

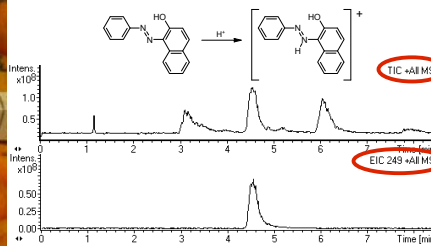
Liquid chromatography tandem mass spectrometry (LC-MSMS) - the primary tool for trace contaminant analysis



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Outline

- Trace analysis
- How to connect LC and MS?
- Electrospray (ESI) ion source
- LC-ESI-MS
- LC-ESI-MSMS

Trace contaminant analysis

- Samples: almost never pure compounds but **(very) complex mixtures** (matrixes)
 - Food, soil, biological samples, ...
- Analytes (compounds that we determine) often at **trace level**
 - pesticides, drug residues, mycotoxins, ...
- Contents often in the range of ppm and ppb
 - **1 ppb \equiv 1 μ g/kg \equiv 1 mg/t**

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The central problem of trace contaminant analysis:

The analysis method has to **find and reliably identify** the trace analyte in the presence of a large number of main components

- Organic traces: chromatography coupled with mass spectrometry



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LC-MS as analysis technique

- Combines two powerful techniques:

LC (liquid chromatograph) **separates** the analyte from other sample components

MS (mass spectrometer) **detects** and **identifies** the analytes and **determines concentration**



LC and MS: strong contrast

- LC separation is carried out in **liquid phase** (mobile phase)
- MS detection proceeds in **high vacuum** ($n \cdot 10^{-6}$ mbar)



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Coupling LC and MS

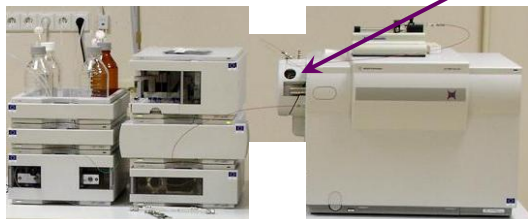
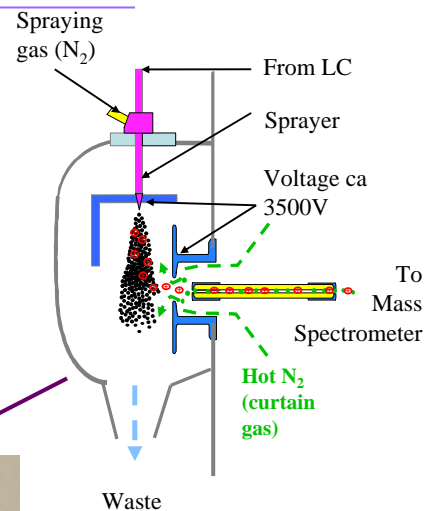
- The connecting interface has to convert compounds in the liquid phase into ions in the gas phase
- Mobile phase must not get into MS
- Developing this interface has been the biggest challenge in development of LC-MS

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Electrospray (ESI) ion source

- The liquid flow is dispersed by electric field into small droplets
- The analyte ions evaporate from droplets
- The ions are directed into MS entrance by the electric field



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ESI in action (Positive ions)

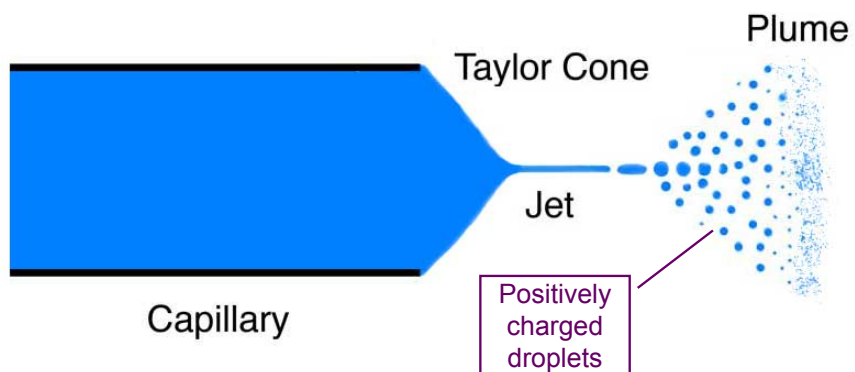
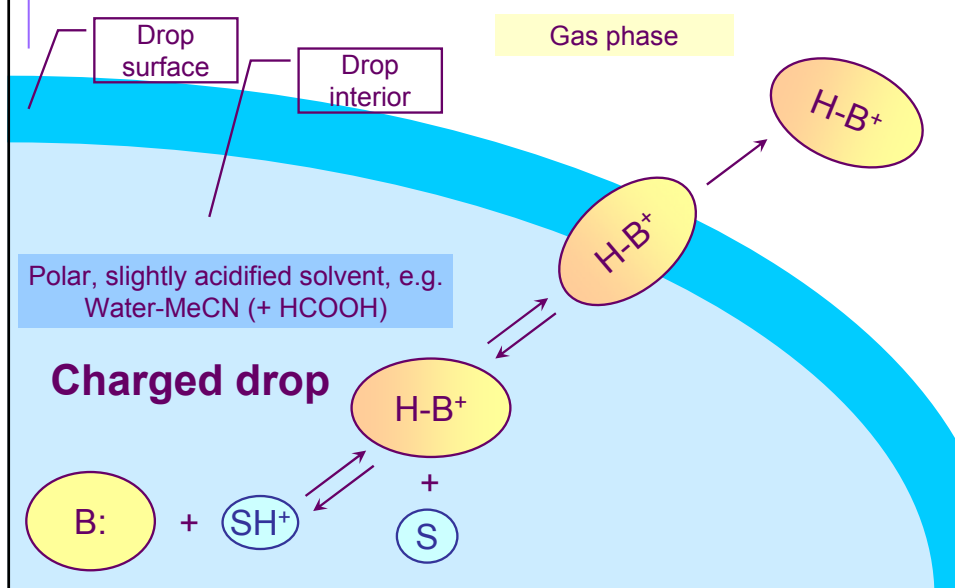


Image by K. K. Murray (via Wikipedia)

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ESI mechanism with small molecules



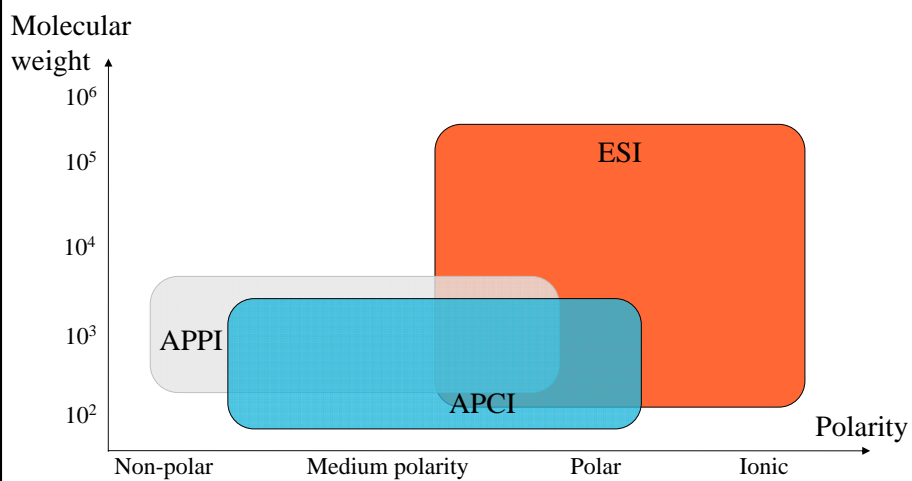
ESI Ionization

- The most widely applied ionization method in MS
 - Nobel prize in 2002 (Fenn, Tanaka, Wüthrich)
- Ionization via
 - Protonation
 - Deprotonation
 - Adduct formation
- Complex mechanism

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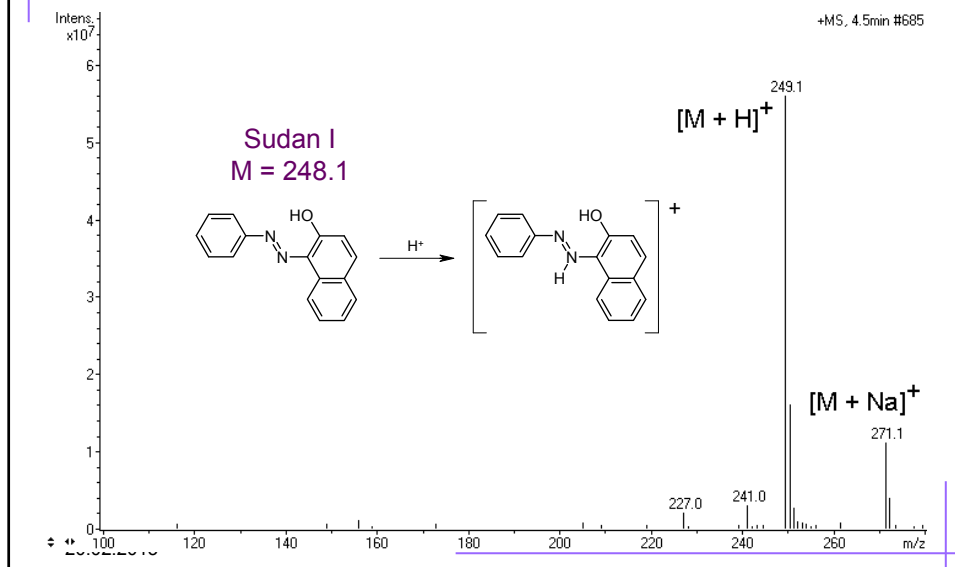
LC-MS usability with different ion sources



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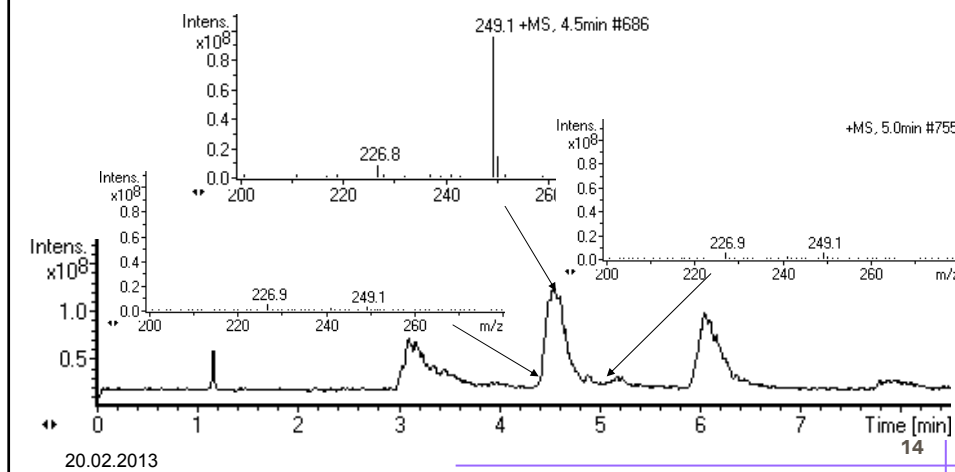
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ESI-MS spectrum



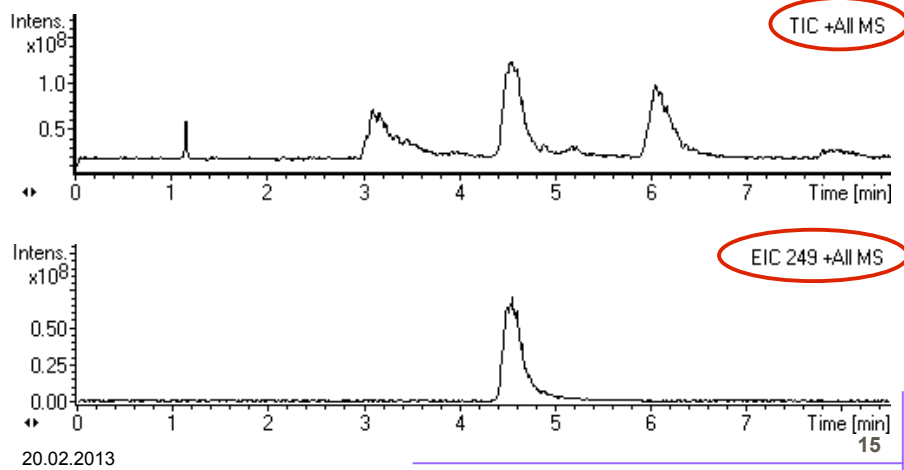
LC-ESI-MS: Mass-chromatogram (TIC)

- Three-dimensional data: Mass spectra can be obtained for LC peaks



LC-ESI-MS: Mass-chromatogram (EIC)

- MS acts as another separation technique!



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Problems with LC-ESI-MS

- ESI mass spectra are **not very characteristic**
 - ESI is soft and ions do not fragment extensively
 - Often just $[M+H]^+$
 - The lower the analyte levels the higher the probability that some interfering compound has the same retention time and gives ions with the same m/z
- ESI mass spectra are **quite noisy**
 - High LoD values

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Solution: Tandem mass spectrometry LC-ESI-MSMS

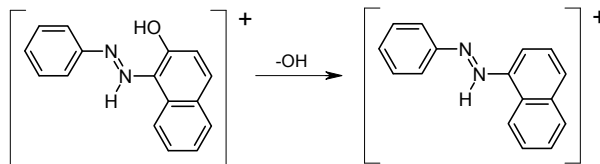
- MSMS workflow:
 - The main ion (usually $[M+H]^+$) is selected
 - All other ions are ejected
 - The selected ion is excited
 - By collision with inert gas molecules
 - The ion fragments
 - By ejecting some part of the molecule
- This is controlled by the MS software
- Needs triple quadrupole or ion trap mass analyzer
 - Cannot be done with single quadrupole

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LC-ESI-MSMS with Sudan I

- Sudan I: main transition $249 \rightarrow 232$



- Now identity is confirmed in **triplicate**
 - Retention time, m/z of parent ion, m/z of fragment ion
 - For higher reliability in identification several transitions can be monitored

Journal of Chromatography A, 1160 (2007) 227–234

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LC-ESI-MSMS

- Almost all quantitative trace analysis is done in the MSMS mode
 - Better signal to noise ratio
 - Couple of orders of magnitude lower LoD
 - Few μl of injected solution with few ppb concentration is sufficient
 - Reliable identification by using different transitions

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LC-ESI-MSMS: standing problems

- Different ionization efficiency of different compounds
 - Also some polar compounds do not ionize well
- Dependence of ionization efficiency on co-eluting compounds – **matrix effect**
- Poor retention of some ionic compounds in LC

See more at:

http://tera.chem.ut.ee/~ivo/Chrom_MS/

http://tera.chem.ut.ee/~ivo/LC-MS_Matrix_Effect_Toolbox/

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LC-MS vs GC-MS

LC-MS “+”

- Almost unlimited M
- No volatility needed
 - Ionic compounds OK
- Thermal stability is not needed

LC-MS “-”

- Expensive
- Not robust
- Predicting ionization is tricky

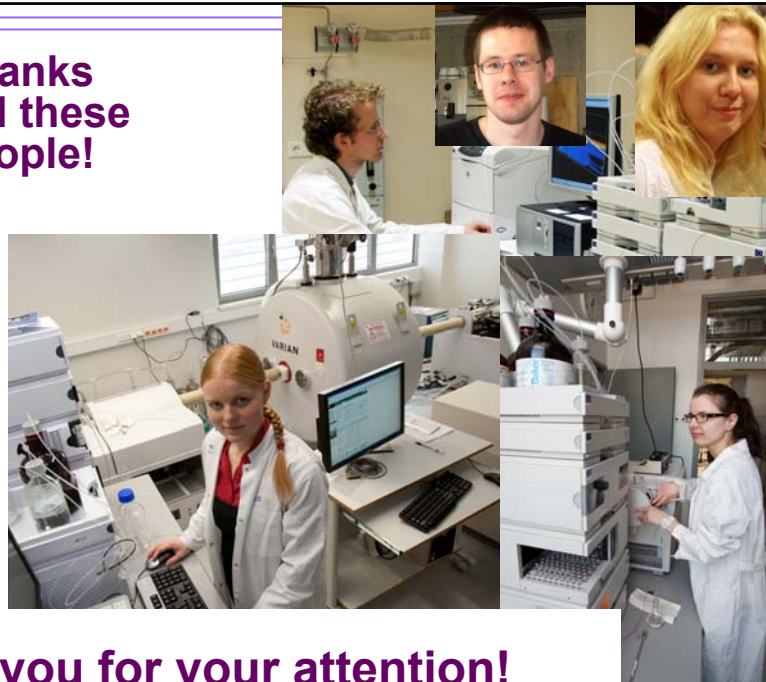
GC-MS “+”

- Better resolution
- Robust

GC-MS “-”

- Low to medium M
- Volatility needed
- Thermal stability needed

**Thanks
to all these
people!**



Thank you for your attention!