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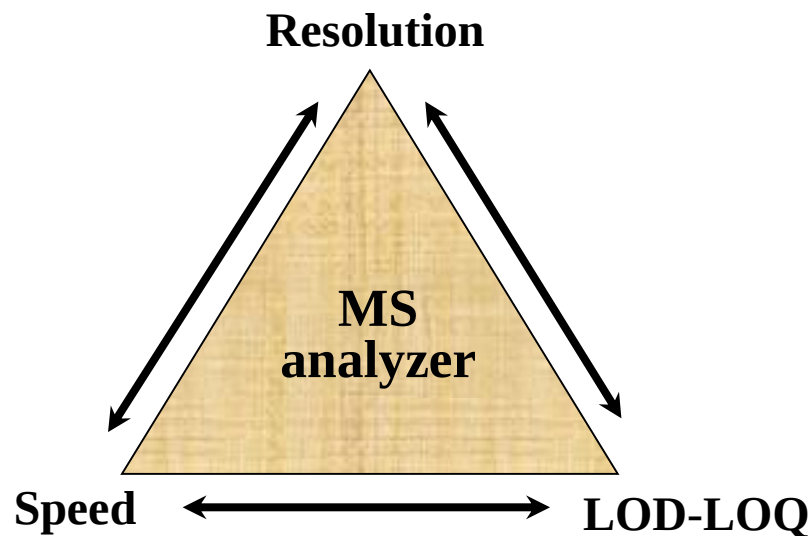
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Methods of ions analysis and detection in MS

Ion analysis (m/z)	Ion detection
range resolution accuracy speed	LOD-LOQ dynamic range stability in time lifetime / durability



Methods of ions analysis in MS

- ion analysis is realized using a mass analyzer
(alternatively called mass filter or mass separator)
- mass analyzer = part of mass spectrometer,
which distinguishes ions
according to their m/z value
- parameters of mass analyzers:
 - a) range of analyzed m/z values
 - b) scan speed, switching of scan modes
 - c) resolution power (resolution) of m/z values
 - d) mass accuracy



Parameters of MS analyzers

➤ range of analyzed m/z values:

a) lower limit: from technical '0' or from a higher value

b) upper limit: up to a defined maximum value
(for multiply charged ions higher M)

➤ scan speed and the ability to switch between different modes:

a) number of scans per second:

< 1	→ low speed
1 - 10	→ medium speed
10 - 500	→ high speed

b) measurement modes: full scan, segment scan, SIM, SRM, MRM

c) display modes:

- TIC (total ion current) → total ion current (including noise)
for the selected measurement mode
- RIC/EIC (reconstructed/extracted ion chromatogram)
→ record for the selected m/z value
- BPC (base peak chromatogram) → record for basic (most intensive)
 m/z values in spectrum at each point of the chromatogram



Parameters of MS analyzers

- resolving power – RP (mass analyzer or spectrometer):
- two different definitions – depends on the type of mass analyzer
- a measure of the mass analyzer's ability to distinguish close m/z values – i.e. what the smallest difference can be recorded; relative data related to m/z values

1. definition for two ions:

$$RP = m_2 / (m_2 - m_1); z_1, z_2 = 1$$

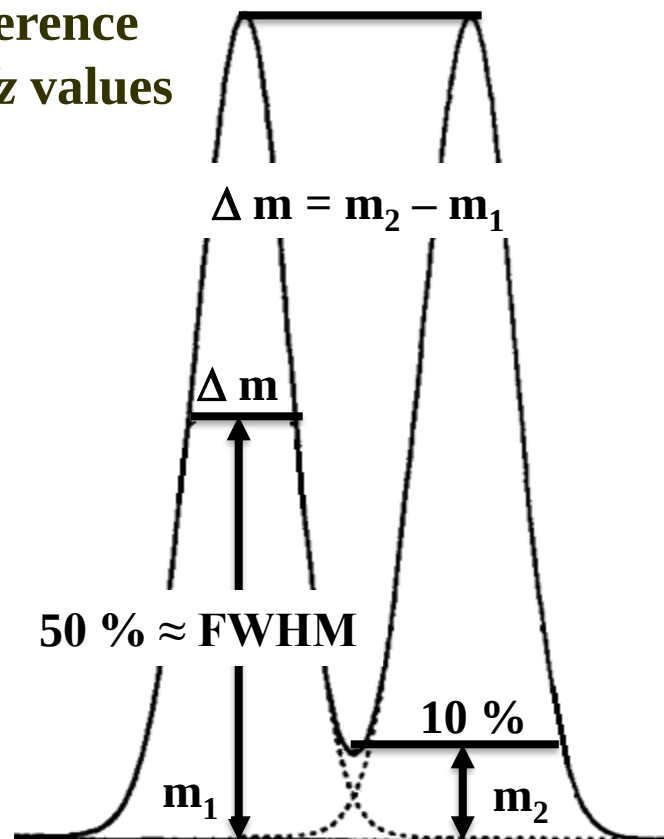
- peaks m_1 and m_2 have 10% overlap at the same height

dimensionless

2. definition for one ion (preferred):

$$RP = m_1 / \Delta m \text{ (alt. } t / 2 \Delta t \text{ - TOF)}$$

- FWHM (full width at half maximum) for the corresponding m/z
- dimensionless*

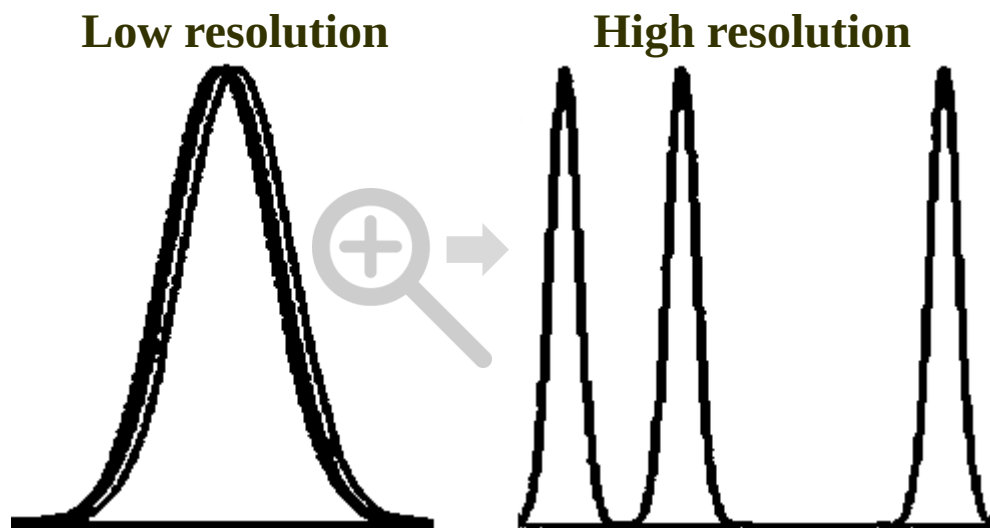


Parameters of MS analyzers

- resolution - $R = 1/RP$ (mass analyzer or spectrometer):
- two different definitions – depends on the type of mass analyzer
- a measure of the mass analyzer's ability to distinguish close m/z values – i.e. what the smallest difference can be recorded; relative data related to m/z values

$$R = (m_2 - m_1) / m_1 = \Delta m / m_1$$

- expressed in ppm for the corresponding m / z



As the resolution increases, the number of compounds corresponding to the determined m/z value decreases.

So the probability of identification increases.

Parameters of MS analyzers

- mass accuracy:
- a measure of agreement between measured (experimental) and calculated (exact / theoretical) m/z
- a measure of the ability of mass analyzer to determine the correct m/z

Relative measure in ppm - depends on m/z value:

$$\text{ppm} = 10^6 \cdot (m/z_{\text{EXP}} - m/z_{\text{CAL}}) / m/z_{\text{CAL}}$$



Parameters of MS analyzers

Examples of relationships between parameters and m/z values

1. Resolving power (RP) = 50000

(often given as a resolution FWHM 50000)

- for $m/z = 1000$ ions distinguished with difference $\pm 0,01$ ($1000/50000 = 0,02$)
- for $m/z = 100$ ions distinguished with difference $\pm 0,001$ ($100/50000 = 0,002$)

2. Resolution (R) for RP 50000 is 20 ppm (i.e. $1/50000 = 20 \cdot 10^{-6}$)

3. Mass accuracy

- a) $\text{ppm} = 10^6 \cdot (266,16315 - 266,16304) / 266,16304 = + 0,41 \text{ ppm}$
- b) $\text{ppm} = 10^6 \cdot (266,16187 - 266,16304) / 266,16304 = - 4,40 \text{ ppm}$
- c) $\text{ppm} = 10^6 \cdot (266,17087 - 266,16304) / 266,16304 = + 29,4 \text{ ppm}$



Methods of ions analysis in MS

Principle of mass analyzer

- ion analysis is realized at low pressure ($< 10^{-4}$ Pa)
- mass analyzer determines m/z value
- multistage mass analysis - combination of several mass analyzers

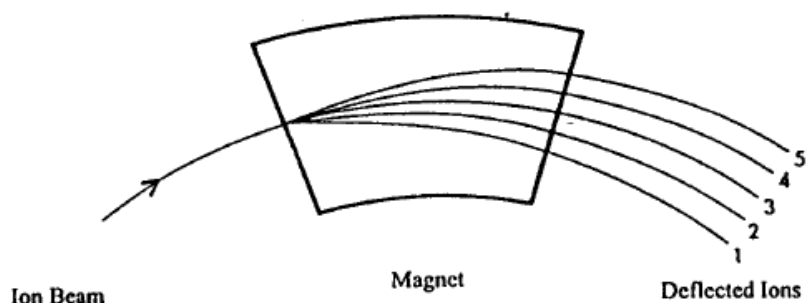
Physical principles of ion analysis

- ❖ curved flight path of ions in a magnetic or electric field
 - magnetic or electrostatic analyzer
- ❖ various stability of ion oscillations in two- or three-dimensional
 - combination of DC and high frequency AC voltage
 - quadrupole or ion trap - Q, IT
- ❖ different time of flight of ions - time of flight analyzer - TOF
- ❖ different frequency of harmonic oscillations - Orbitrap
- ❖ different energy absorption during cycloidal motion of ions in a combined magnetic and electric field - ionic cyclotron resonance - ICR

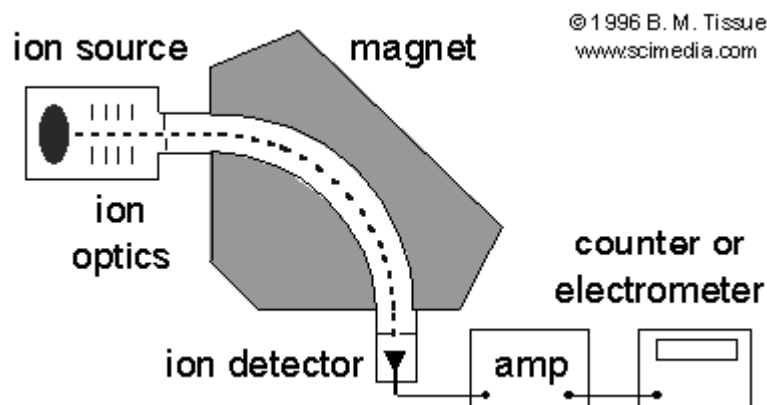


Principles of mass analyzers

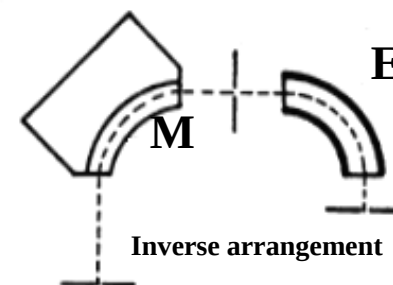
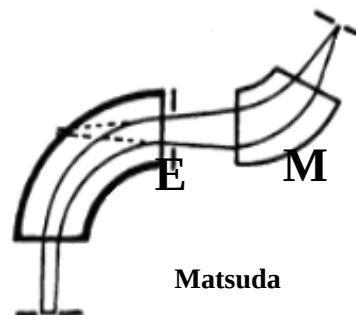
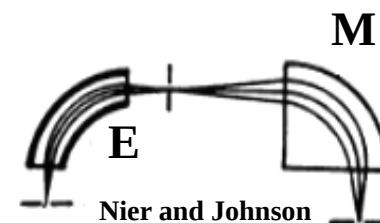
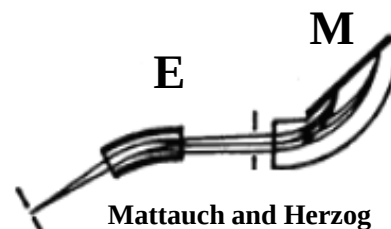
Magnetic sector



$$(m/z)_1 < (m/z)_2 < \dots < (m/z)_5$$

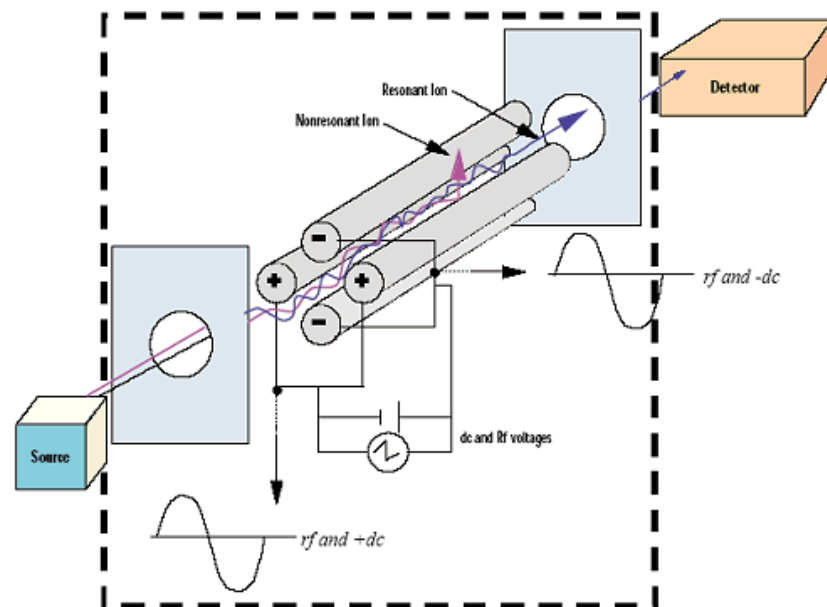


Double focusing: electrostatic (E) and magnetic (M)

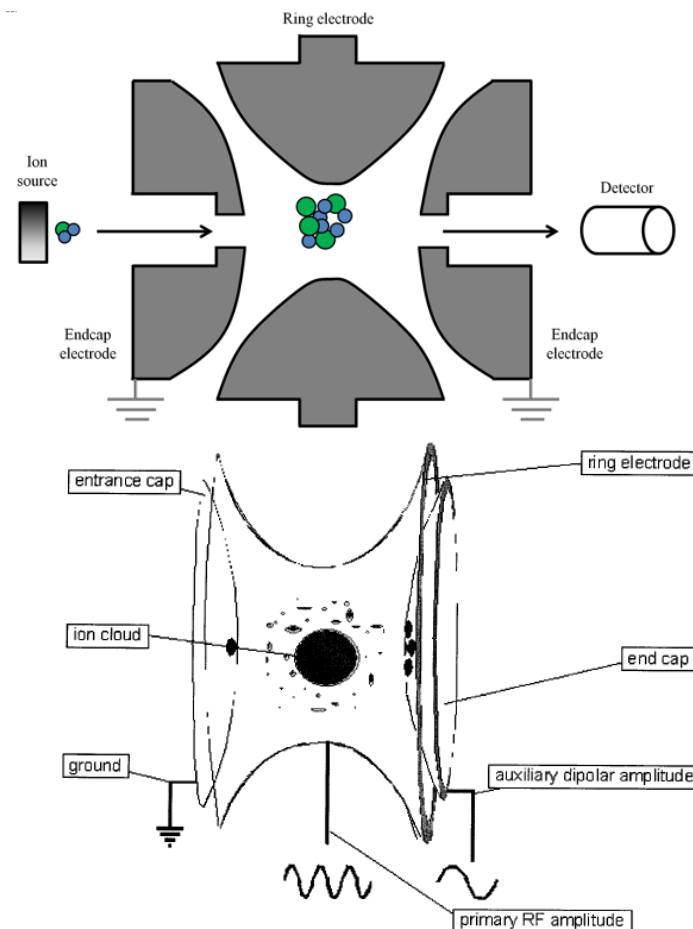


Principles of mass analyzers

Quadrupole: a) straight (Q)



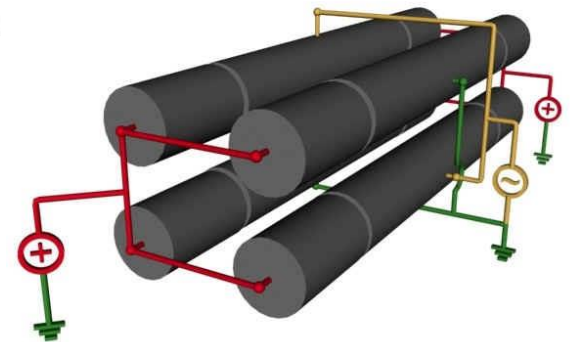
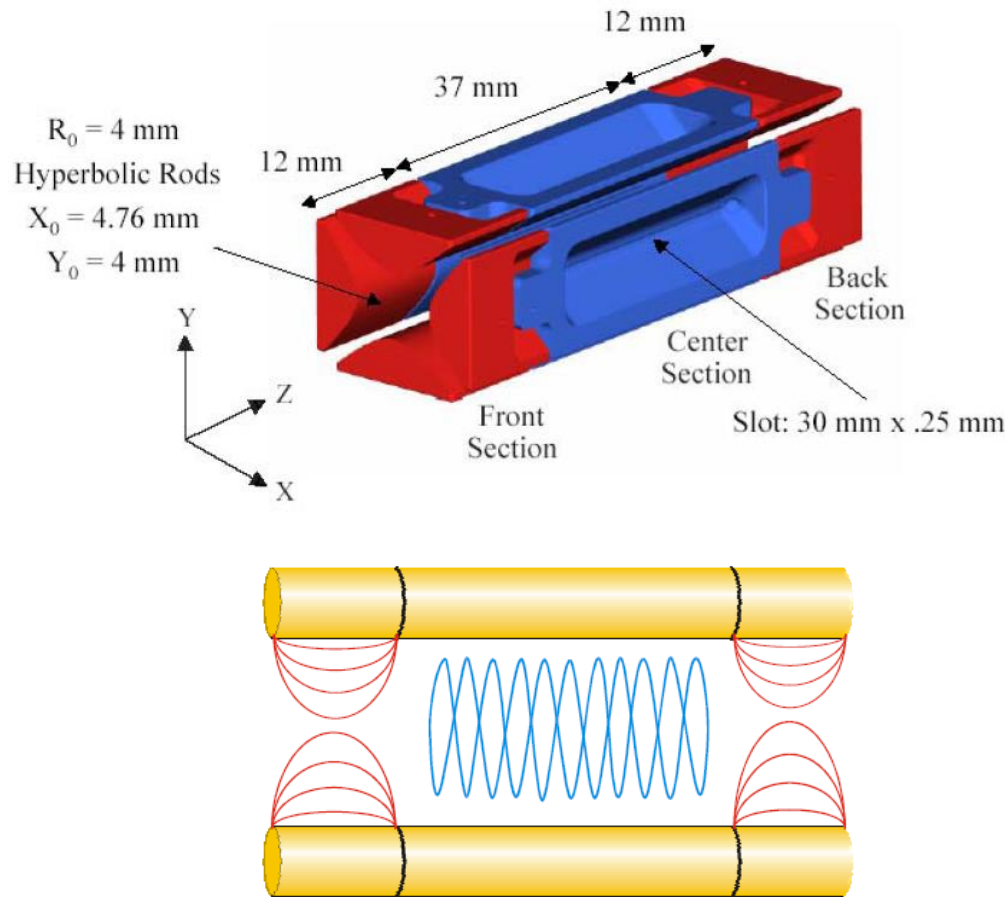
b) spherical - 3D trap (IT)



FIGURES: Agilent Technologies nad Thermo

Principles of mass analyzers

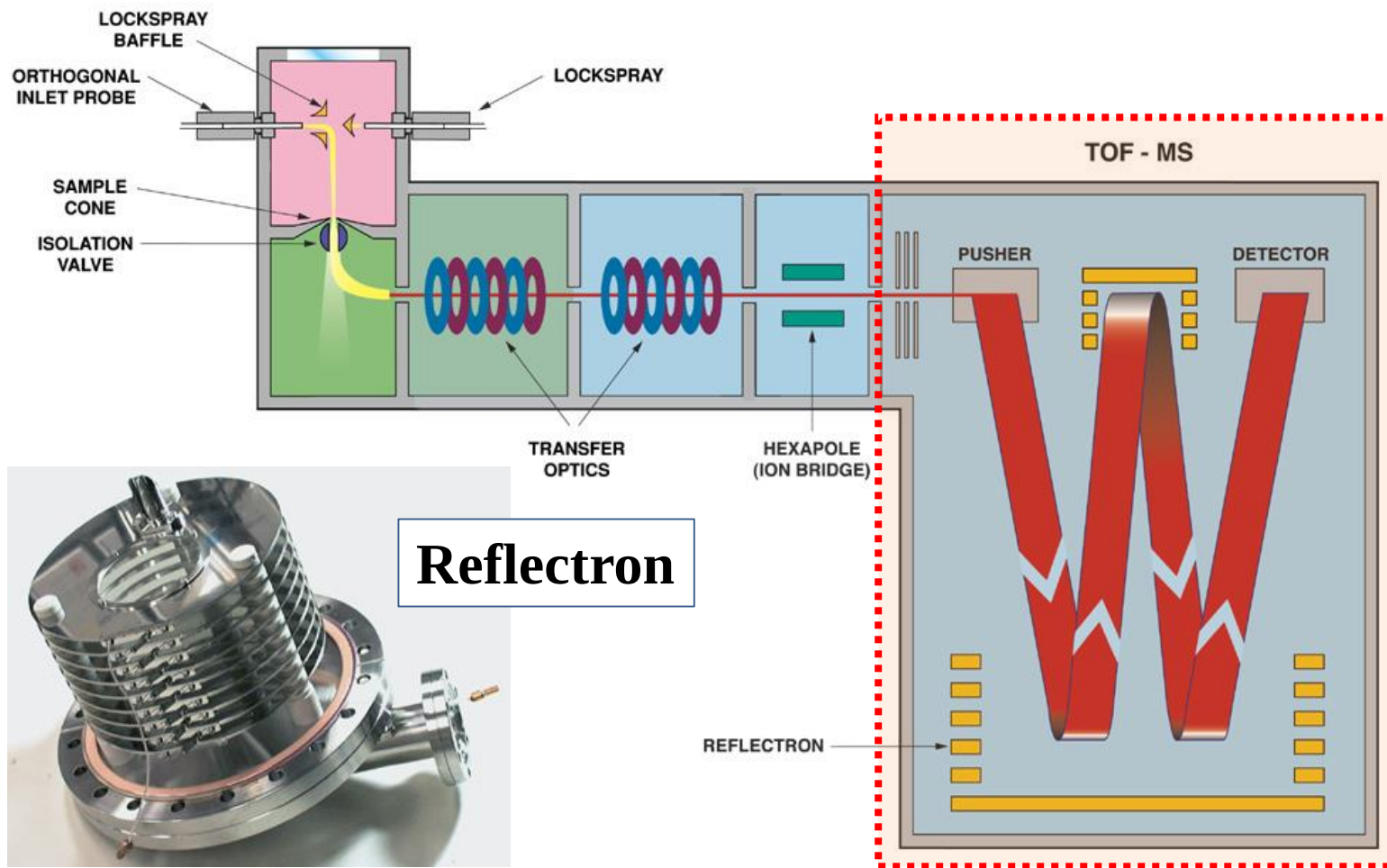
Quadrupole: linear ion trap (segmented quadrupole) 2D trap (IT)



FIGURES: AB Sciex

Principles of mass analyzers

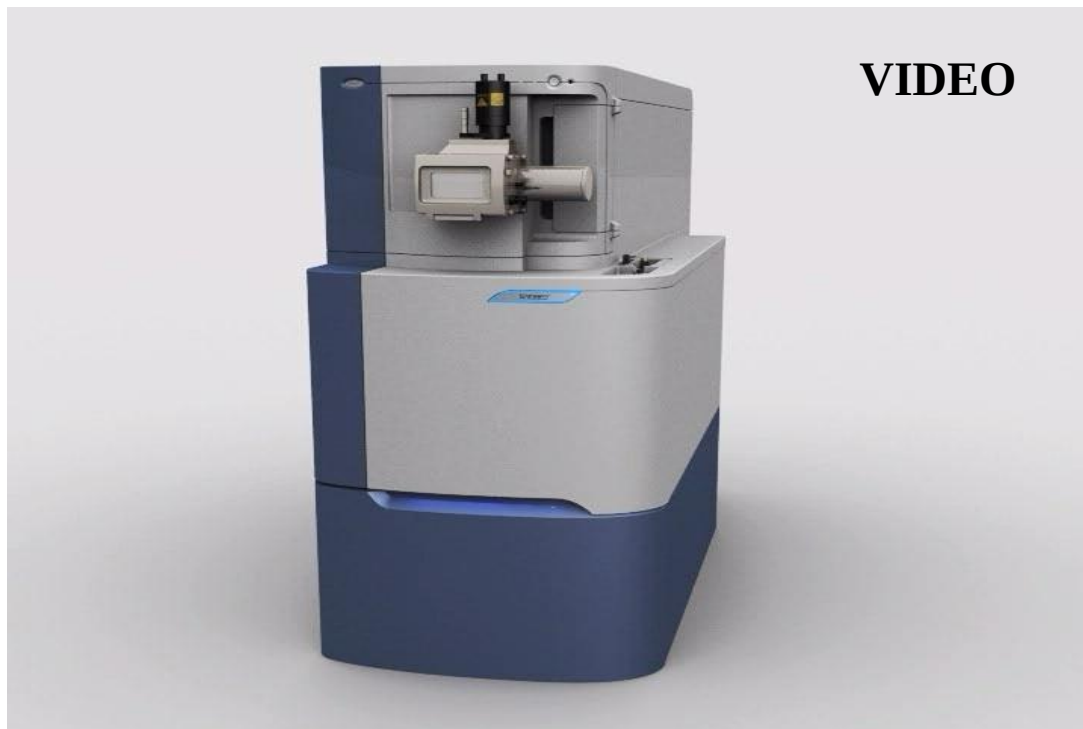
Time Of Flight (TOF)



FIGURES: Waters

Principles of mass analyzers

Time Of Flight (TOF)

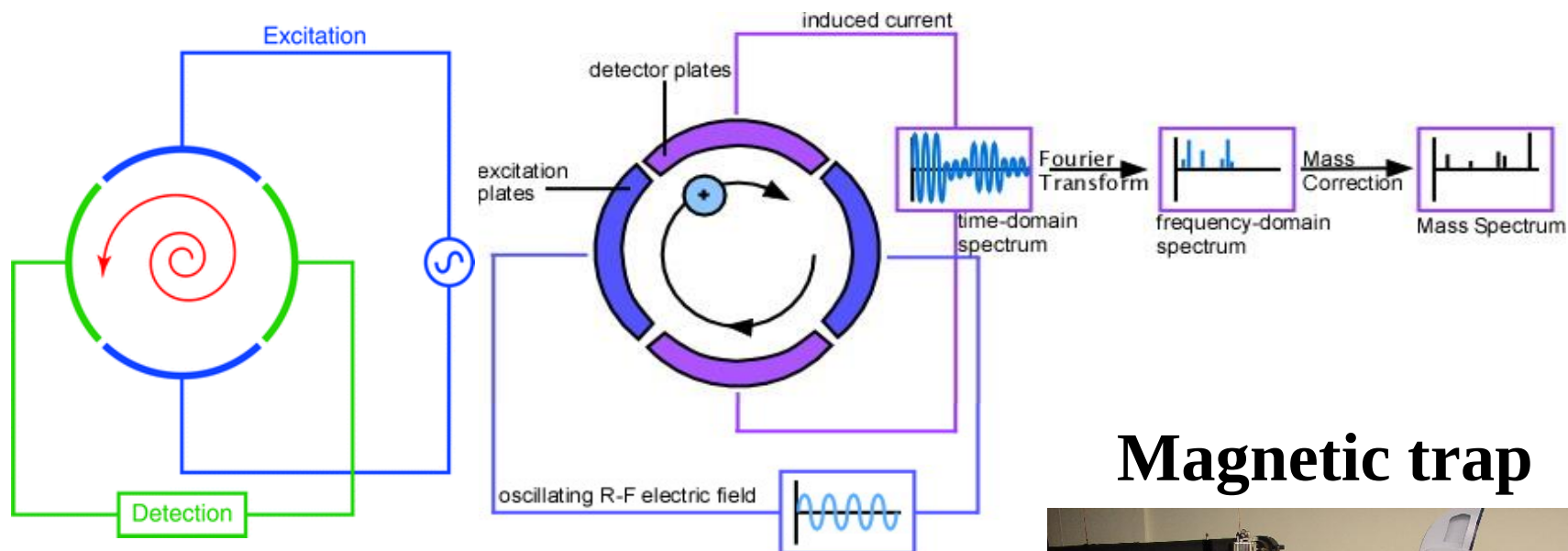


**Ion Mobility (travelling wave)
- TOF MS (with reflectron)**



Principles of mass analyzers

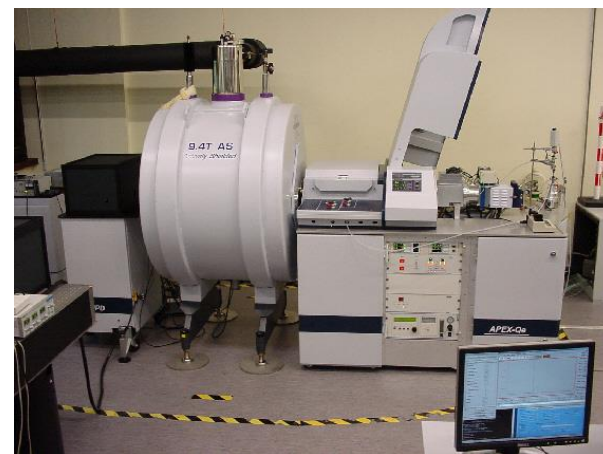
Fourier Transform Ion Cyclotron (FT-ICR)



Mark P. Barrow, William I. Burkitt
and Peter J. Derrick
Analyst, 2005, 130, 18-28

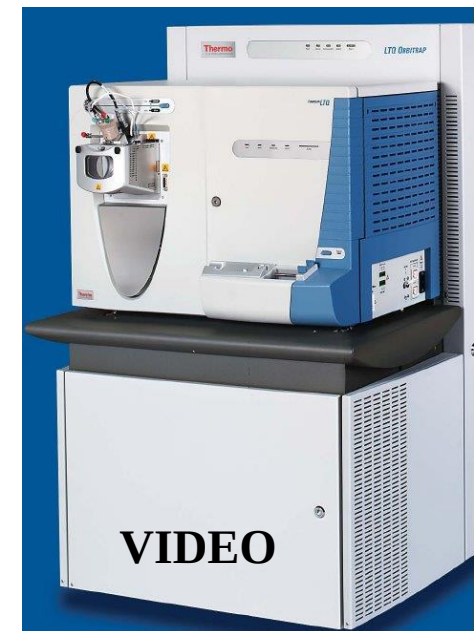
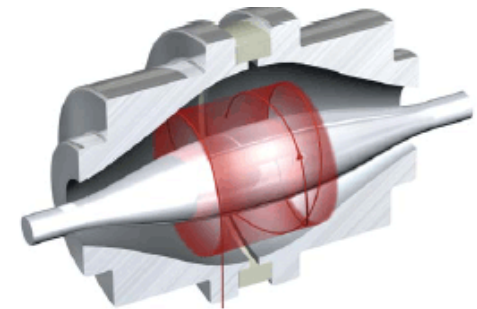
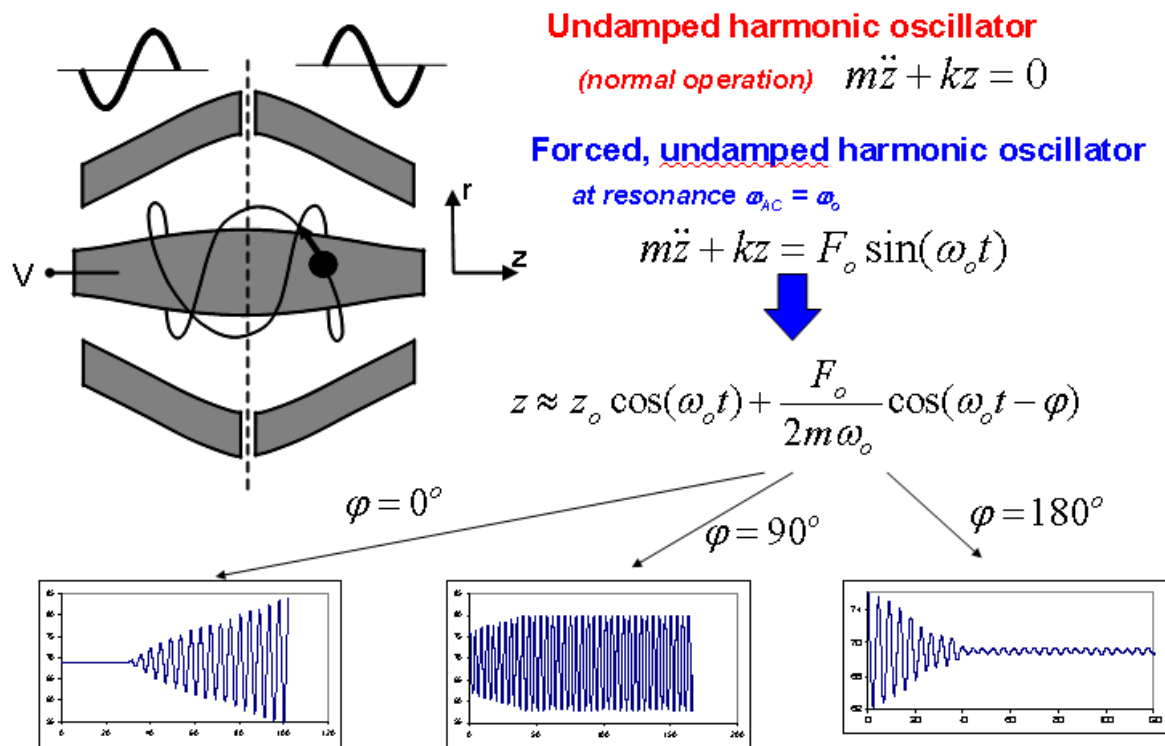
Nicole James
© Dunnivant & Ginsbach, 2008

Magnetic trap



Principles of mass analyzers

Electrostatic orbital trap - ORBITRAP



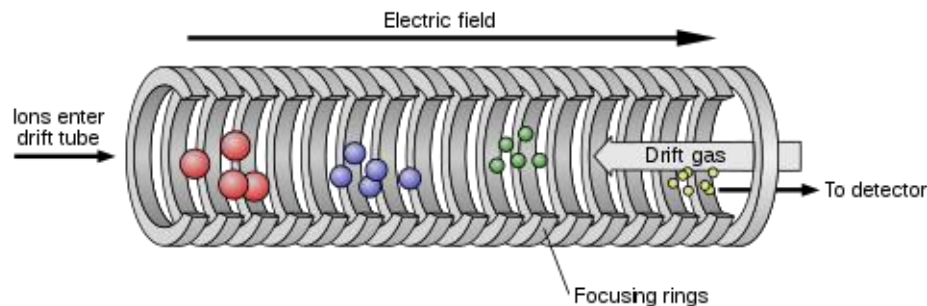
VIDEO

FIGURES: Thermo

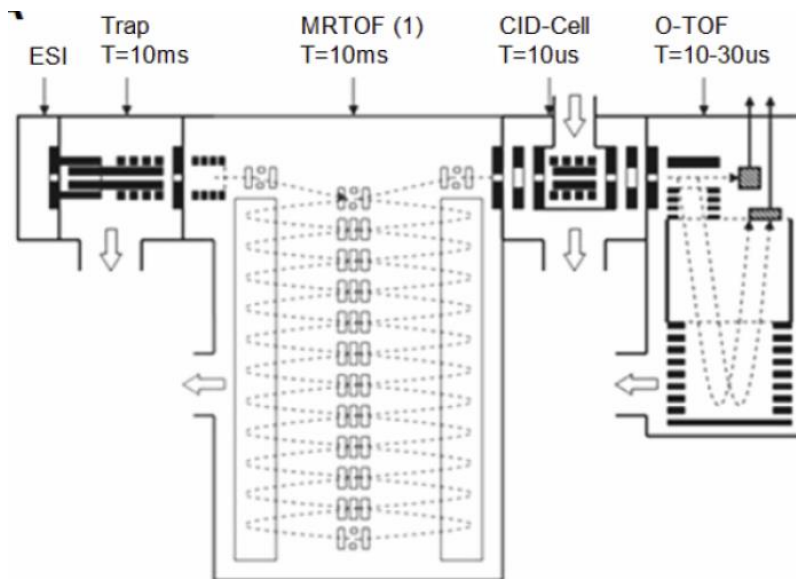


Principles of additional MS parts

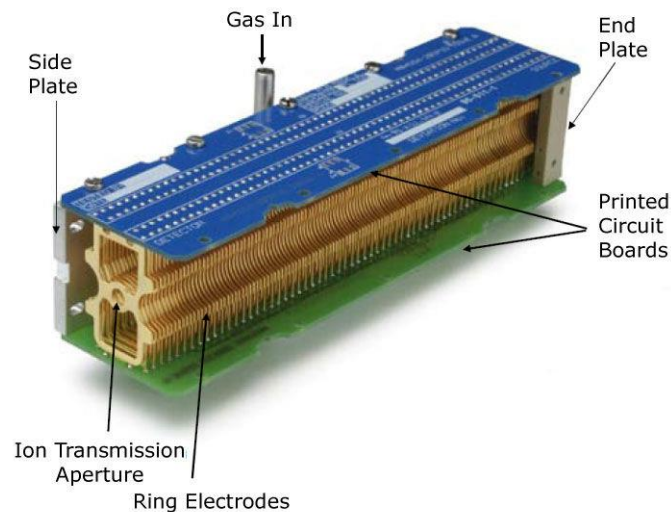
Ion mobility



Multi-reflective TOF



Traveling wave



FIGURES: Agilent Technologies, LECO, Waters,

Comparison of parameters of mass analyzers*

ANALYZER	Max. RP FWHM	Accuracy <i>ppm</i>	Range <i>m/z</i>	Scan speed	Price
Magnetic sector	10^5	< 5	10^4	low	high
Q	10^3	≈ 1000	10^3	medium	low
IT	10^3	≈ 1000	10^4	medium	medium
TOF	10^4	< 5	10^4	very high	medium
Orbitrap	10^5	< 3	10^3	high	high
FT-ICR	10^6	< 1	10^4	low	very high

* All parameters vary depending on technical progress and experimental settings



Realization of multistage (multiple) mass analysis

➤ multistage or multiple mass analysis

→ $MS/MS = MS^2 = \text{tandem MS}$; $MS/MS/MS = MS^3$

Principle: in the first stage we select primary (parent/precursor) ions, which break down in collision cell to secondary (daughter/product/fragment) ions, which are analyzed in the second stage. It can be followed by process repetition for MS^3 realization etc.

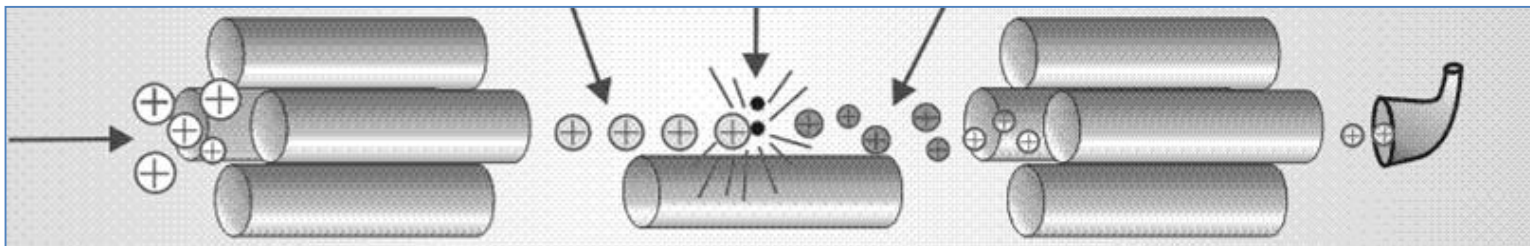
Technical realization:

- a) **in space** – serial combination of the same or different mass analyzers (QqQ, Q-TOF)
(chaining of experiments in space)
- b) **in time** – takes place in one mass analyzer by timed parameter changes (IT)
(chaining experiments in time)



Realization of multistage (multiple) mass analysis

a) MS² in space

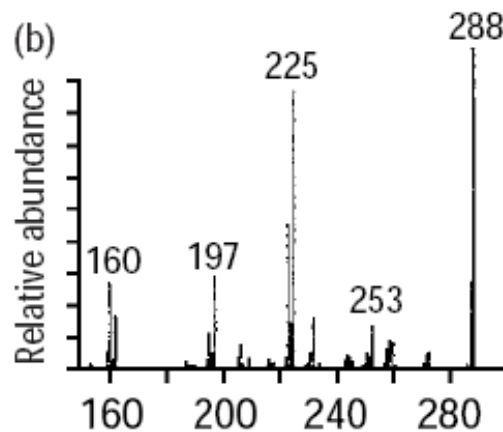
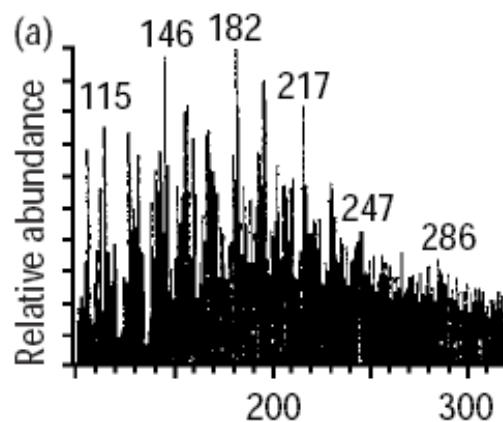
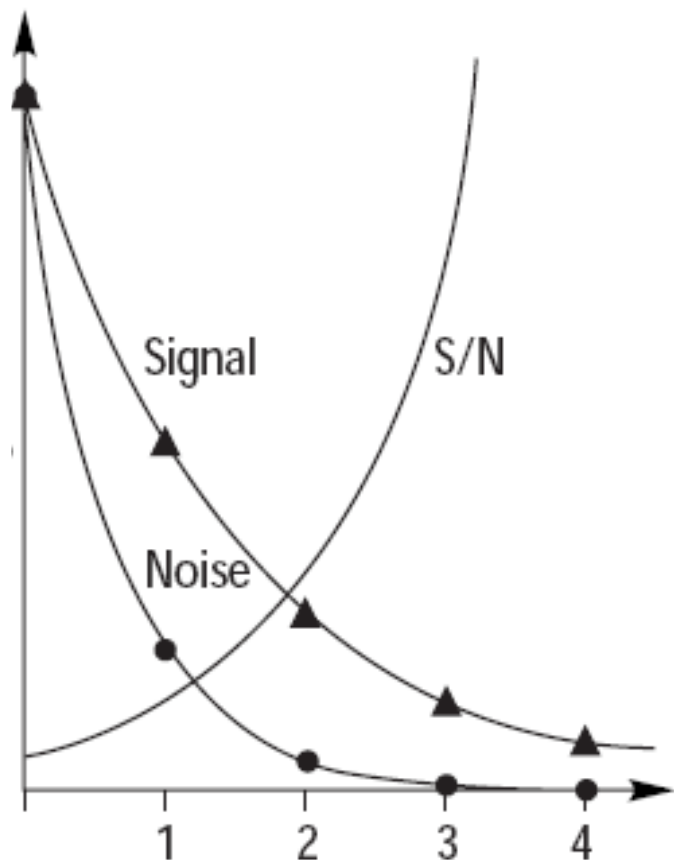


b) MS² in time



Realization of multistage (multiple) mass analysis

Noise reduction with number of MS analyzes $\rightarrow$ S / N increase



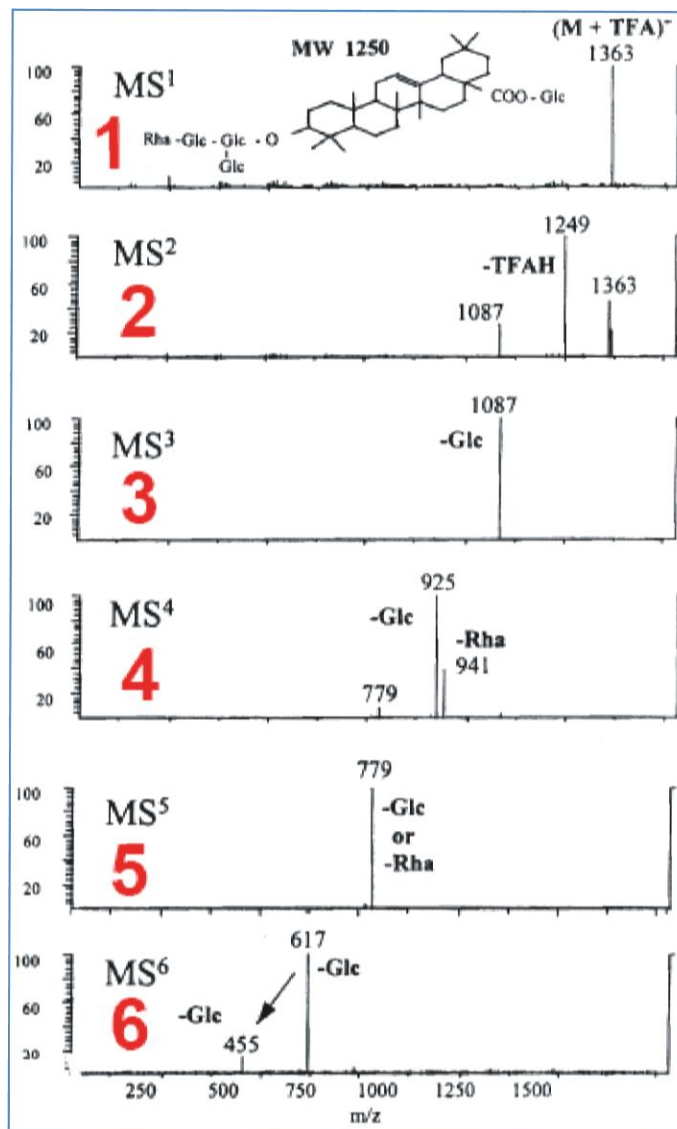
MS¹

MS²



Realization of multistage (multiple) mass analysis

Gradual fragmentation
in ion trap



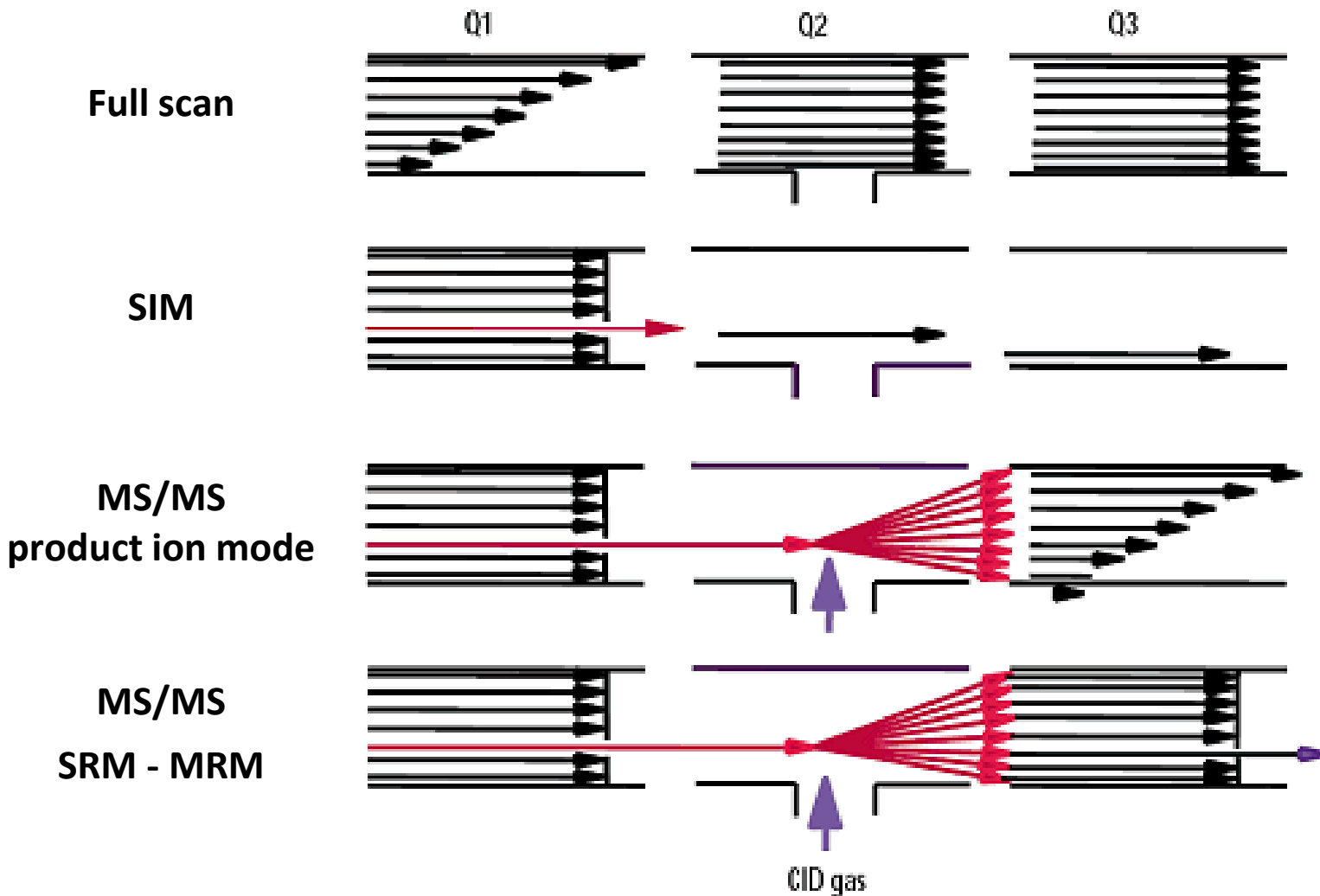
MS¹

MS⁶



Realization of multistage (multiple) mass analysis

Available measurement modes when using tandem MS



Ion detection in MS

**Ion detector in MS records numbers of ions,
which pass through the mass analyzer**

➤ **the incident ion induces electron(s) whose current is amplified by impacts on another surface of electron multiplier (HED - high energy dynodes)**

a) plate dynodes

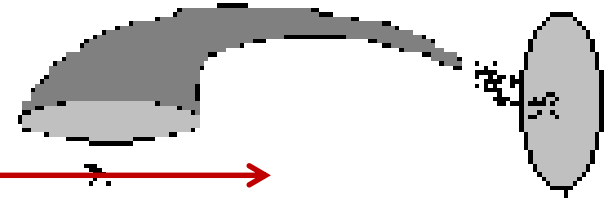
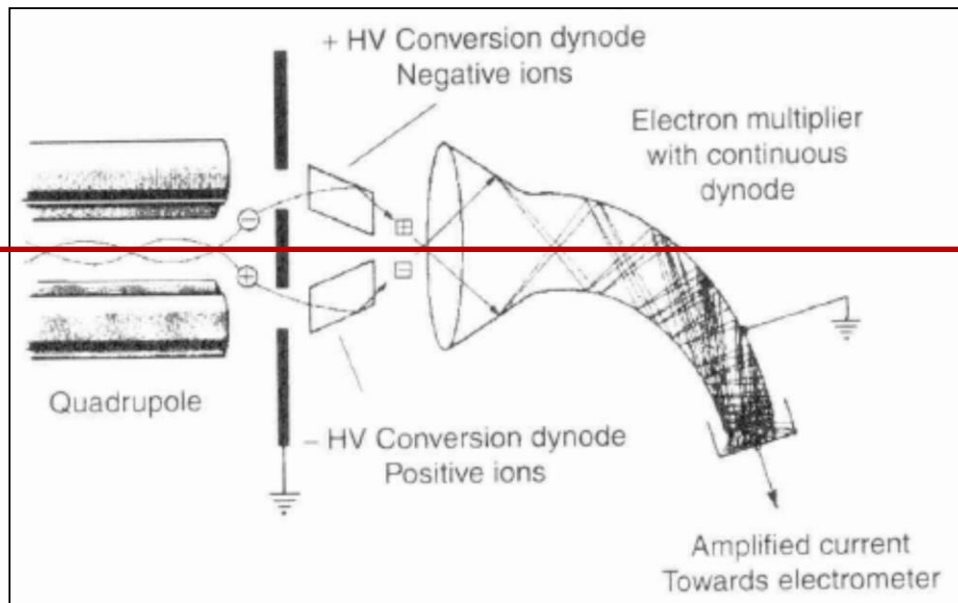
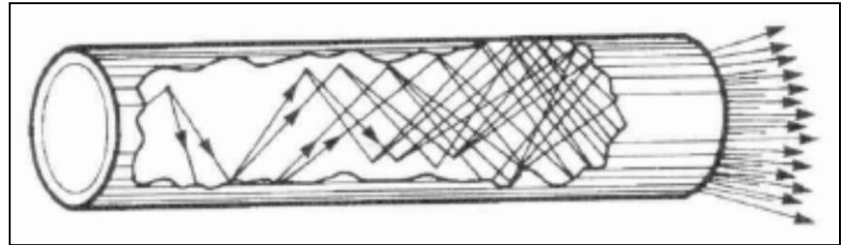
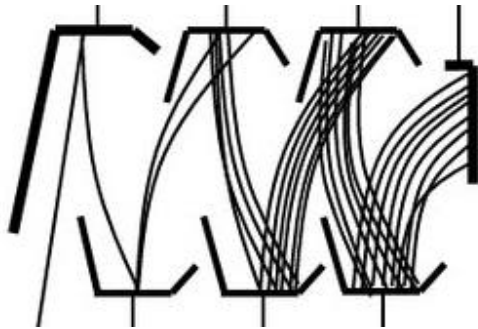
b) continuous dynodes - tubes (straight, curved)

c) parallel dynodes - microchannel plate

**Parameters: amplification, lifetime, detection of $[\pm]$ ions,
detection of unwanted fragments**



Ion detection in MS



Ion detection in MS

