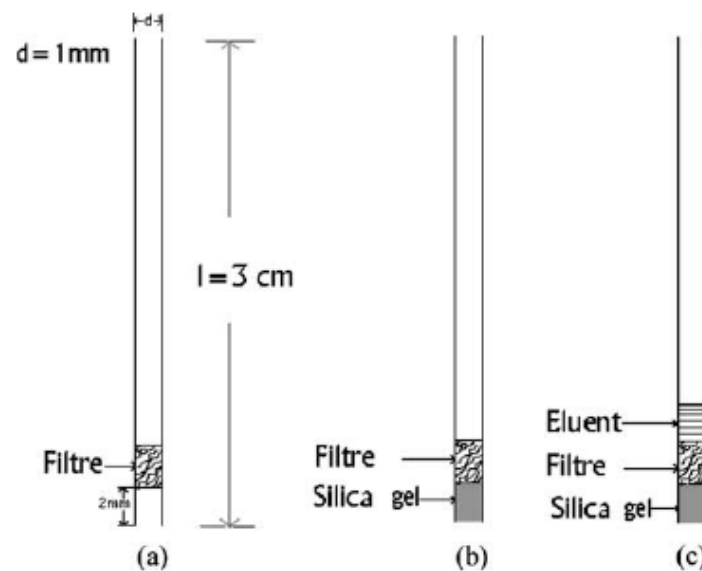


Fig. 2. Capillary transfer technique, showing the steps of transfer technique: (a) with the filter plug; (b) with filter plug and silica gel containing adsorbed compounds; (c) after eluting with the eluent, the upper eluent layer contains separated component.

W. He et al., *Vibrational Spectroscopy* 30 (2002)

W. He et al., *Spectrochimica Acta A* 61 (2005)

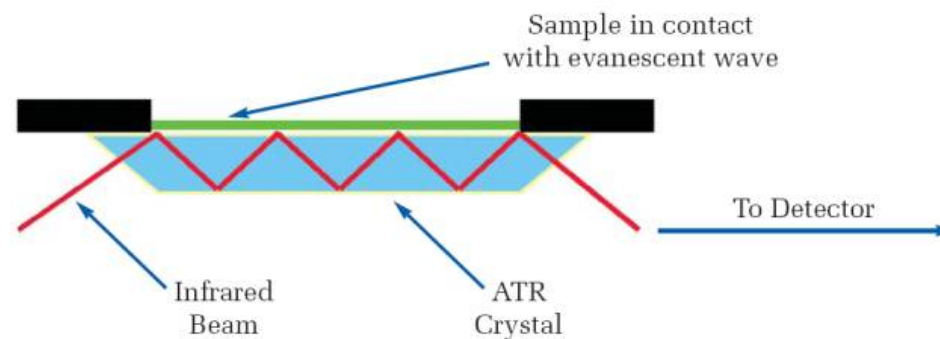
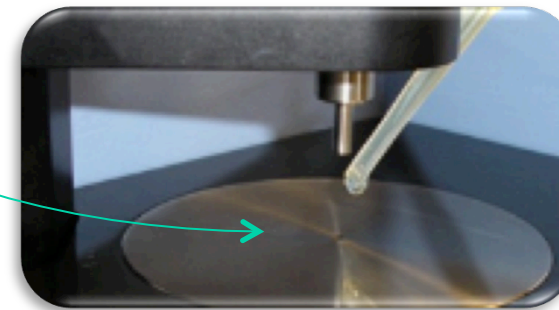




# FTIR spectrum

## Elution-based FTIR techniques

- ☞ Transfer to an ATR unit (attenuated total reflection)

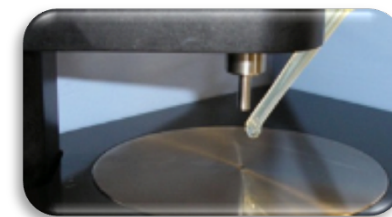




# FTIR spectrum

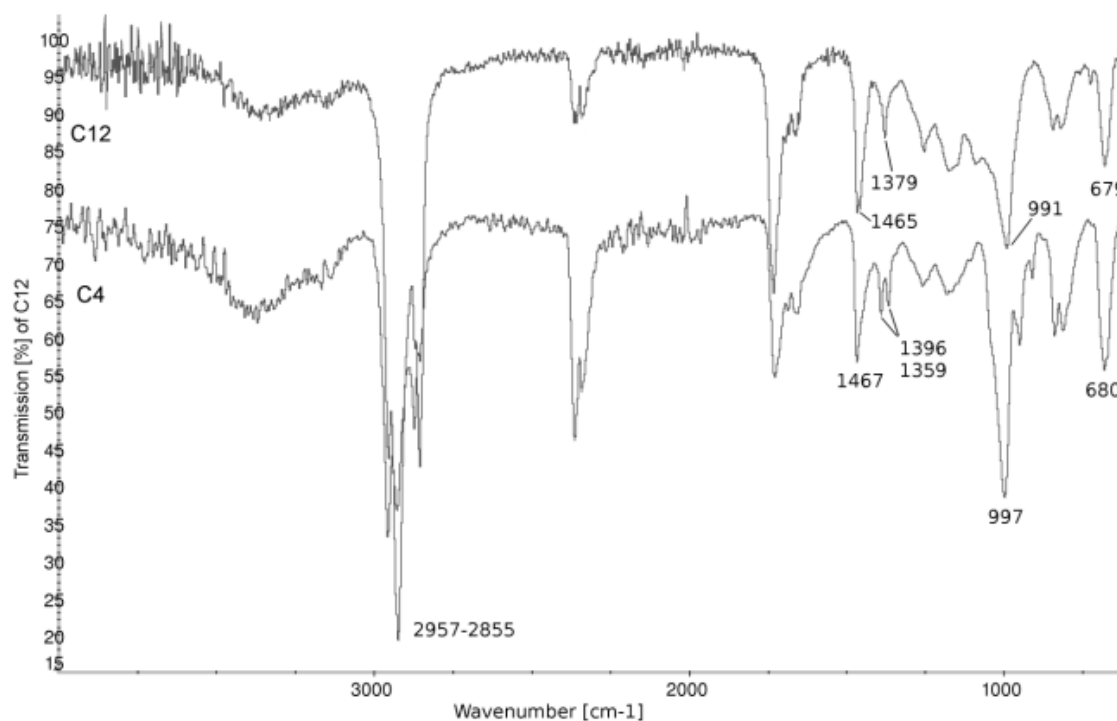
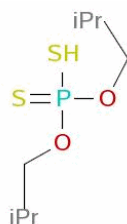
## Attenuated Total Reflection (ATR)

☞ Additives in mineral oils



Zinc-bis(O,O'-didodecyl)-  
dithiophosphate

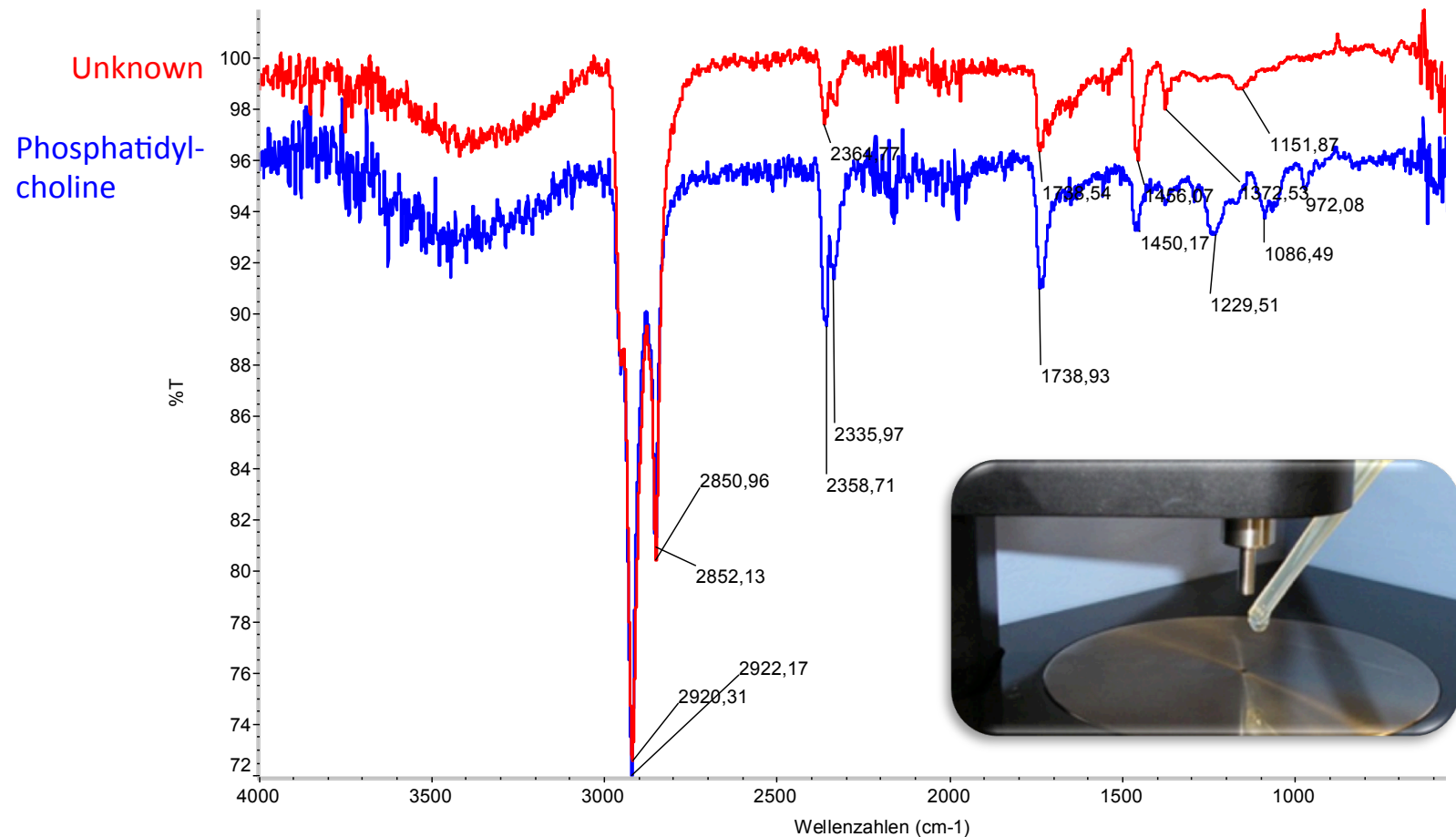
Zinc-bis(O,O'-diisobutyl)-  
dithiophosphate





# FTIR spectrum

## Bioactive components of *Lactobacillus fermentum*



G. Morlock et al. (in prep.)



# FTIR spectrum

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## Additional structural information from IR: carboxylic acid

☞ ChemSpider search ([www.chemspider.com](http://www.chemspider.com))

$$[M+H]^+ = 319.22677$$

- Monoisotopic mass  $\pm 0.00032$  (1 ppm)  
=> 1060 hits (all with  $C_{20}H_{30}O_3$ )
- Including name fragment (substring): „acid“  
=> 185 hits

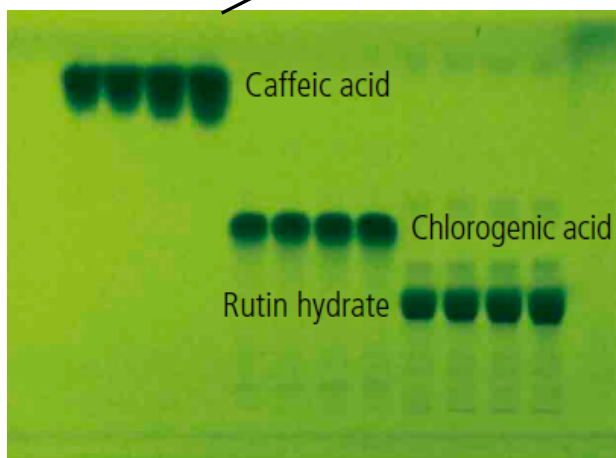
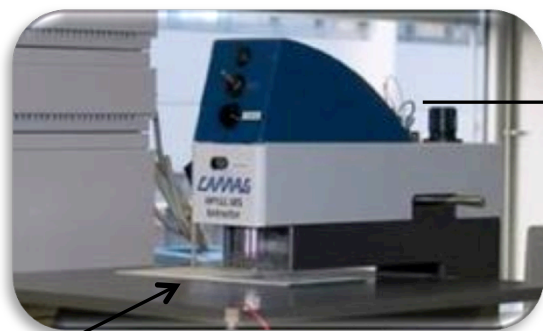
☞ Additional structural information: NMR ?



# $^1\text{H}$ -NMR spectrum

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## $^1\text{H}$ -NMR: proton spin systems



A. Gössi, U. Scherer, G. Schlotterbeck, *Chimia* (2012)  
CBS 110 (2013)



# $^1\text{H}$ -NMR spectrum

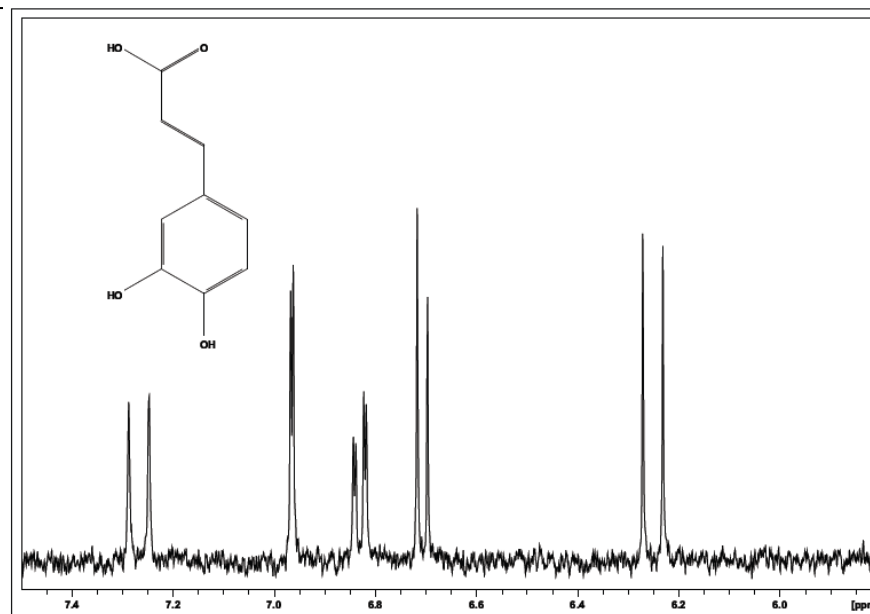


Fig. 1. Aromatic region of  $^1\text{H}$  NMR Spectrum of 15.6  $\mu\text{g}$  caffeic acid extracted from a TLC plate after development and measured at 400 MHz.

600  $\mu\text{L}$   $\text{d}_4$ -methanol (400 MHz)

$^1\text{H}$ NMR	Measurements in solution			After elution from the plate		
	Linearity $r^2$	LOD ( $\mu\text{g}$ )	LOQ ( $\mu\text{g}$ )	Quantity ( $\mu\text{g}/\text{band}$ )	Recovery (%)	%RSD (n=3)
Rutin	0.9976	2.3	6.9	20.3	101.8 $\pm$ 4.0	3.9
Caffeic acid	0.9978	2.5	7.3	17.1	103.4 $\pm$ 1.0	1.5
Chlorogenic acid	0.9991	3.3	10.1	19.6	100.5 $\pm$ 3.1	3.1



# NMR spectrum

$^1\text{H}$ -NMR/ $^{13}\text{C}$ -NMR including 2-D correlation spectra

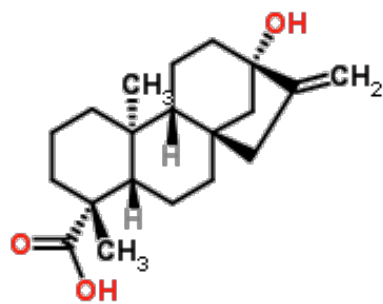
☞ More substance than some  $\mu\text{g}$  is needed

Due to the highest yield and purity, zones 1 and 6 were isolated for NMR spectroscopy and determination of their biological activity. For this purpose fractions A5 and A6 were applied on HPTLC plates as 16 cm lines with concentrations of 5.0  $\mu\text{g}/\text{mm}$ . From 20 HPTLC plates 1.58 and 1.75 mg of the substances 1 and 6 were obtained, respectively. The structures were characterised by one- and two-dimensional  $^1\text{H}$ - and  $^{13}\text{C}$ -NMR spectroscopy and mass spectrometry. The accurate molecular masses of the two oily substances were determined in an UHPLC-QTOF MS system. Flow injection analysis was performed with 0.3  $\mu\text{L}$  sample solution and a flow rate of the mobile phase of 0.4 mL/min in an isocratic mode. The mobile



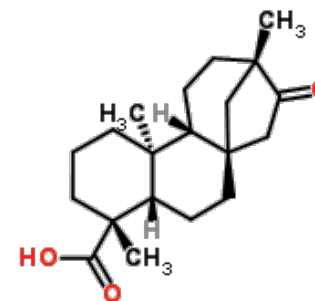


# $^1\text{H-NMR}$ spectrum



Steviol

or



iso-Steviol

$\delta$  (ppm) =

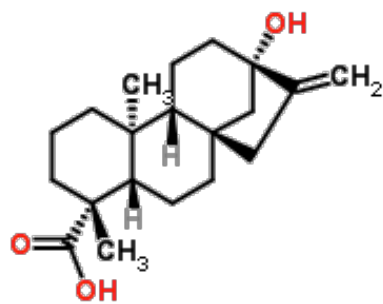
0.9 (s, 3 H,  $-\text{CH}_3$ )

1.0 (s, 3 H,  $-\text{CH}_3$ )

4.9 (s, 2 H,  $=\text{CH}_2$ )

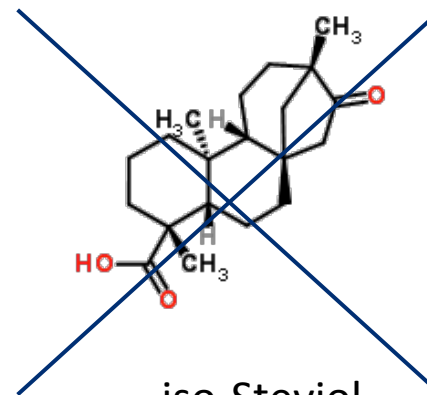


# $^1\text{H-NMR}$ spectrum



Steviol

or



iso-Steviol

$\delta$  (ppm) =

0.9 (s, 3 H,  $-\text{CH}_3$ )

1.0 (s, 3 H,  $-\text{CH}_3$ )

4.9 (s, 2 H,  $=\text{CH}_2$ )



# Conclusions

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- HPTLC-MS - the most sensitive interface?
- Scanning MS devices ☞ selective detectors
- From a detected zone to the chemical structure?
  - ☞ High-resolution MS alone does not simply provide the structure of an unknown compound.
  - ☞ Strong support by IR and NMR spectra is required.
- TLC-MS interface
  - ☞ mass spectra, ATR-FTIR spectra,  $^1\text{H-NMR}$ -spectra
  - ☞ analytical scale (MS: ng/zone, FTIR/ $^1\text{H-NMR}$ :  $\mu\text{g/zone}$ )
- Good luck for your study from the detected zone to the chemical structure!



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Many thanks to Gerda Morlock for providing literature and materials for my presentation and for helpful discussions.

Many thanks to you for your kind attention!



# Liquid microjunction

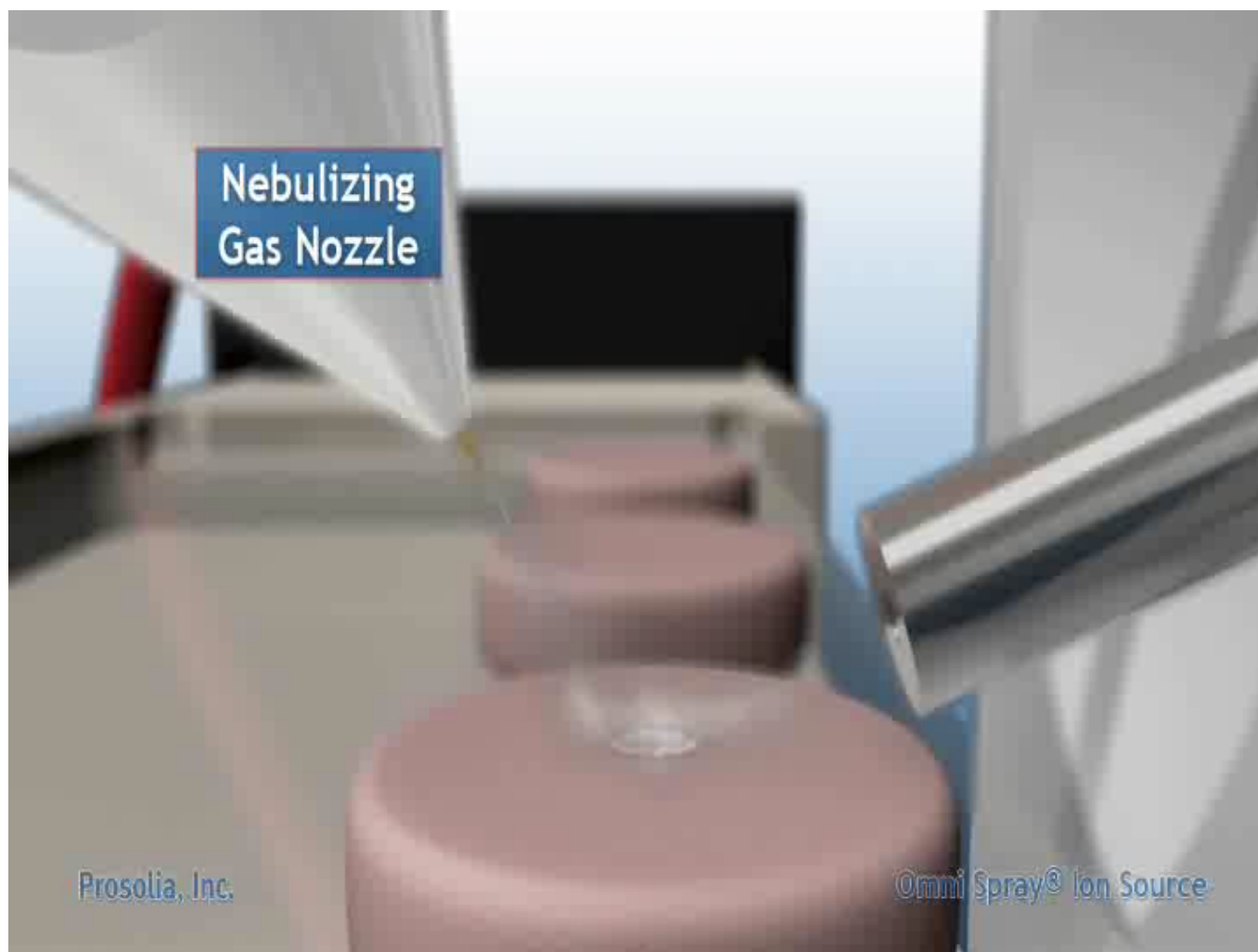
## In situ microextraction (flowprobe™): Prosolia





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# DESI ion sources





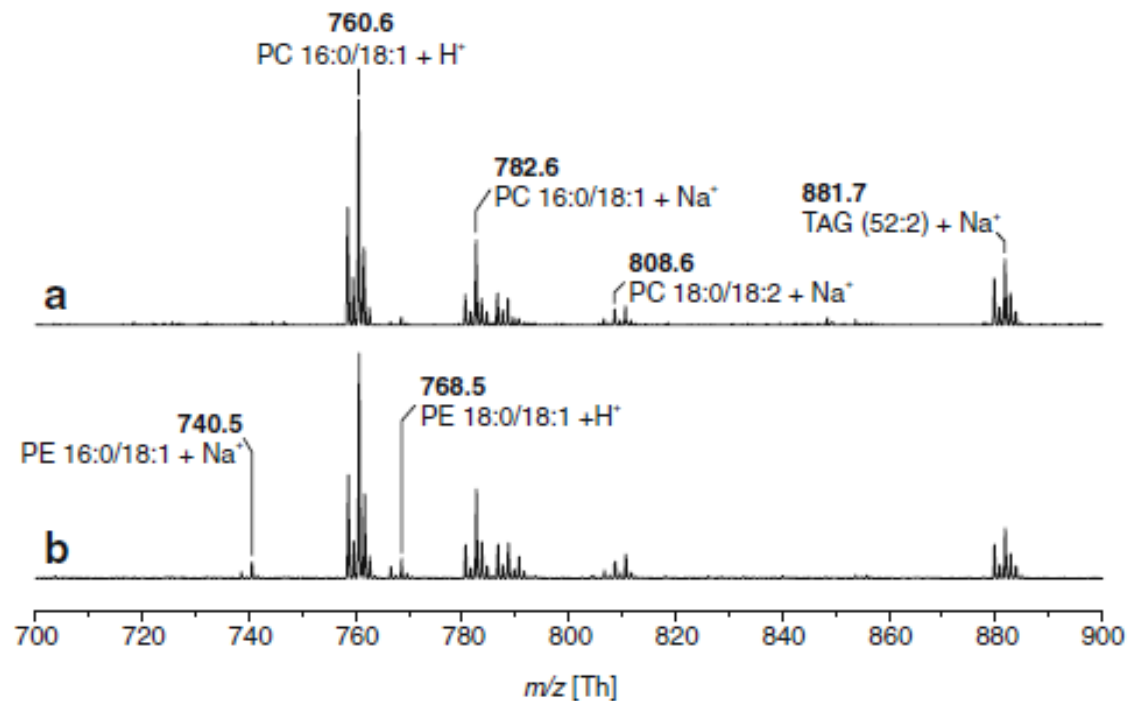
# TLC-MALDI-TOFMS

## Offline TLC-MALDI-TOFMS:

1. TLC and primuline staining
2. Scrape off and extract
3. Mix with matrix
4. MALDI-TOFMS

Hen yolk extract

- a) directly
- b) after HPTLC







# TLC-MALDI-TOFMS

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## TLC-scanning MALDI-TOFMS: Bruker Daltonics (2009)

