

# Mass spectrometry – part II

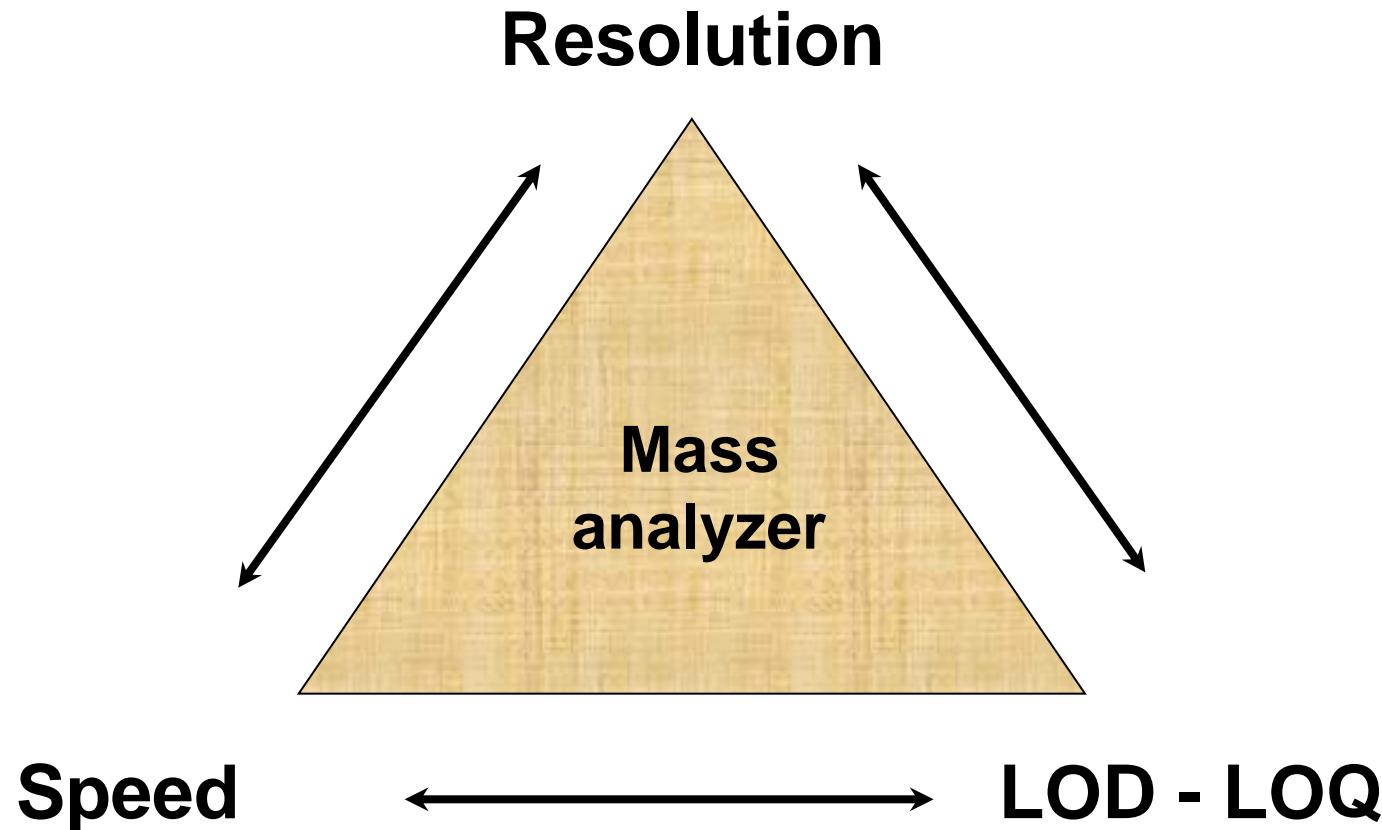
## SEPARATION and DETECTION of IONS

<i>Ion separation</i>	<i>Ion detection</i>
<b>Spectrum quality</b> - stability, range - resolution - accurate mass <b>true mass</b>	<b>LOD / LOQ</b>  <b>Stability</b>
<b>Speed of measurement</b>	<b>Lifetime / Durability</b>
<b>LOD / LOQ</b>	

# Principles of ion separation

- \* separation is realized by mass analyzer (mass filter)
- \* mass analyzer = part of mass spectrometer,  
which distinguishes ions  
according to their value of  $m/z$
- \* *basic parameters of mass analyzers:*
  - a) range of  $m/z$
  - b) scan speed, switching of scan mode
  - c) resolution power X resolution
  - c) accurate mass X true mass
  - e) multistage mass analysis ( $MS^2$  etc.)

# Contradictory of parameters of mass analyzer



# Parameters of mass analyzers (1)

\* range of m/z:

- a) lower limit: from technical "0" or from higher values
- b) upper limit: up to a certain maximum value  
(singly charged X multiply charged ions)

\* scan speed and possibility of switching of scan modes:

- a) speed = number of scans per second:  
    < 1 - slow  
    1 to 10 - medium  
    10 to  $\approx$  500 - fast
- b) mode: full scan, segment scan, SIM, SRM, MRM

## Parameters of mass analyzers (2)

\* resolving power - RP ( $10^3$  –  $10^6$ );  $RP = 1/R$

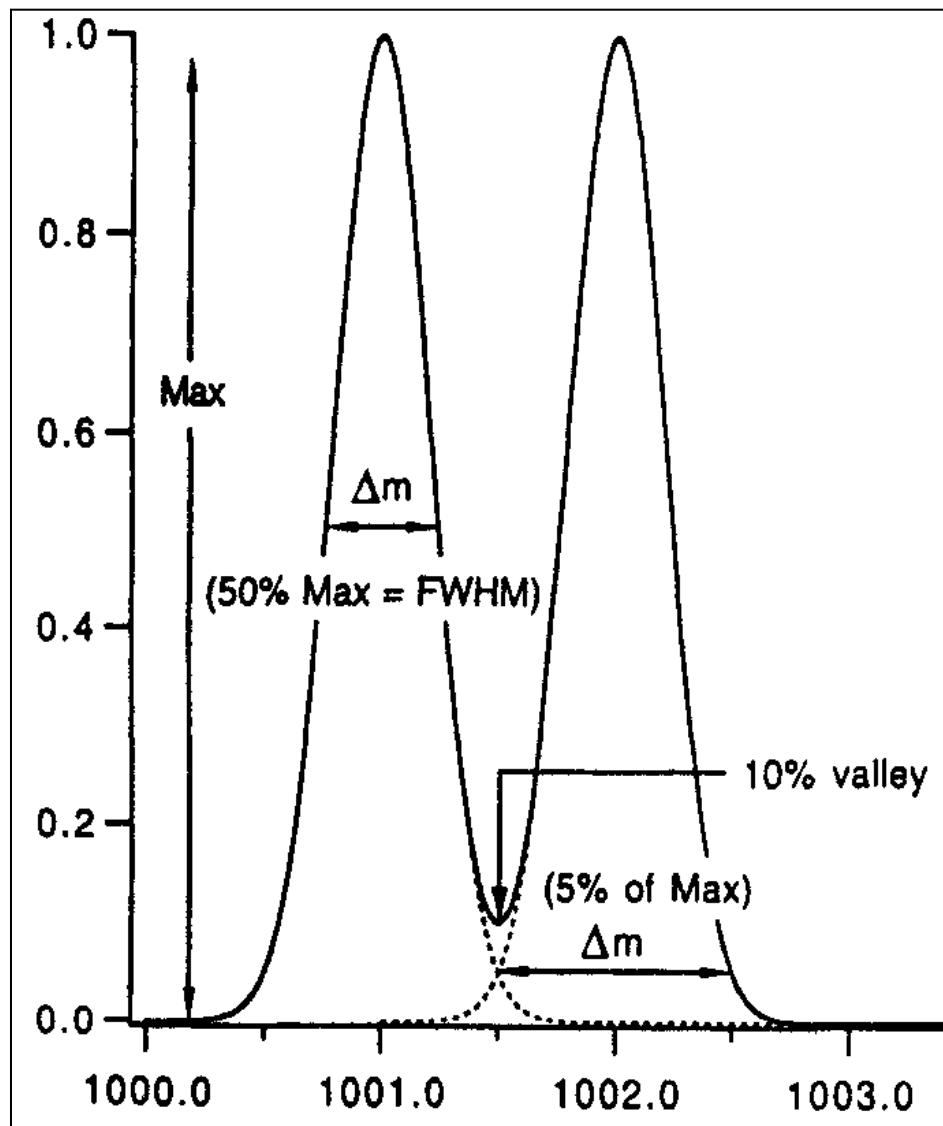
\* a measure of the ability of mass analyzer to distinguish between close m/z values, i.e. what the smallest difference can be record

Relative measure - depends on m/z value

a) definition for two close ions:  $RP = m_1 / (m_1 - m_2)$ ;  $z_1, z_2 = 1$   
peaks  $m_1$  and  $m_2$  with 10 %  
overlaid at the same height

b) definition for one ion:  $m / \Delta m$  (or  $t / 2 \Delta t$  for TOF)  
based on concept of FWHM (full width at half maximum)

# Parameters of mass analyzers (3)



## Parameters of mass analyzers (4)

\* resolution - R ( $10^{-1}$  –  $10^2$  ppm);  $R = 1/RP$

\* a measure of the ability of mass analyzer to distinguish between close  $m/z$  values, i.e. what the smallest difference can be record

**Relative measure - depends on  $m/z$  value**

Expressed in ppm;  $R = m_1 - m_2 / m_1$

## Parameters of mass analyzers (5)

\* accurate mass

\* a measure of match between measured and calculated (exact)  $m/z$

\* a measure of the ability of mass analyzer to determine the correct  $m/z$

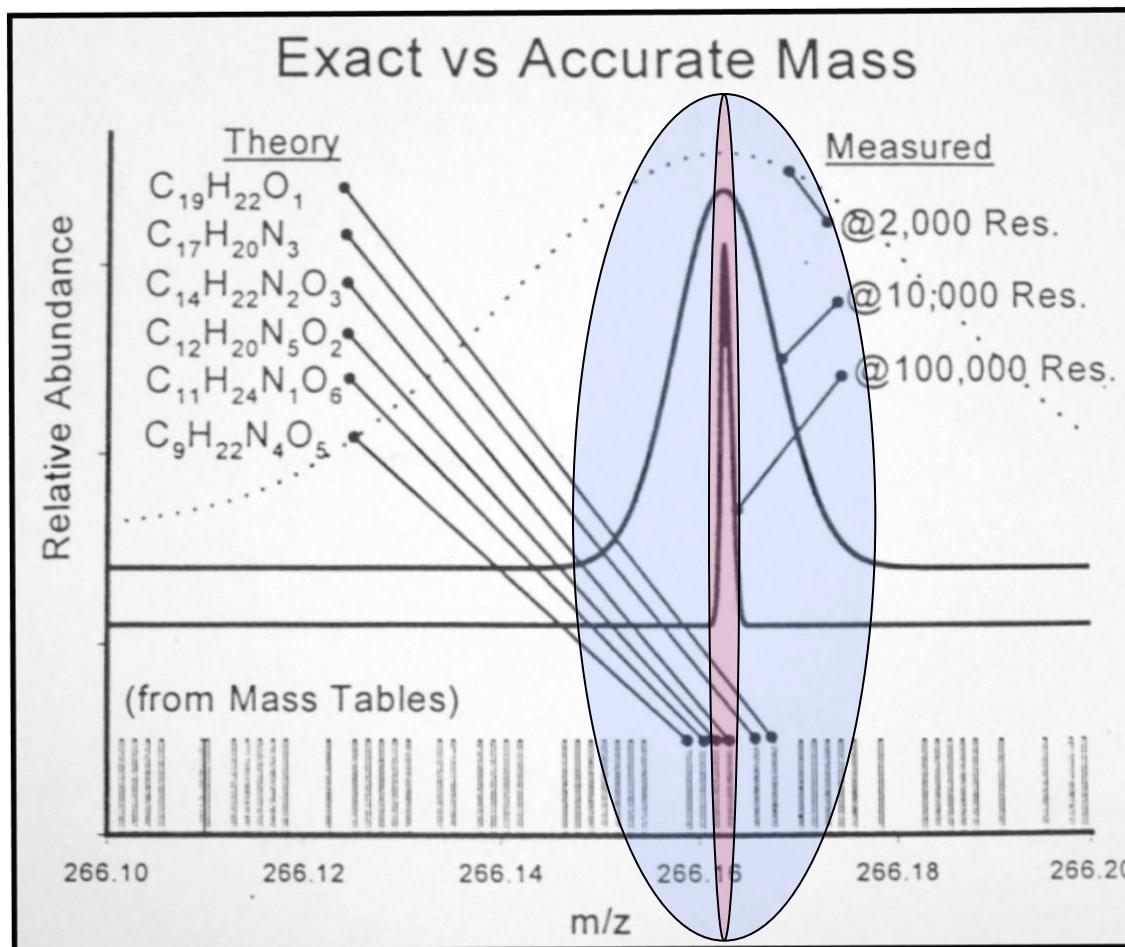
Relative measure - depends on  $m/z$  value

Expressed in ppm:  $\text{ppm} = 10^6 \cdot \Delta m / m$

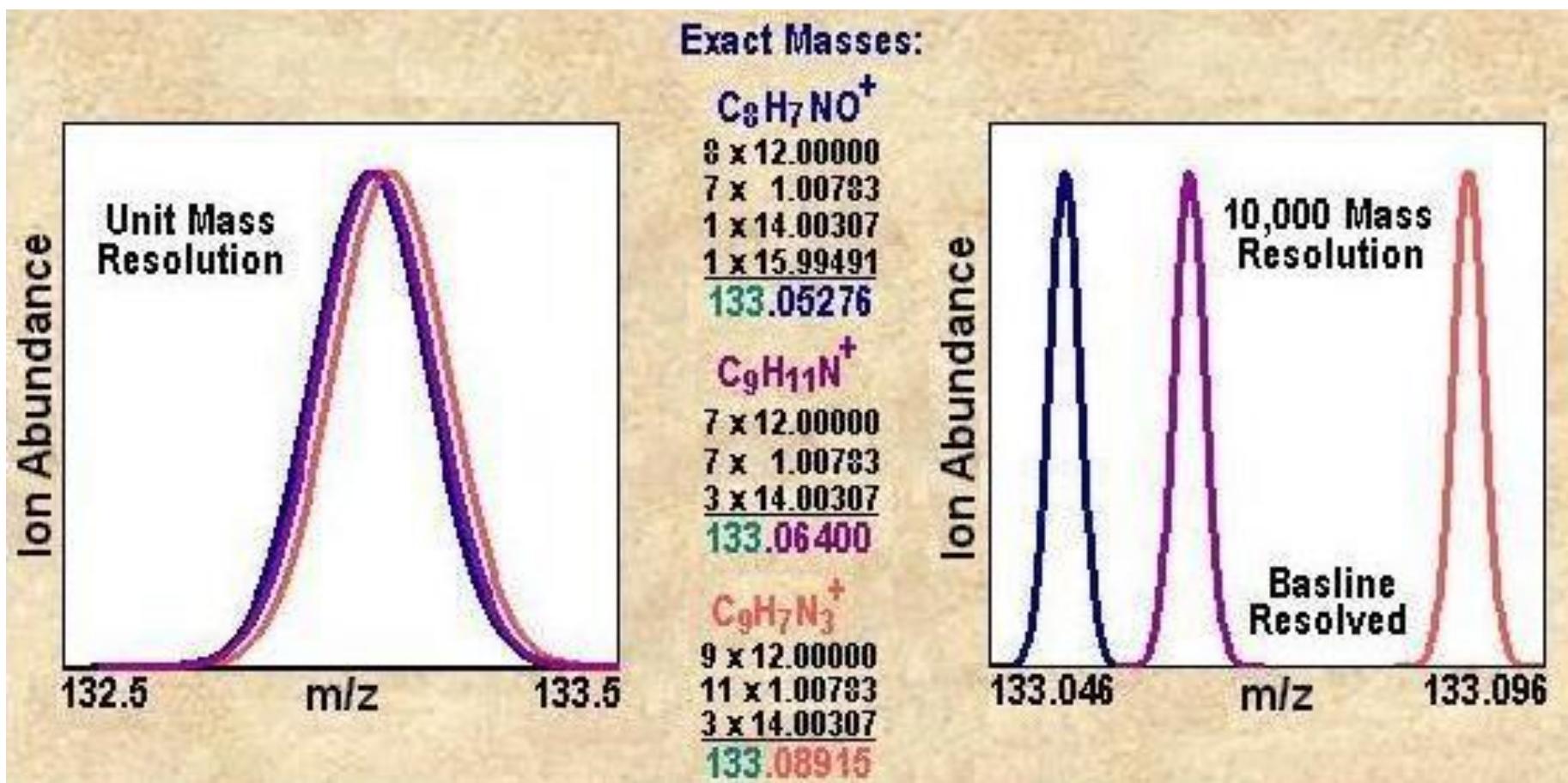
$10^6 \cdot (\text{exact value} - \text{measured value} / \text{exact value})$

e.g.:  $\text{ppm} = 10^6 \cdot 212,14124 - 212,14072 / 212,14124 = 2,45 \text{ ppm}$

# Parameters of mass analyzers (6)



# Parameters of mass analyzers (7)



# Parameters of mass analyzers (8)

## Examples

Resolving power (RP) = 20000 (usually given as FWHM resolution)

- for  $m/z = 1000$  ion distinguishing with difference  $\pm 0.05$  ( $1000/20000 = 0.05$ )
- for  $m/z = 100$  ion distinguishing with difference  $\pm 0.005$  ( $100/20000 = 0.005$ )

Resolution (R) for RP above is 50 ppm ( $1/20000 = 50 \cdot 10^{-6}$ )

## Accurate mass

- a) high resolution with accuracy 3 ppm  
 $100 \pm 0.0003$  or  $1000 \pm 0.003$  etc.
- b) Low resolution with accuracy 100ppm  
 $100 \pm 0.01$  or  $1000 \pm 0.1$  etc.

## Parameters of mass analyzers (9)

- \* multistage or multiple mass analysis  
→  $\text{MS/MS} = \text{MS}^2$ ,  $\text{MS/MS/MS} = \text{MS}^3$  etc.

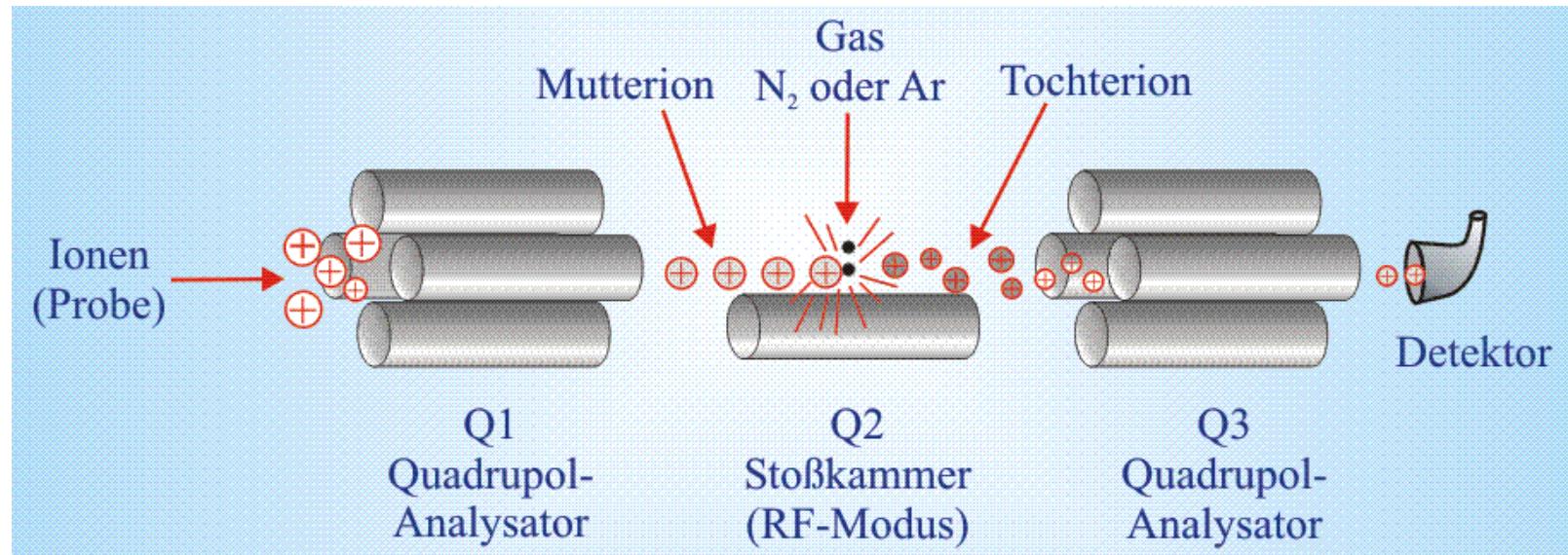
a) in space – combination of the same or various mass analyzers  
in series:  $Q_1 \rightarrow Q_2 \rightarrow Q_3$  vs.  $Q_1 \rightarrow T\text{-wave} \rightarrow Q_2$

analyzer → collision cell → analyzer

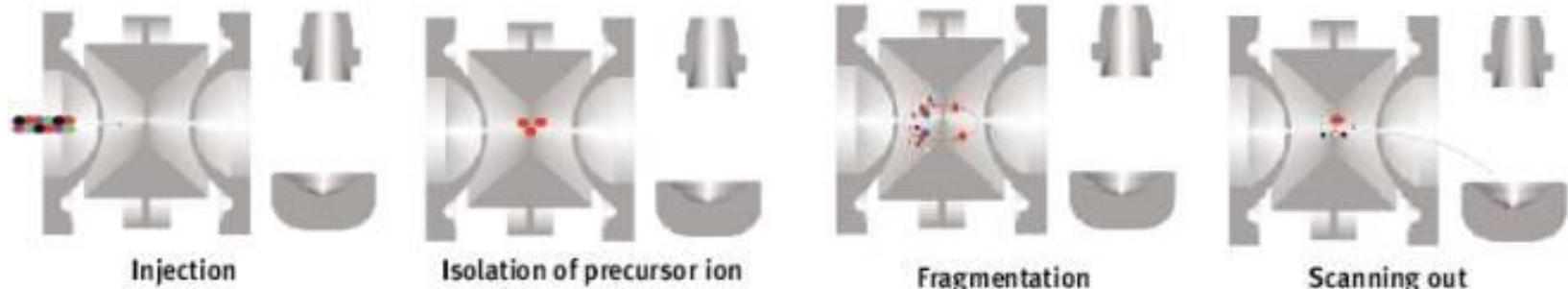
b) in time – realized in the same space (analyzer)  
with changes of parameters of the analyzer - ion trap  
primary analysis → collision → secondary analysis

# Parameters of mass analyzers (10)

## In space

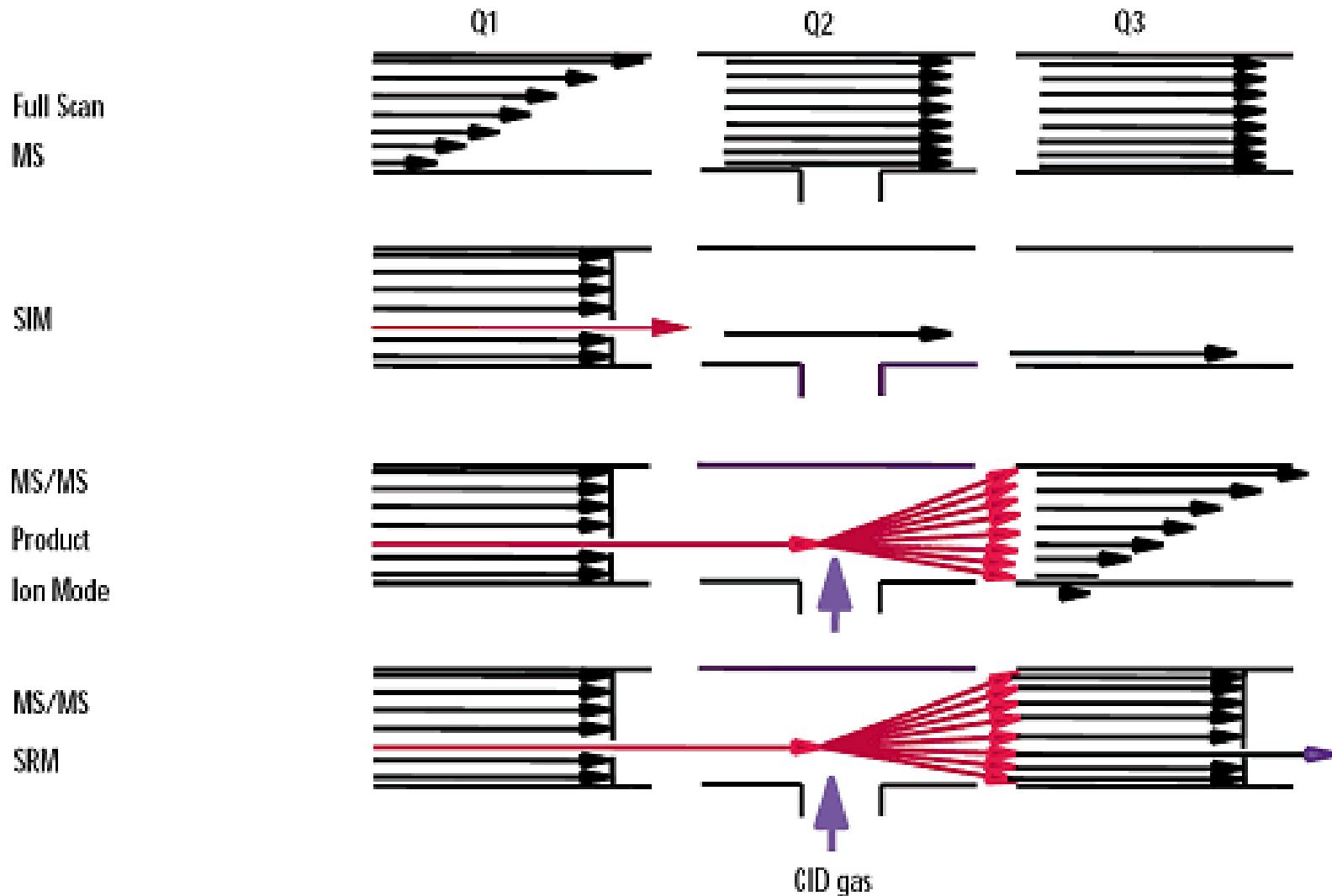


## In time



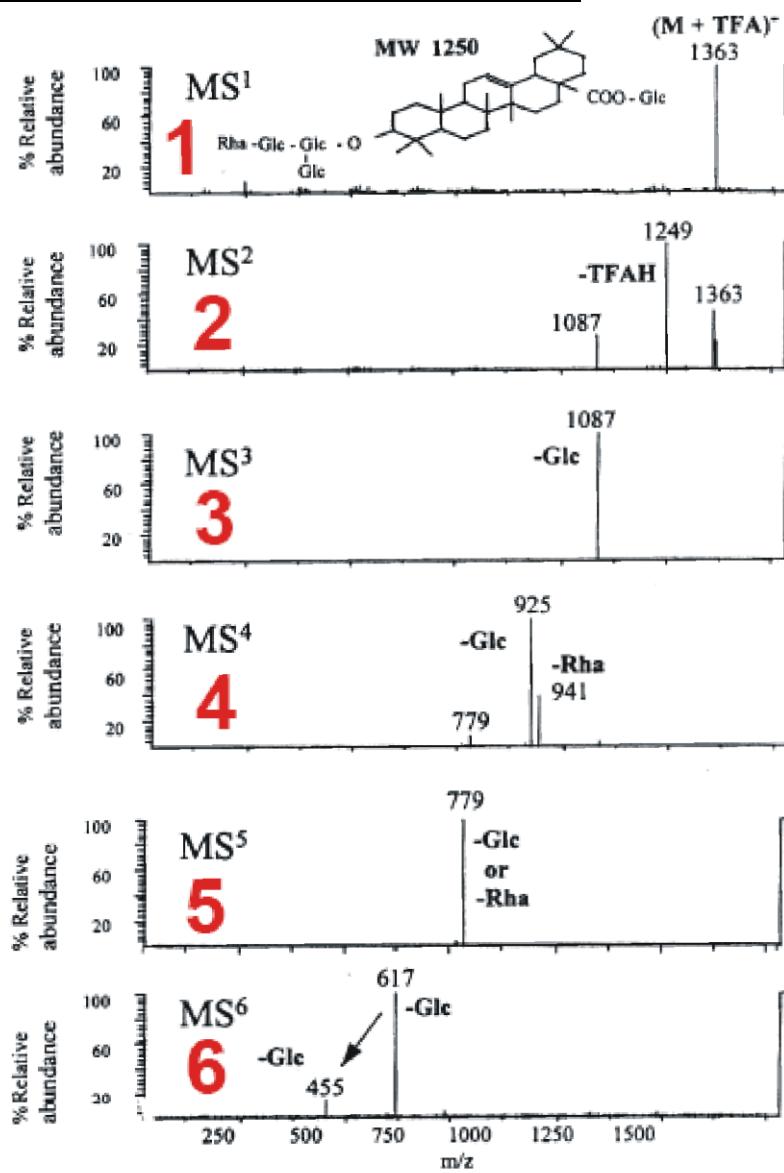
# Parameters of mass analyzers (11)

## Scan functions of triple quadrupole

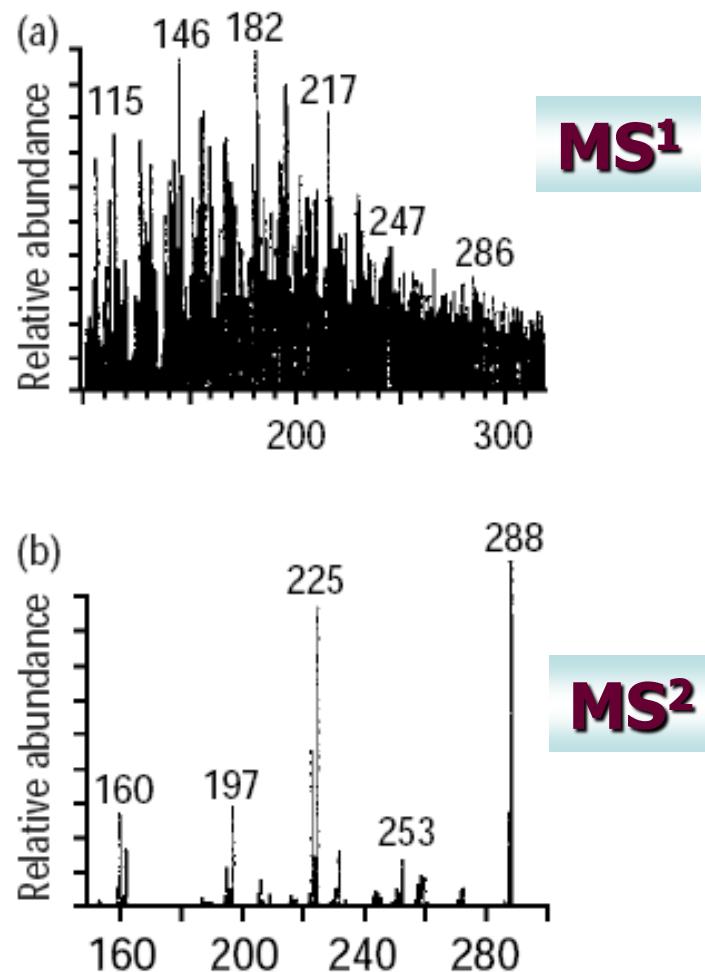
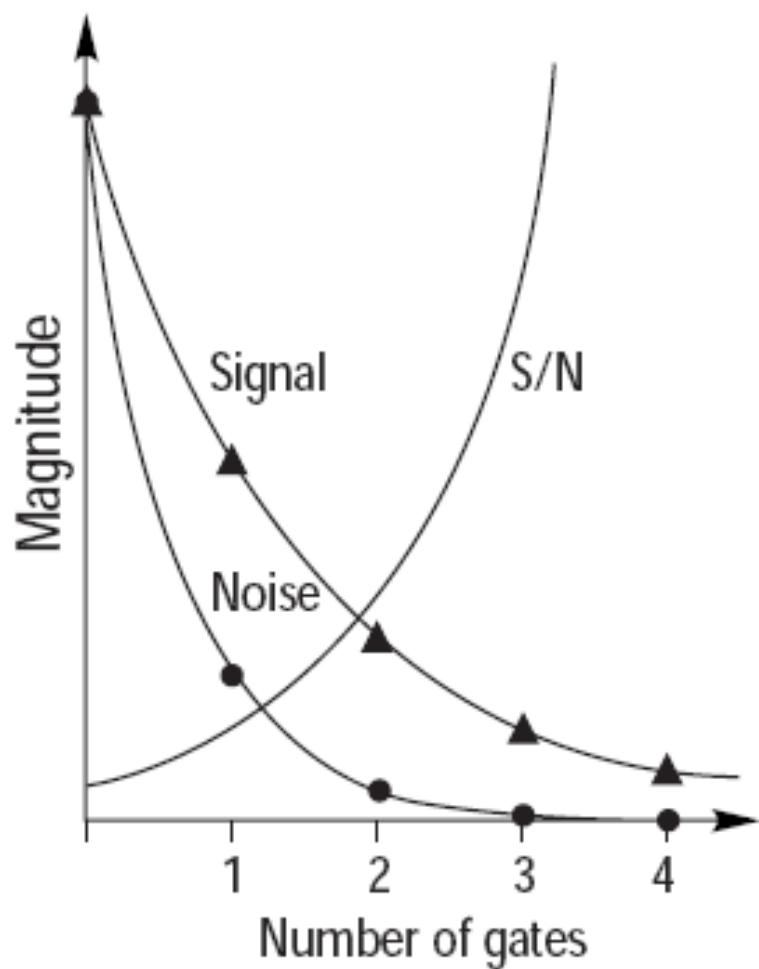


# Parameters of mass analyzers (12)

Sequential  
fragmentation  
the ion trap



# Parameters of mass analyzers (13)



# Technical principles of mass analyzers (1)

\* different principles allowing ion separation

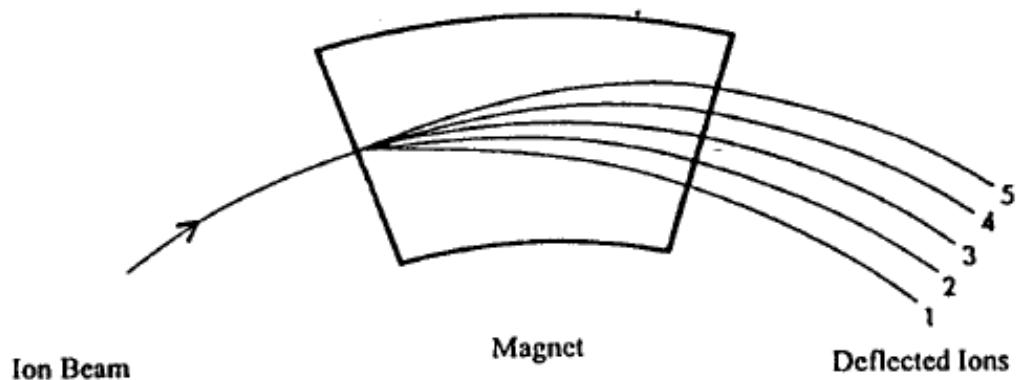
- a) magnetic analyzer with single focusation
- b) magnetic sector analyzer with dual focusation
- b) quadrupole analyzer: four rods X segment
  - ion trap (3D, 2D - linear)
- e) time of flight: straight X reflex (includes reflectron)
- f) ion mobility - transport waves (travelling wave) X drifting tube
- g) ion cyclotron resonance X orbitrap

\* mass analyzers used in ion optics applied for ion focusation:

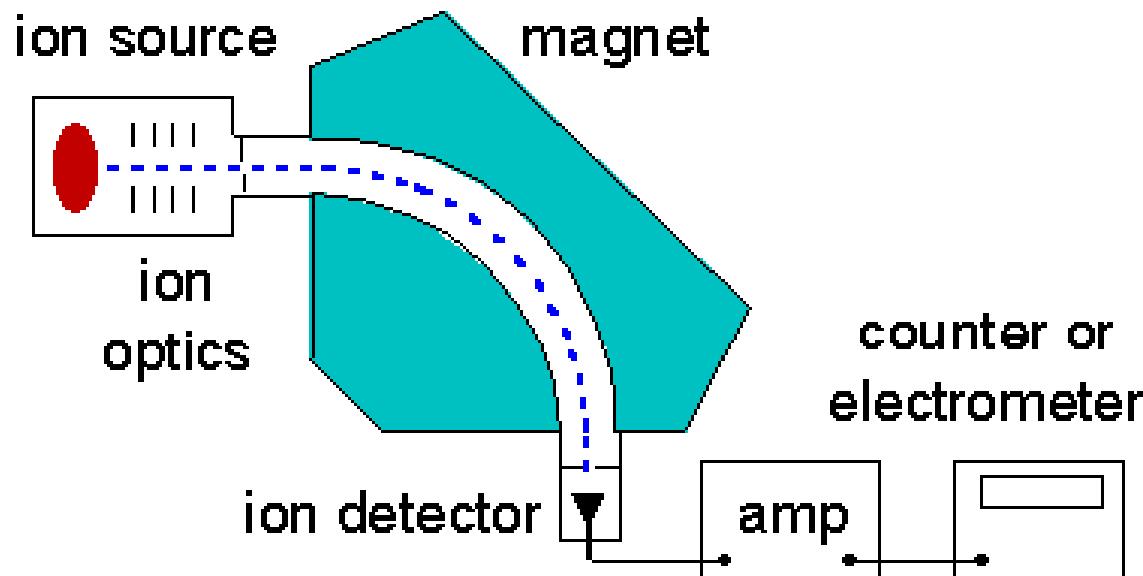
- a) multipole analyzers: hexa-, octa-pole
- b) ion optics - plates, rings

## Technical principles of mass analyzers (2)

### Magnetic sector

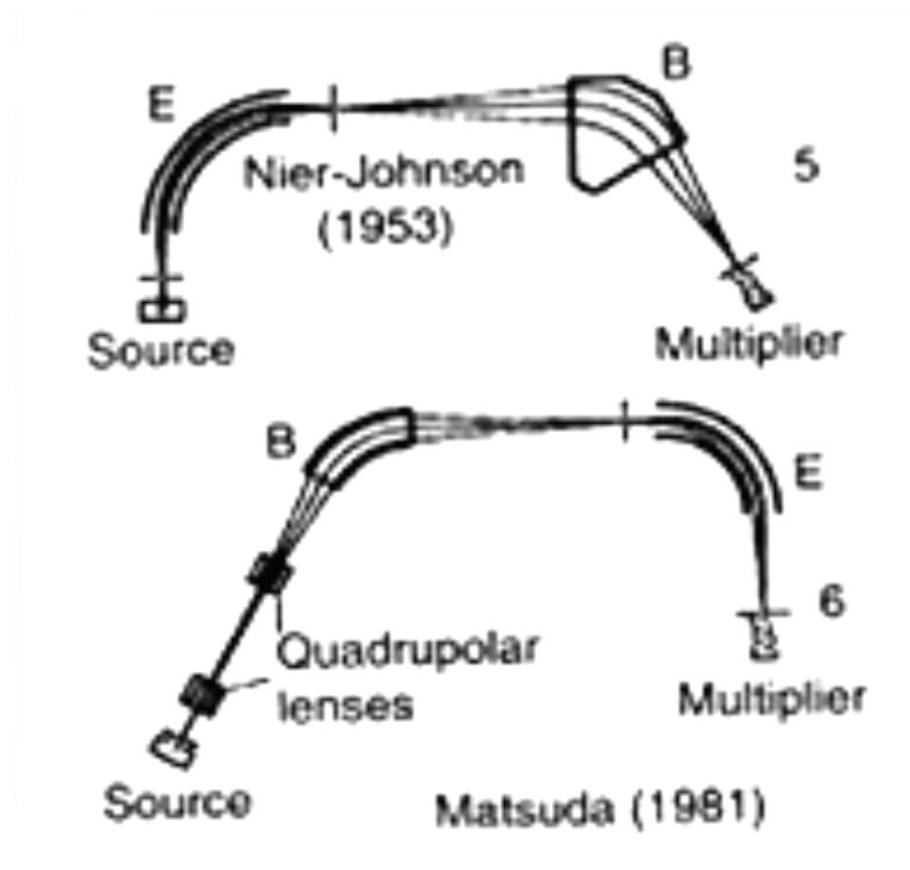


$$(m/z)_1 < (m/z)_2 < \dots < (m/z)_5$$



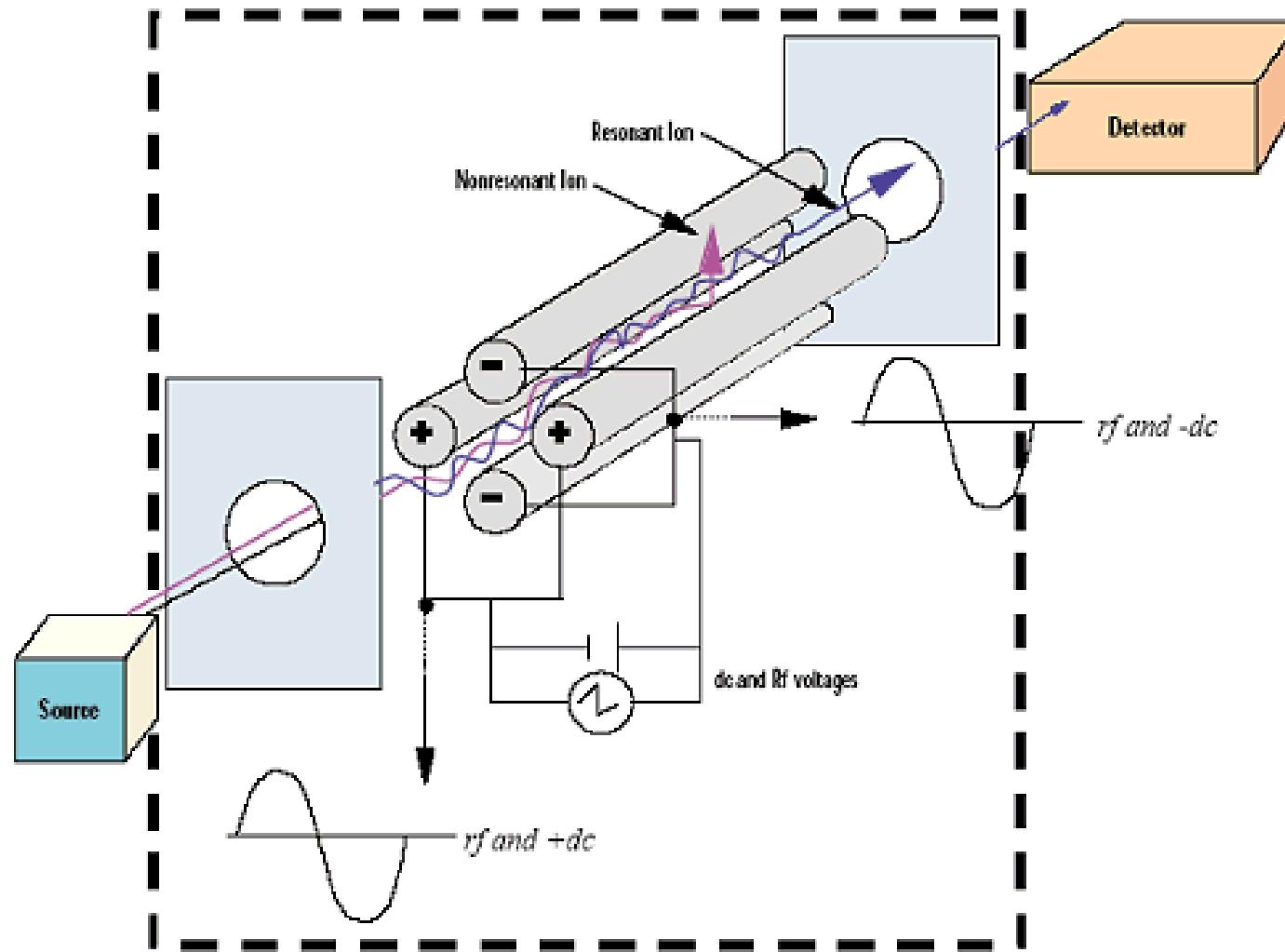
# Technical principles of mass analyzers (3)

Magnetic sector - dual focusation  
→ electrostatic / magnetic



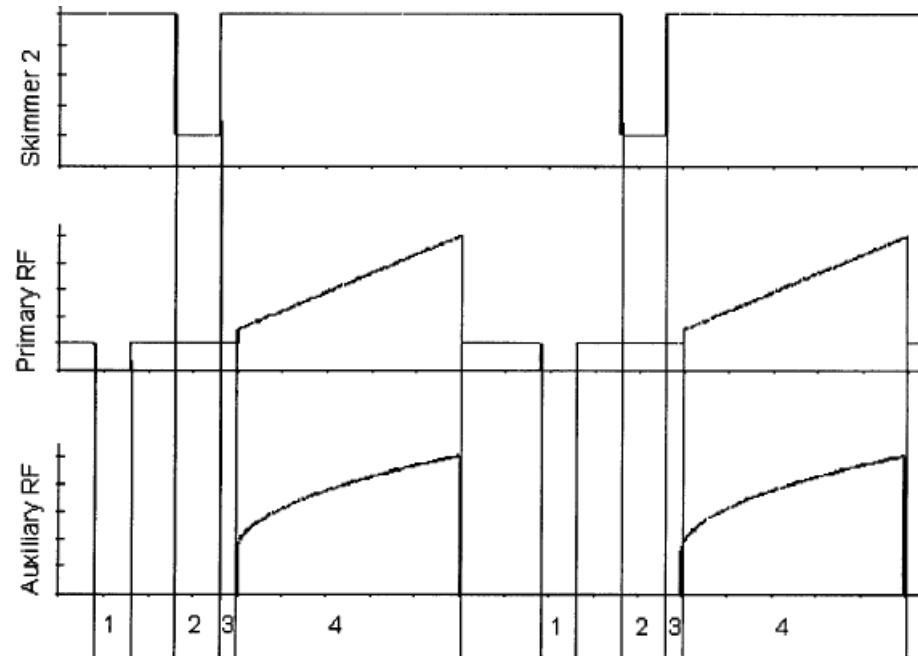
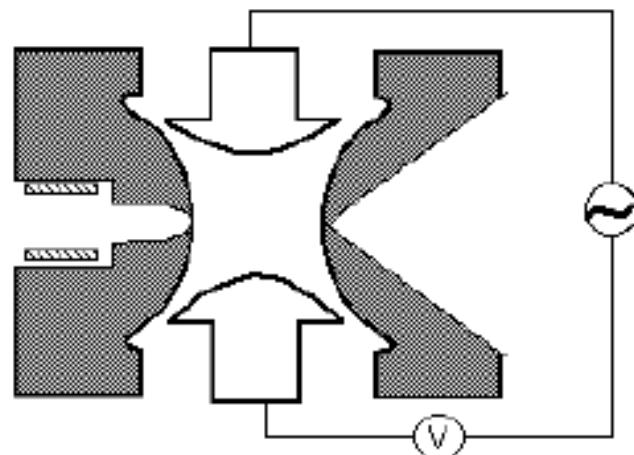
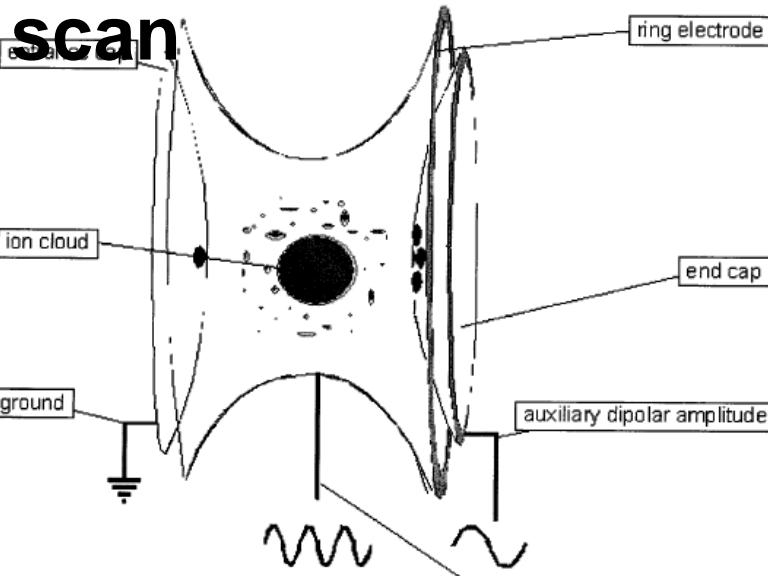
# Technical principles of mass analyzers (4)

## Quadrupole: straight - 4 rods



# Technical principles of mass analyzers (5)

## Quadrupole: ion trap (3D) - MS

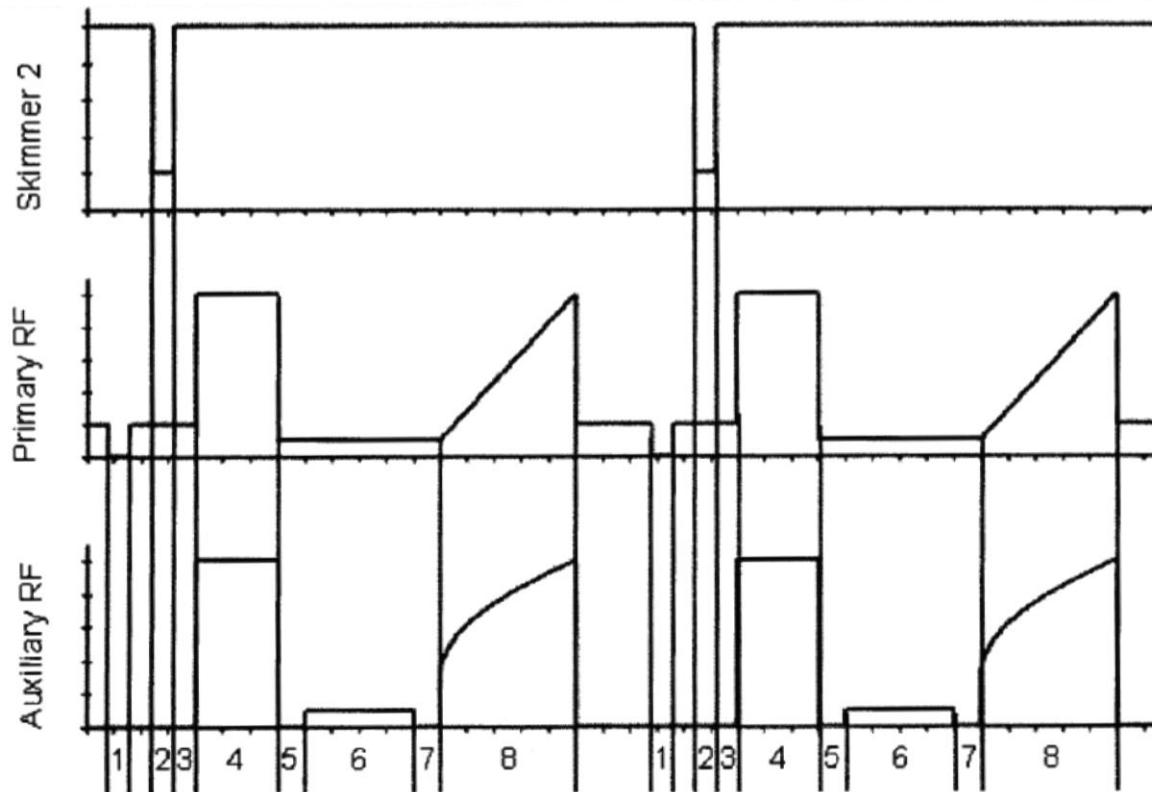


### **Ion trap - MS scan**

1. Trap emptying
2. Accumulation of ions
3. Scan dwell
4. Mass analysis

# Technical principles of mass analyzers (6)

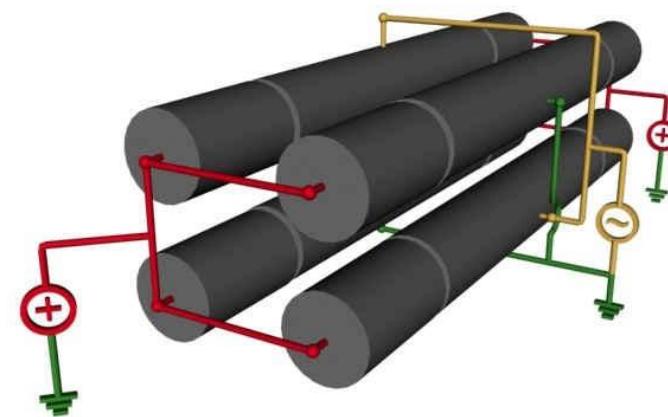
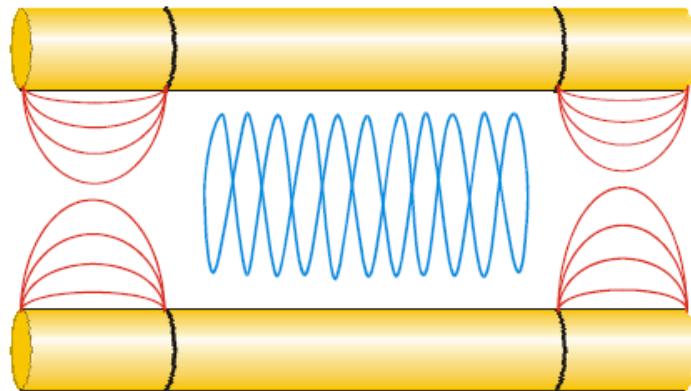
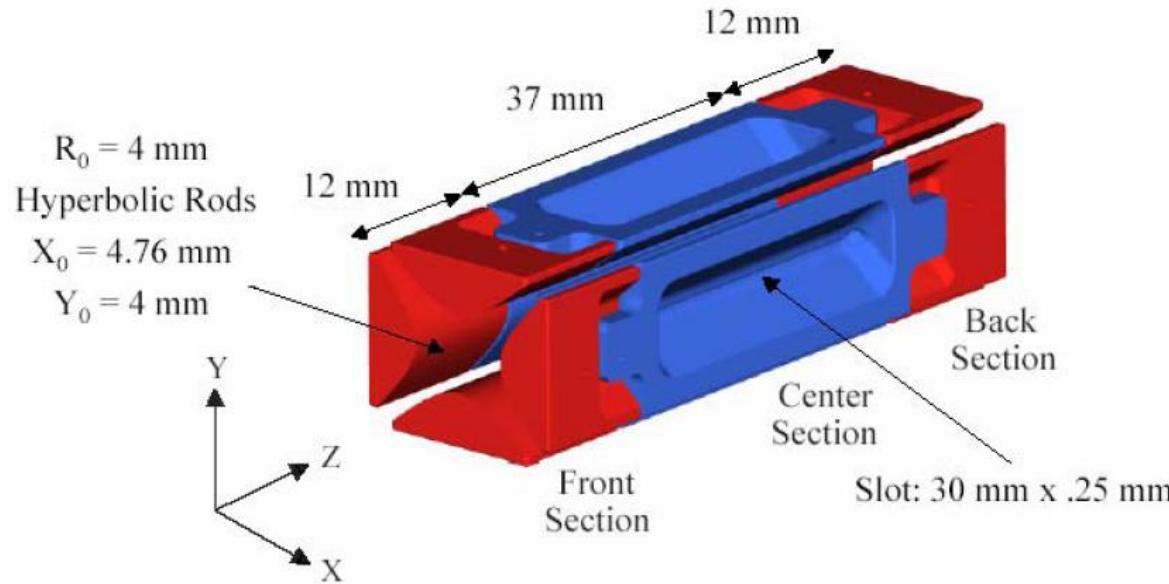
## Quadrupole: ion trap - MS/MS scan



- |   |                               |
|---|-------------------------------|
| <b>1. Trap emptying</b>                         | <b>5. Fragmentation dwell</b> |
| <b>2. Accumulation of ions</b>                  | <b>6. Fragmentation</b>       |
| <b>3. Isolation dwell</b>                       | <b>7. Scan dwell</b>          |
| <b>4. Isolation of parent ions (precursors)</b> | <b>8. Mass analysis</b>       |

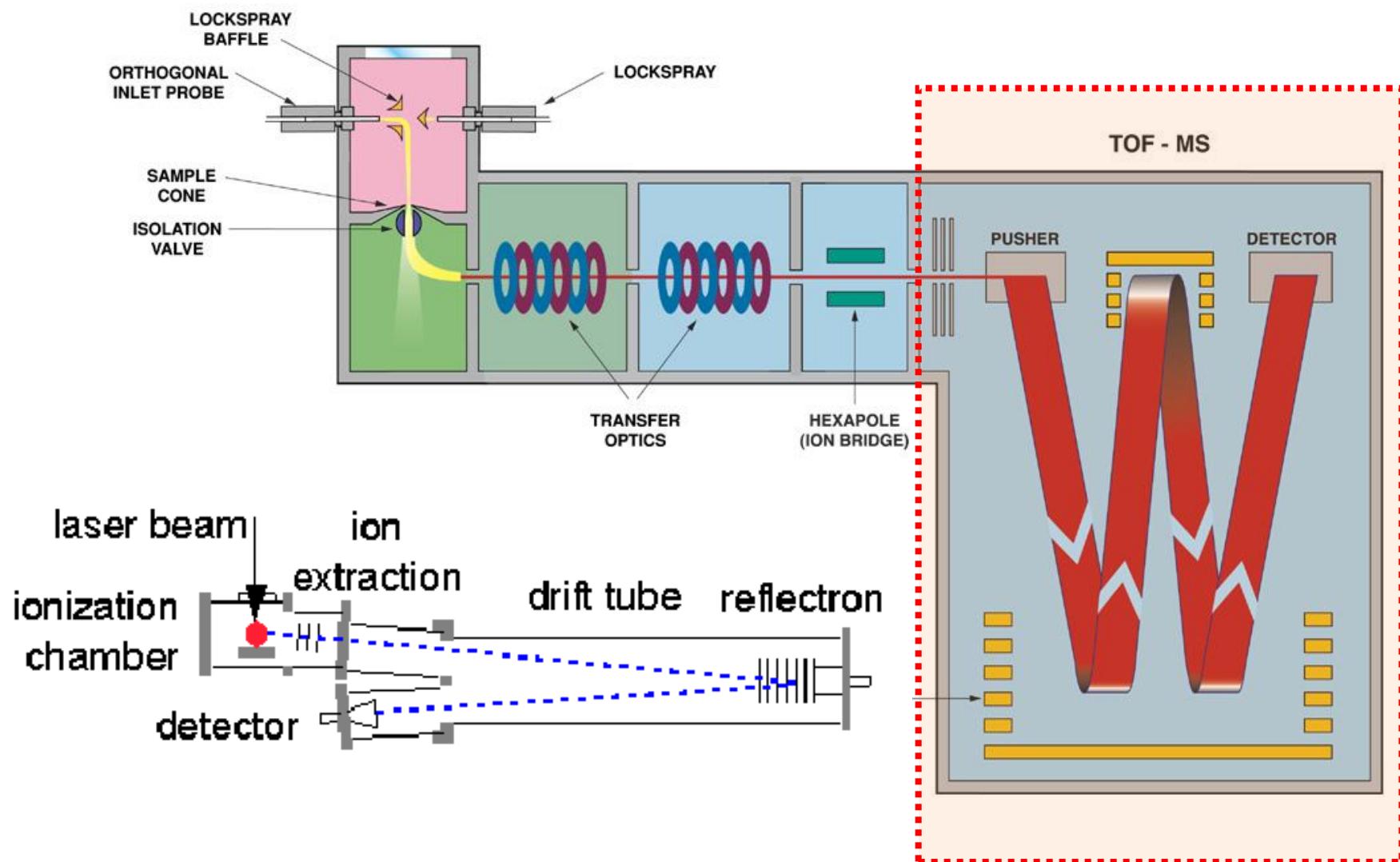
# Technical principles of mass analyzers (7)

## Quadrupole: linear ion trap (segmented quadrupole)



# Technical principles of mass analyzers (8)

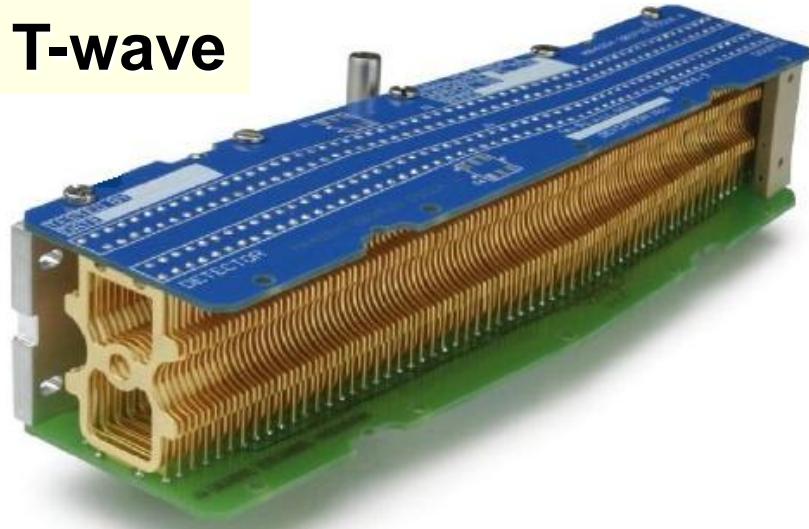
## Time Of Flight (TOF)



# Technical principles of mass analyzers (9)

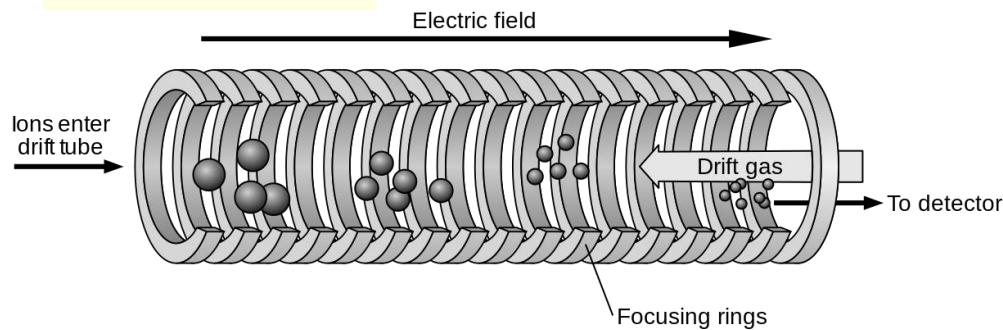
## Travelling wave

T-wave



Drift tube

Ion mobility



Reflectron - TOF



# Technical principles of mass analyzers (8&9)

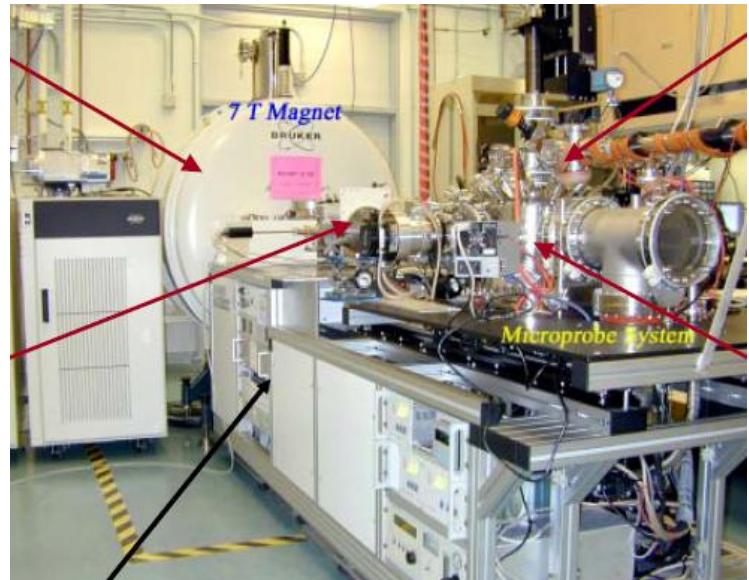
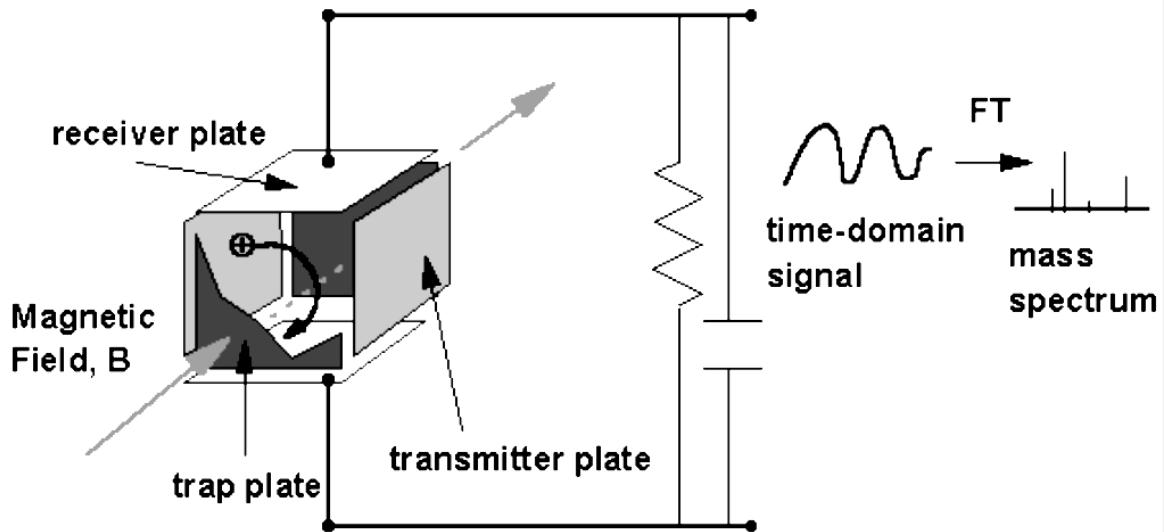
**Ion Mobility (Travelling wave) - TOF MS (with reflectron)**

**VIDEO**



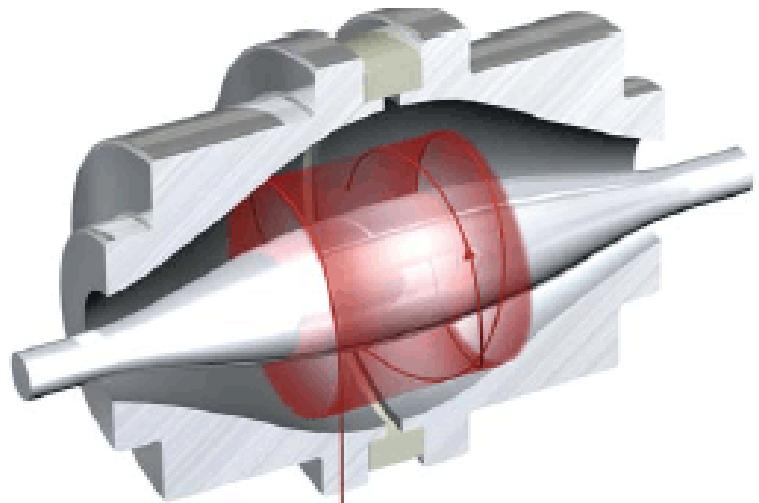
# Technical principles of mass analyzers (10)

## Fourier transform ion cyclotron resonance



# Technical principles of mass analyzers (11)

## ORBITRAP



# Comparison of mass analyzers

Mass analyzer	RP (FWHM)	RANGE <i>m/z</i>	SCAN SPEED	PRICE [mil. CZK]
Magnet-sector	$10^5$	$10^4$	low	10 – 15
Quadrupole – straight	$10^3$	$10^3$	medium	2 – 4
Quadrupole – 3D trap	$10^3$	$10^4$	high	3 – 6
TOF	$10^4$	$10^3 (10^6)$	very high	6 - 8
ICR	$10^6$	$10^8$	high	30
Orbitrap	$10^5$	$10^4$	high	10

# Ion detectors and signal multipliers (1)

## Direct detection:

- a) photographic plate (past)
- b) Faraday cup - accurate number of ions

## Detection using signal multipliers:

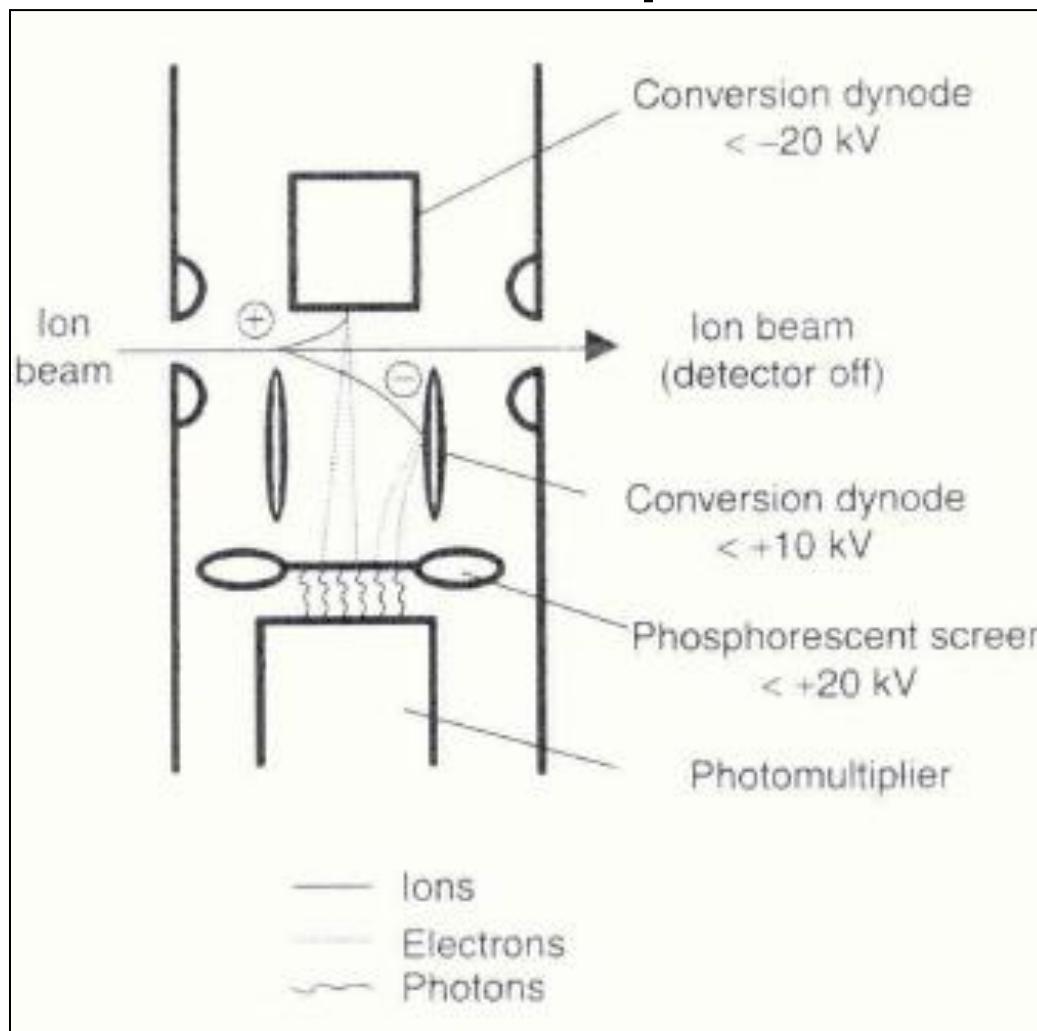
- a) multiplier with conversion - photomultiplier
- b) electron multiplier producing measurable current
  - dynode plates
  - dynode continuous (tube - straight, curved)
  - microchannel plate

## Parameters:

amplification, durability, undesirable detection (neutral fragments)

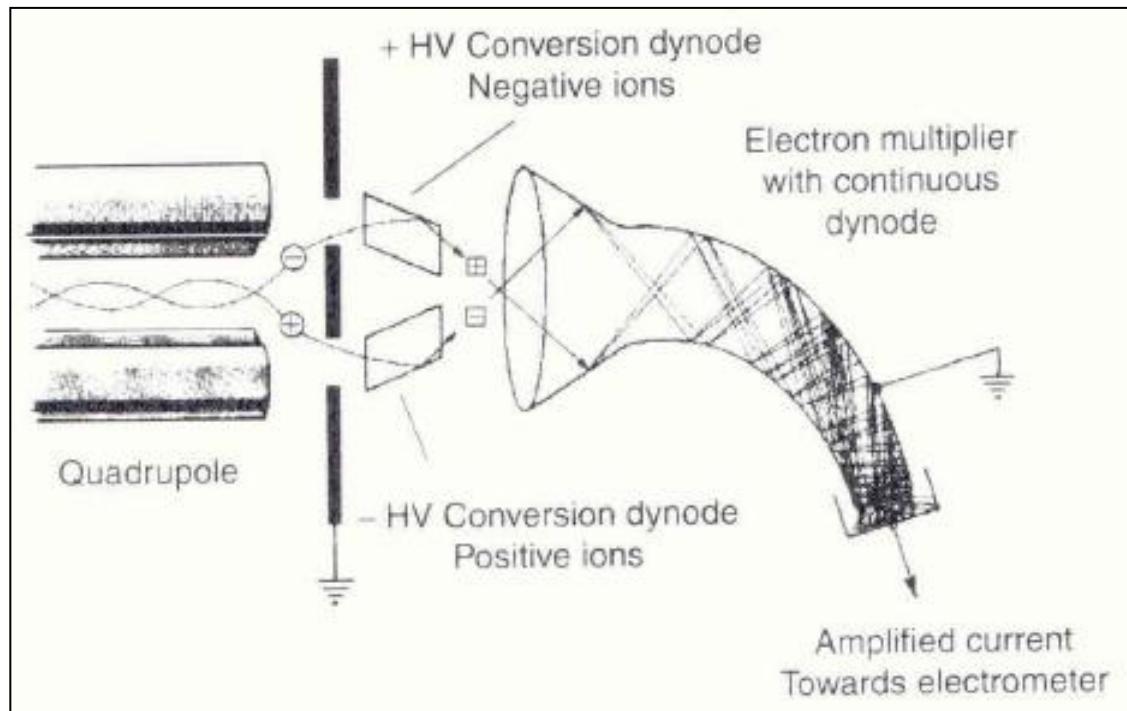
# Ion detectors and signal multipliers (2)

## Photomultiplier



# Ion detectors and signal multipliers (3)

## Electron multiplier



# Ion detectors and signal multipliers (4)

