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CEE 772: Instrumental Methods in Environmental Analysis

Lecture #20

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Mass Spectrometry and Instrumentation

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Content

- **A brief introduction to mass spectrometry**
- **Mass spectrometry instrumentation**
 - Important MS instrument performance factors
 - Types of mass spectrometers:
 - (Triple) Quadrupole Mass Spectrometer
 - Quadrupole Ion Traps (QIT)
 - Time-of-flight (TOF) Mass Analyzers

Ion source ➡ Mass filtration/separation ➡ Detection

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Mass Spectrometry Introduction

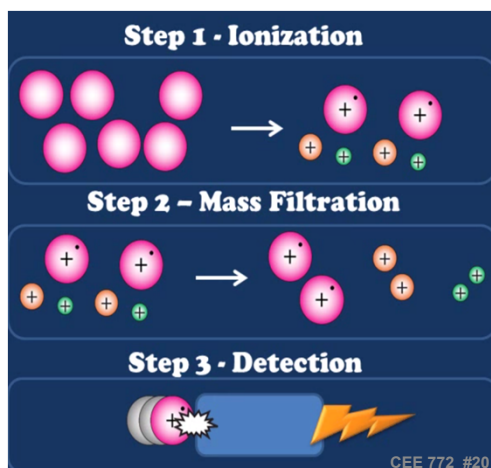
Ion Formation



Mass Analysis



Detection



- **Ionization:**

- Electrospray Ionization
- Electron Impact/Chemical Ionization
- MALDI (matrix assisted laser desorption ionization)

- **Mass Analysis/Separation:**

- Use electric and magnetic fields to apply a force on charged particles to control the trajectories of ions

- **Detection:**

- Electron multiplier

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The effect of electromagnetic fields on ions

- The relationship between force, mass, and the applied fields can be summarized in Newton's second law and the Lorentz force law.

1. **Newton's second law:**

$$\mathbf{F} = m\mathbf{a}$$

the force causes an acceleration that is **MASS** dependent

2. **Lorentz force law:**

$$\mathbf{F} = e(\mathbf{E} + \mathbf{v}\mathbf{B})$$

the applied force is also dependent on the **IONIC CHARGE**

F = the force applied to the ion

m = the mass of ion

a = acceleration

e = ionic charge

E = the electric field

$\mathbf{v}\mathbf{B}$ = the vector cross product of the ion velocity and the applied magnetic field

- Mass spectrometers separate ions according to their mass-to-charge ratio (m/z) rather than by their mass alone**

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Tandem Mass Spectrometry

- Tandem mass spectrometry**, also known as MS/MS or MS² (MSⁿ only by ion traps), involves **multiple steps of mass spectrometry selection**, with some form of fragmentation occurring in between the stages.
- Select and fragment ions of interest to provide **structural information** (e.g. large molecules, such as proteins, polypeptides with a great number of residues, etc.)

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Important MS Instrument Performance Factors

- **Mass Accuracy:** How accurate is the mass measurement?
- **Resolution:** How well separated are the peaks from each other?
- **Sensitivity:** How small an amount can be analyzed?
(More about sample preparation and which type of instrumentation applied.)

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Mass Accuracy

- **How is mass defined?**

Assigning numerical value to the intrinsic property of “mass” is based on using carbon-12, ^{12}C , as a reference point.

One unit of mass is defined as a Dalton (Da); thus **one Dalton is defined as 1/12 the mass of a single ^{12}C atom.**

By definition, one ^{12}C atom has a mass of 12.0000 Da

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Mass Accuracy

- **Isotopes**

Most elements have more than one stable isotope.

For example, most carbon atoms have a mass of 12 Da, but in nature, 1.1% of carbon atoms have an extra neutron, making their mass 13 Da.

Why do we care?

Mass spectrometers “see” the isotope peaks provided the resolution is high enough.

If an MS instrument has resolution high enough to resolve these isotopes, better mass accuracy is achieved.

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Stable Isotopes of Some Common Elements

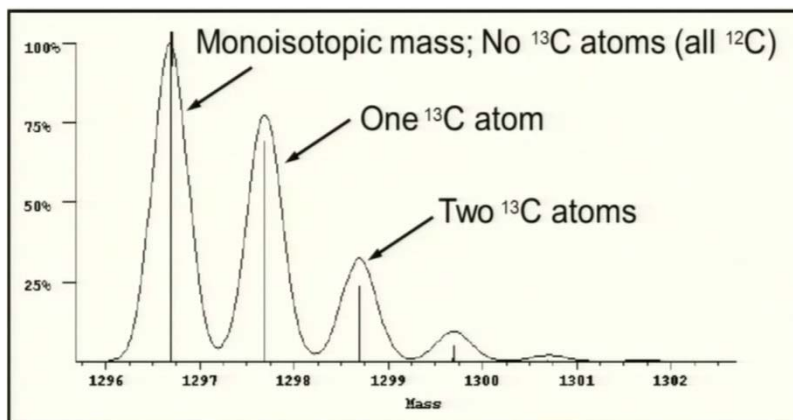
Z	Name	Symbol	Mass of Atom (u)	% Abundance	Z	Name	Symbol	Mass of Atom (u)	% Abundance
1	Hydrogen	¹ H	1.007825	99.9885	15	Phosphorus	³¹ P	30.973762	100
	Deuterium	² H	2.014102	0.0115	16	Sulphur	³² S	31.972071	94.93
	Tritium	³ H	3.016049	*			³³ S	32.971458	0.76
2	Helium	³ He	3.016029	0.00137			³⁴ S	33.967867	4.29
		⁴ He	4.002603	99.99863			³⁶ S	35.967081	0.02
3	Lithium	⁶ Li	6.015122	7.59	17	Chlorine	³⁵ Cl	34.968853	75.78
		⁷ Li	7.016004	92.41			³⁷ Cl	36.965903	24.22
4	Beryllium	⁹ Be	9.012182	100	18	Argon	³⁶ Ar	35.967546	0.3365
5	Boron	¹⁰ B	10.012937	18.9			³⁸ Ar	37.962732	0.0632
		¹¹ B	11.009305	80.1			⁴⁰ Ar	39.962383	99.6003
6	Carbon	¹² C	12.000000	98.93	19	Potassium	³⁹ K	38.963707	93.2581
		¹³ C	13.003355	1.07			⁴⁰ K	39.963999	0.0117
		¹⁴ C	14.003242	*			⁴¹ K	40.961826	6.7302
7	Nitrogen	¹⁴ N	14.003074	99.632	20	Calcium	⁴⁰ Ca	39.962591	96.941
		¹⁵ N	15.000109	0.368			⁴² Ca	41.958618	0.647
8	Oxygen	¹⁶ O	15.994915	99.757			⁴³ Ca	42.958767	0.135
		¹⁷ O	16.999132	0.038			⁴⁴ Ca	43.959481	2.086
		¹⁸ O	17.999160	0.205			⁴⁶ Ca	45.955993	0.004
9	Fluorine	¹⁹ F	18.998403	100			⁴⁸ Ca	47.952534	0.187
10	Neon	²⁰ Ne	19.992440	90.48	21	Scandium	⁴⁵ Sc	44.955910	100
		²¹ Ne	20.993847	0.27	22	Titanium	⁴⁶ Ti	45.952629	8.25
		²² Ne	21.991386	9.25			⁴⁷ Ti	46.951764	7.44
11	Sodium	²³ Na	22.989770	100			⁴⁸ Ti	47.947947	73.72
12	Magnesium	²⁴ Mg	23.985042	78.99			⁴⁹ Ti	48.947871	5.41
		²⁵ Mg	24.985837	10.00			⁵⁰ Ti	49.944792	5.18
		²⁶ Mg	25.982593	11.01	23	Vanadium	⁵¹ V	49.947163	0.250
13	Aluminum	²⁷ Al	26.981538	100			⁵² V	50.943964	99.750
14	Silicon	²⁸ Si	27.976927	92.2297	24	Chromium	⁵² Cr	49.946050	4.345
		²⁹ Si	28.976495	4.6832			⁵³ Cr	51.940512	83.789
		³⁰ Si	29.973770	3.0872			⁵⁴ Cr	52.940654	9.501
							⁵⁶ Cr	53.939885	2.365
					25	Manganese	⁵⁵ Mn	54.938050	100
					26	Iron	⁵⁶ Fe	55.934936	91.754
							⁵⁷ Fe	56.935399	2.125
							⁵⁸ Fe	57.933276	0.281
							⁵⁹ Fe	58.934896	0.008

Element	Mass	Abundance
H	1.0078	99.985%
	2.0141	0.015
C	12.0000	98.89
	13.0034	1.11
N	14.0031	99.64
	15.0001	0.36
O	15.9949	99.76
	16.9991	0.04
	17.9992	0.20

The decimal component is referred to as the mass defect

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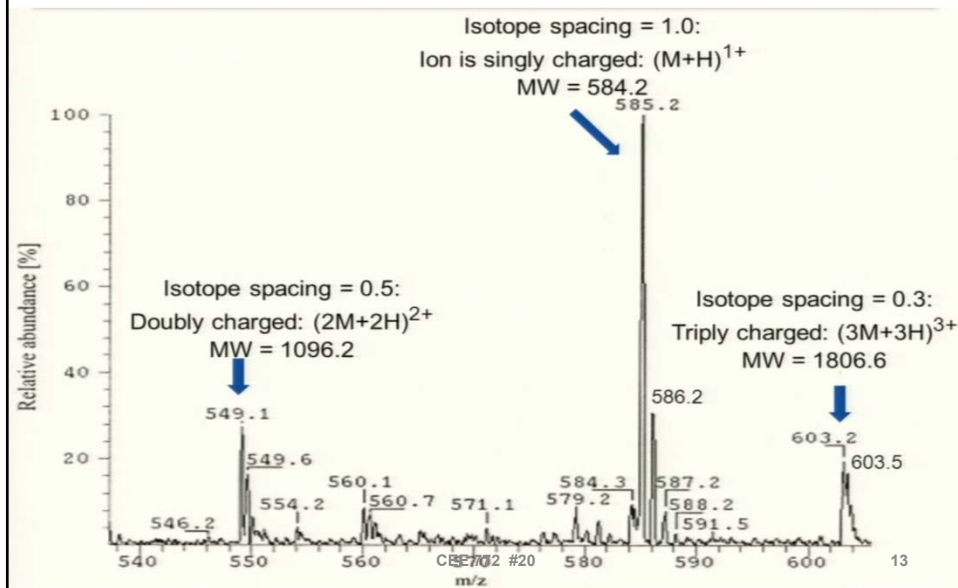
Monoisotopic Mass and Isotopes



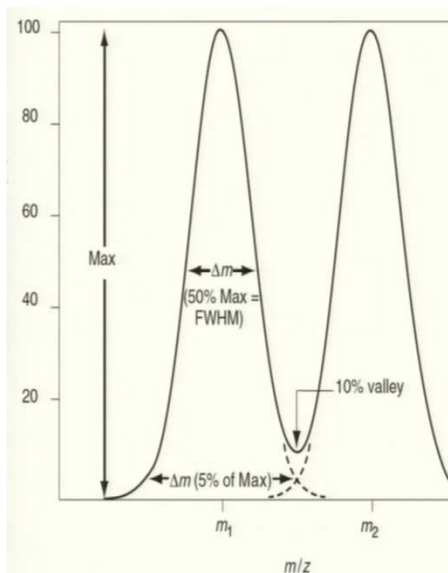
The **monoisotopic mass** of a molecule is the sum of the accurate masses for the most abundant isotope of each element present. As the number of atoms of any given element increases, the percentage of the population of molecules having one or more atoms of a heavier isotope of this element also increases.

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Mass Spectrometers measure mass-to-charge, not molecular mass



Mass Resolution



Schematic representation of the two common definitions of resolution used in mass spectrometry:

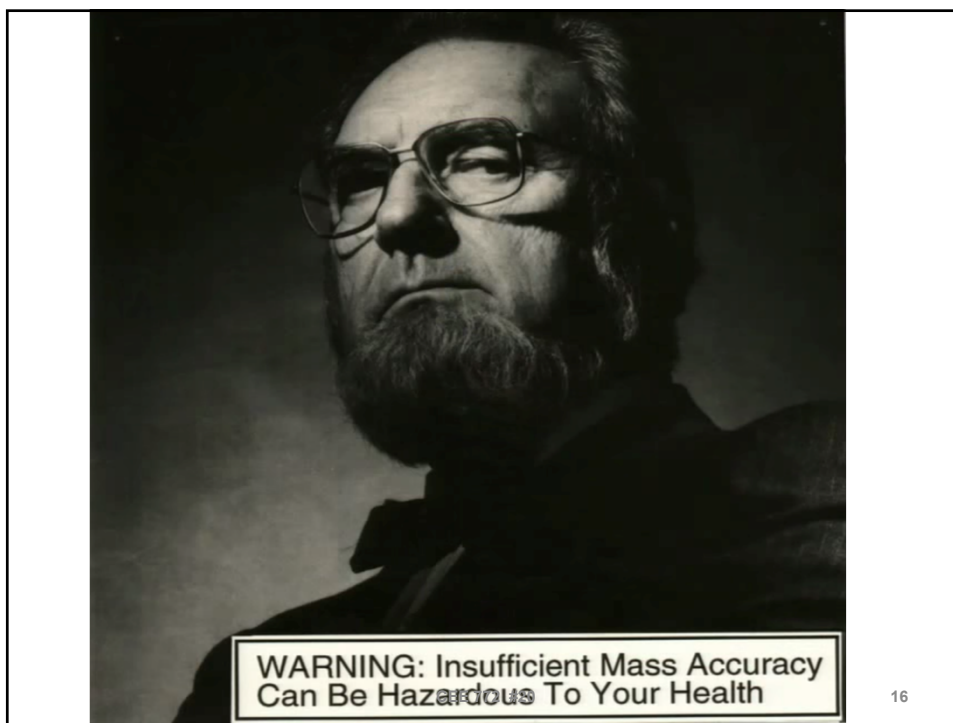
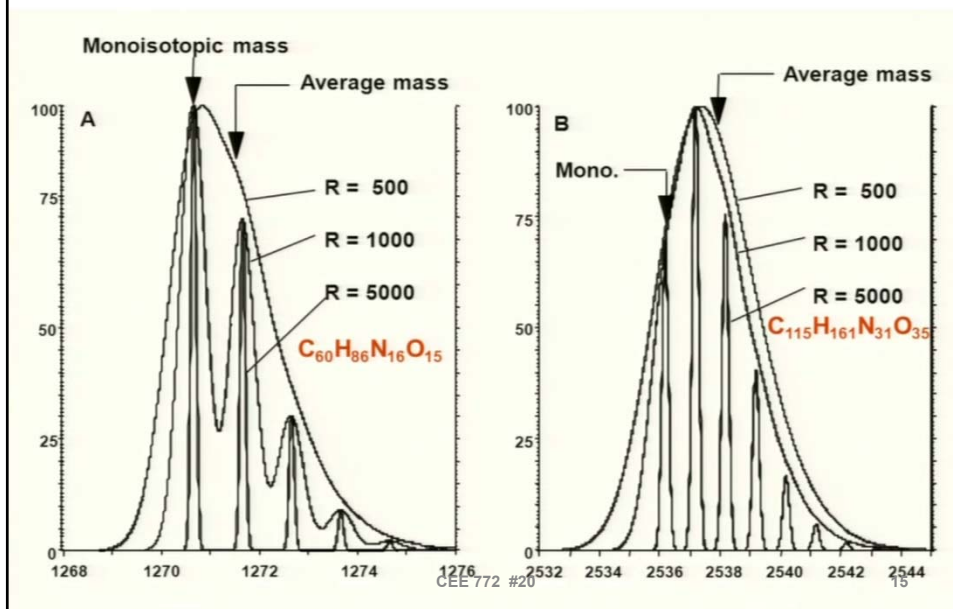
- 10% valley
- Full width, half-maximum (FWHM)

The peak width definition considers a single peak in a mass spectrum made up of singly charged ions at mass m . The resolution R is expressed as $m/\Delta m$, where m is the width of the peak at a half-height (50%).

R is usually a function of m , $m/\Delta m$ should be given for a

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Mass Resolution



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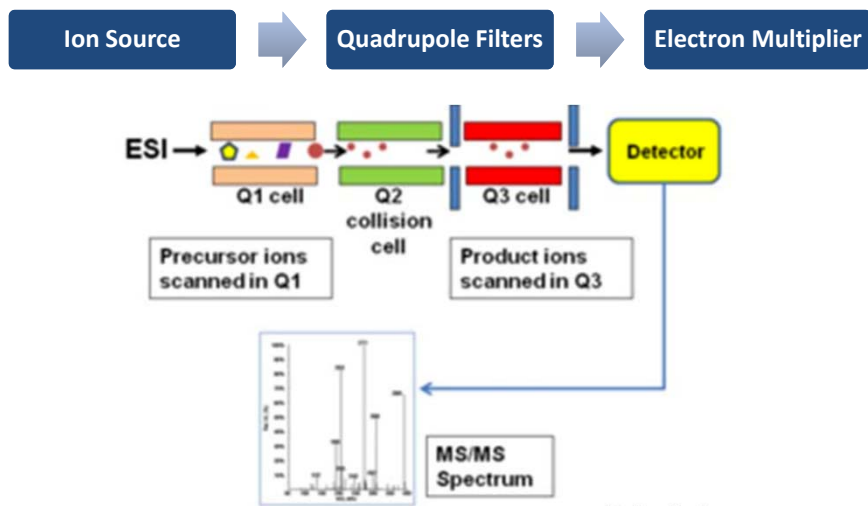


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Types of Spectrometers

1. Triple Quadrupole Mass Spectrometer



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Ion Formation: Electrospray Ionization (ESI)

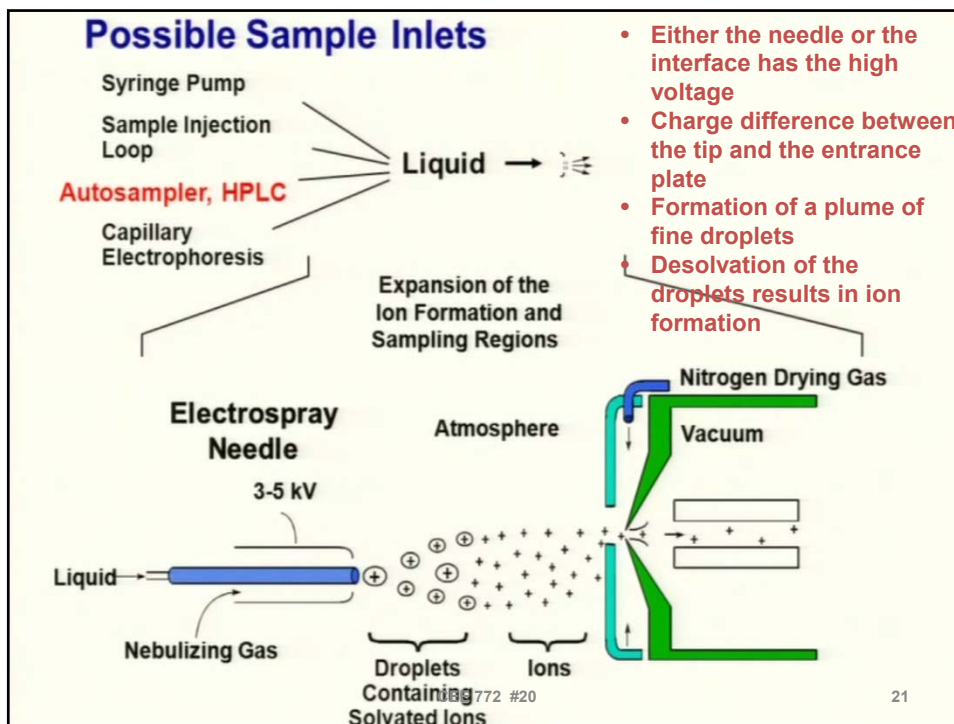
- **Liquid-phase** dissolved analytes eluting from a chromatographic separation system (LC column) must be converted into **GAS-PHASE** under high vacuum.

• Conversion Process

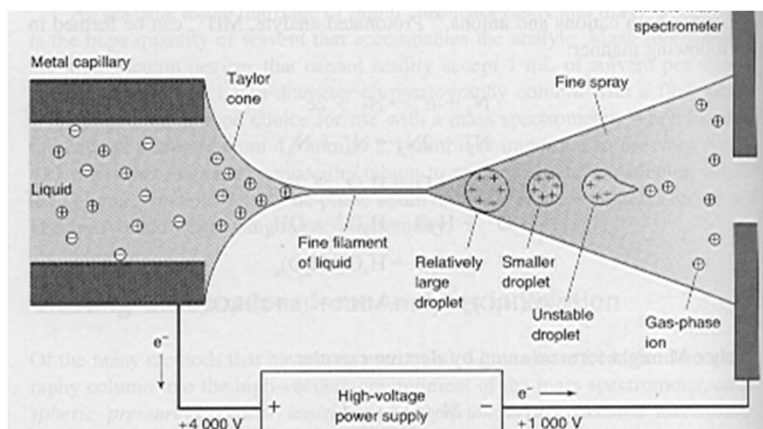
State-of-matter:	Liquid-phase	Evaporation	Gas Phase
Pressure:	Atmospheric	Pressure Reduction	High Vacuum
Charge State:	Neutral(Ionic)	Ionization	Ionic

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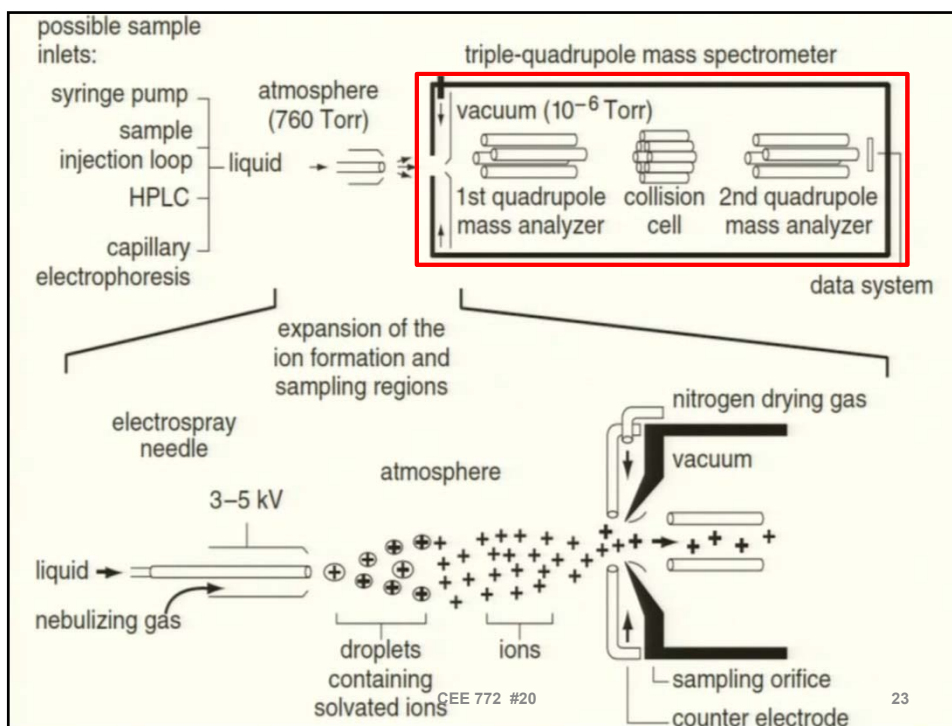


Electrospray Ionization

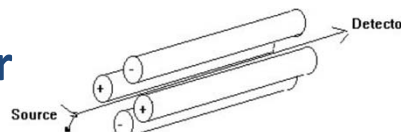


ESI Types:

1. **Positive** – Use volatile proton donor (e.g. 0.1% formic acid)
2. **Negative** – Use volatile proton acceptor (e.g. 0.3%



Quadrupole Mass Filter



- The quadrupole mass analyzer is a **“mass filter”**.
- Quadrupole is made up of four parallel rods. The rods are **electrodes**, with electric fields around them.
- **Combined DC and RF potentials** on the quadrupole rods can be set to pass only a **selected mass-to-charge ratio**. All other ions do not have a stable trajectory through the quadrupole will collide with the quadrupole rods, never reaching the detector.

Quadrupole Mass Filter

Configuration of Voltages

- Each pair of rods is connected; rods have exactly the same voltage as the one directly opposite.

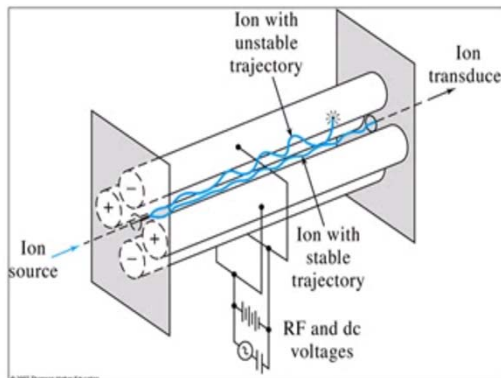
- One pair of rods have voltage:

$$+V_{DC} + V_{RF} \cos(\omega t)$$

The other pair have voltage:

$$-V_{DC} - V_{RF} \cos(\omega t)$$

- Ions either make it through

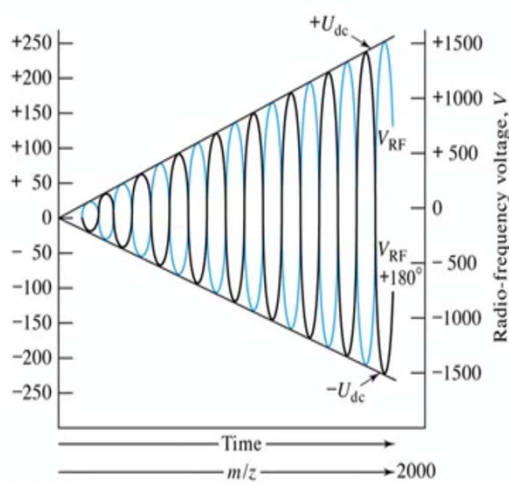


Only ions of a specific mass-to-charge ratio make it through quadrupole based on magnitudes of V_{DC} and V_{RF}

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Voltage Relationships During a Mass Scan

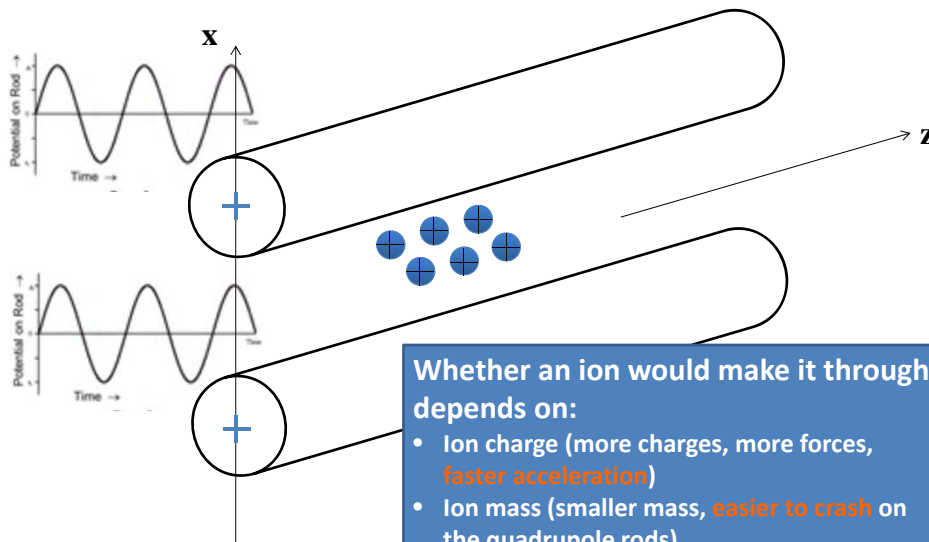


- Amplitude of the DC and alternating fields both increase in time.
- Amplitude of the alternating fields is ~6x strength of the DC fields.
- RF/DC ratio is constant

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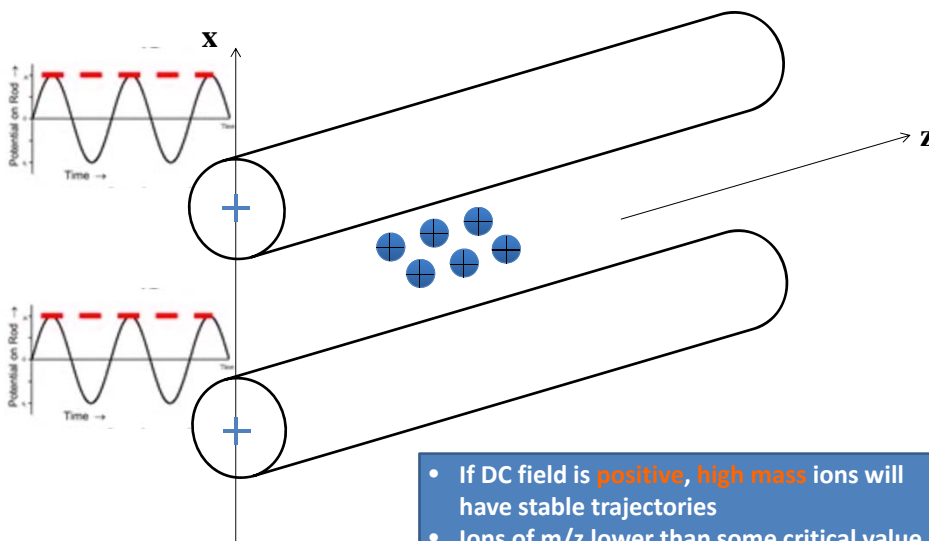
How Do RF Fields Affect Ion Trajectories ?



Whether an ion would make it through depends on:

- Ion charge (more charges, more forces, **faster acceleration**)
- Ion mass (smaller mass, **easier to crash** on the quadrupole rods)
- Strength of field and frequency of

How Do Combined DC & RF Fields Affect Ion Trajectories ?



- If DC field is **positive**, **high mass** ions will have stable trajectories
- Ions of m/z lower than some critical value **will crash**

How Do Combined DC & RF Fields Affect Ion Trajectories ?

- If DC field is **negative**, **low mass ions** will have stable trajectories (**respond quickly**)
- Ions of m/z greater than some critical **value** will crash

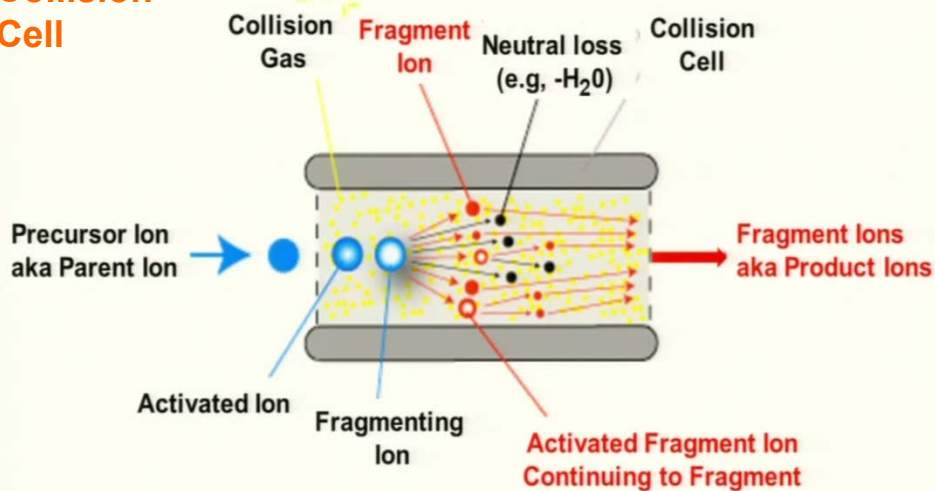
Quadrupole Is A Double Mass Filter

- In **Y dimension (+DC voltage)**, ions make it through unless their mass is too low – **low mass filter**
- In **X dimension (-DC voltage)**, ions make it through unless their mass is too high – **high mass filter**

At certain values of voltage, only a narrow range of m/z will make it through the quadrupole.

Tandem MS – Collision-induced Dissociation (CID)

Collision Cell

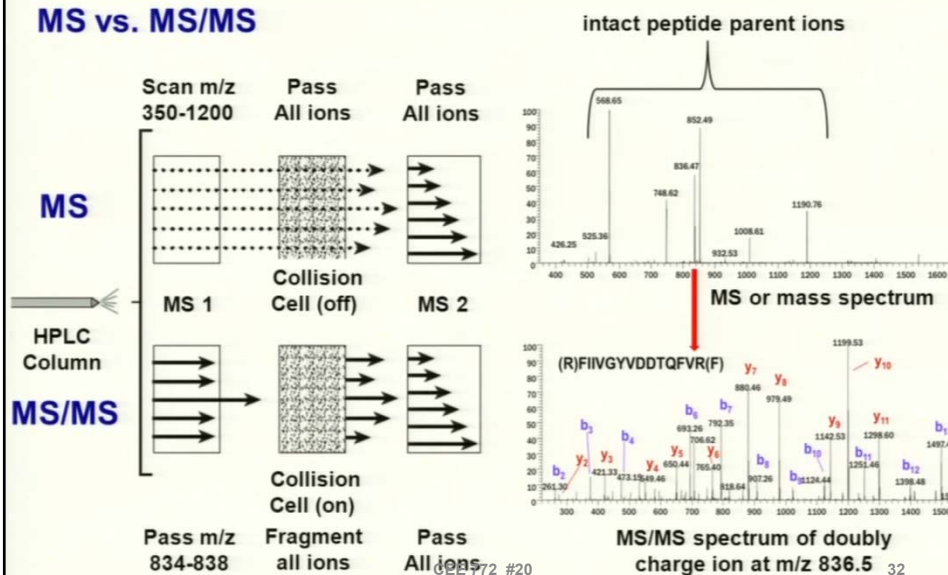


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Structural Information from Tandem Mass Spectrometry

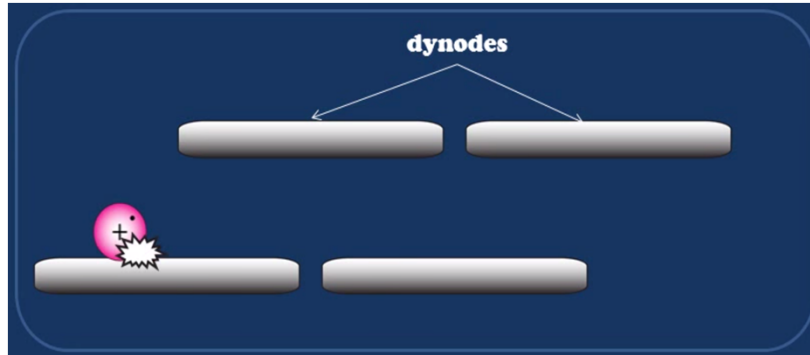
MS vs. MS/MS



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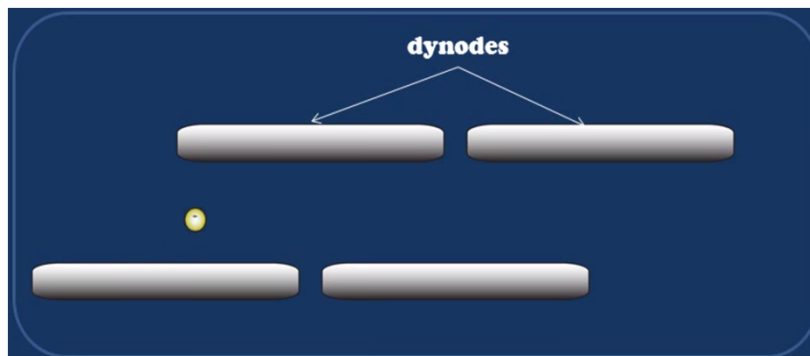
Detector – Electron Multiplier



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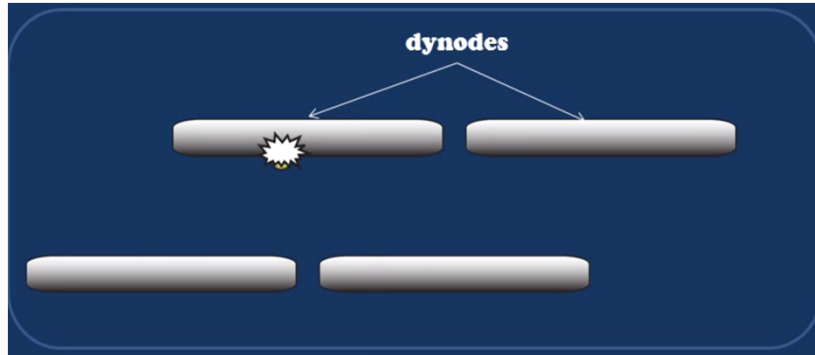
Detector – Electron Multiplier



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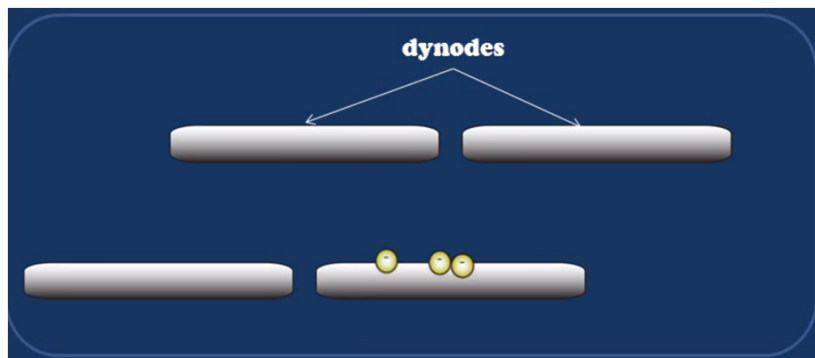
Detector – Electron Multiplier



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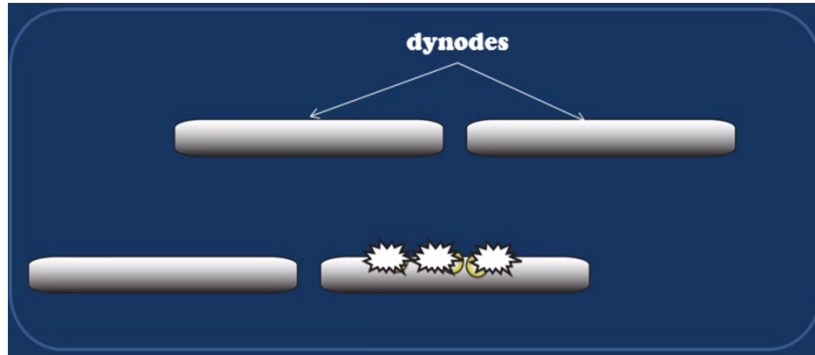
Detector – Electron Multiplier



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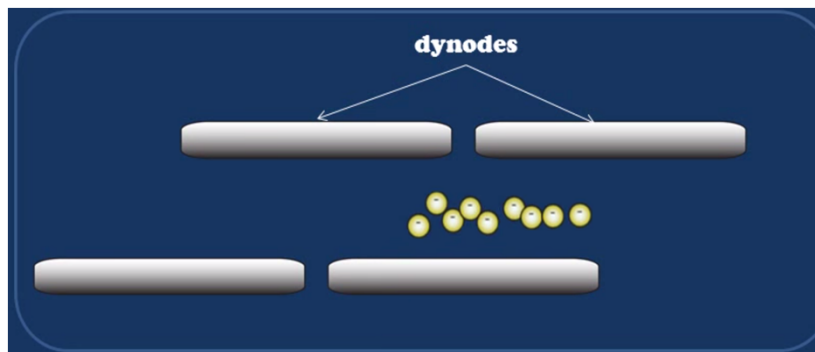
Detector – Electron Multiplier



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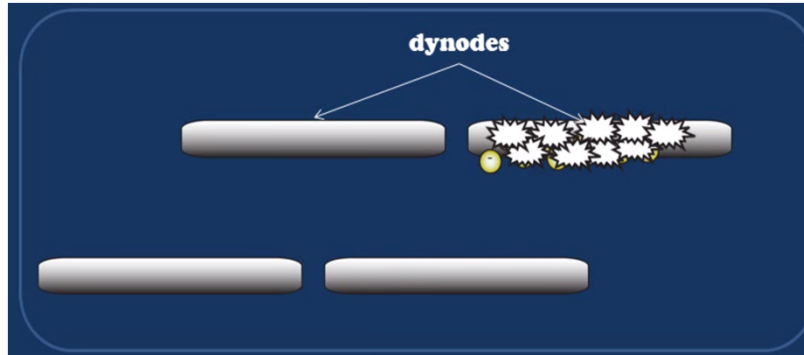
Detector – Electron Multiplier



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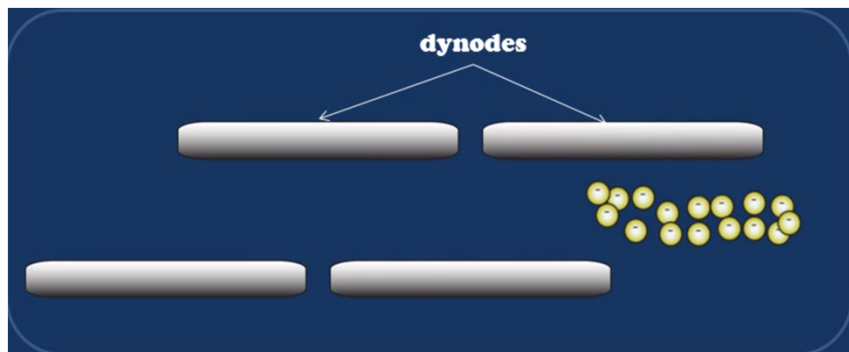
Detector – Electron Multiplier



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Detector – Electron Multiplier



One Impact



$\sim 10^6$ electrons

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Benefits and Limitations of Quadrupole Mass Filter

- **Benefits**

Classical mass spectrometer
 Good reproducibility
 Relatively small and low-cost system
 Low-energy collision-induced dissociation (CID) has efficient conversion of parent to daughter ions

- **Limitations**

Limited resolution
 Not well suited for pulsed ionization methods
 CID depends strongly on energy, collision gas, pressure and other factors

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Types of Spectrometers

2. Quadrupole Ion Traps



***Ions are created inside the trapping volume**

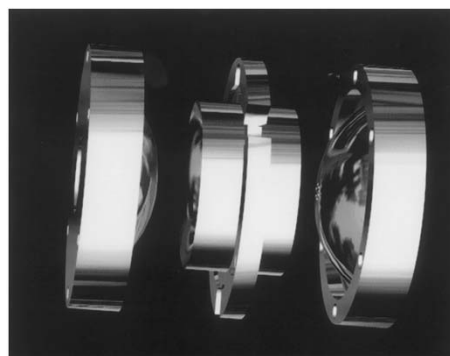
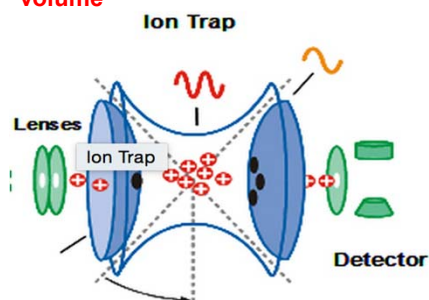
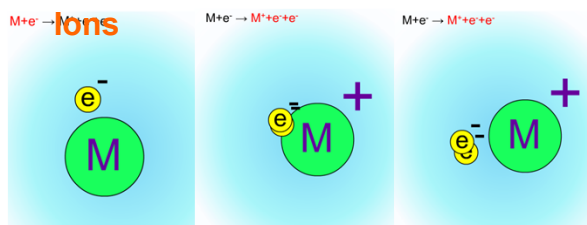


FIG. 1. Rendering of the ion trap electrode assembly showing the ring electrode and the two endcap electrodes.

Jonscher, Karen R., and John R. Yates III. "The quadrupole ion trap mass spectrometer—a small solution to a big challenge." *Analytical biochemistry* 244.1 (1997): 1-15.

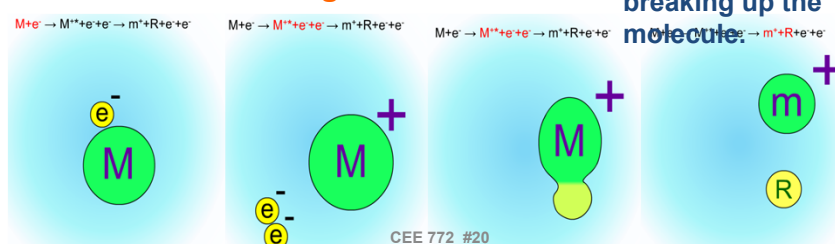
Ion Formation: Electron Impact

1. Production of Molecular Ions



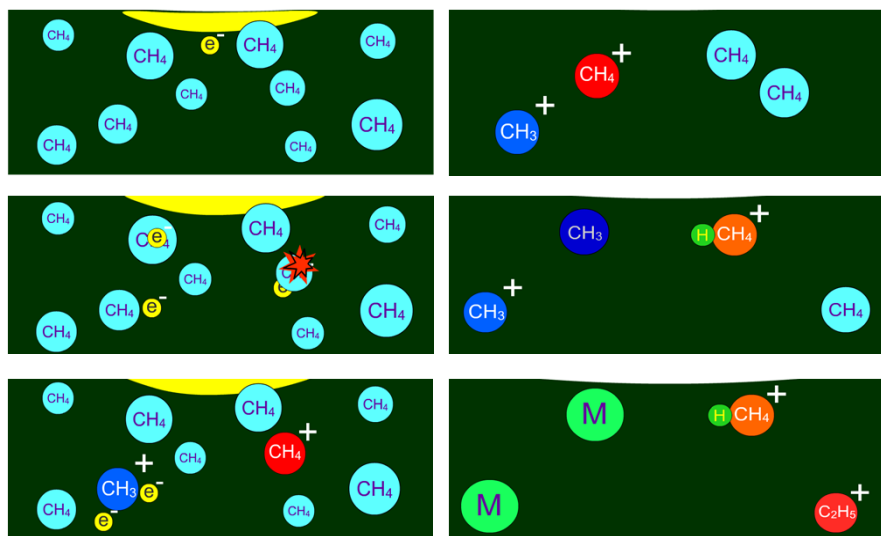
In electron impact ionization (EI), electrons from filament ionizes a compound, in a certain case, by only knocking out one electron without breaking up the molecule.

2. Production of Fragment Ions



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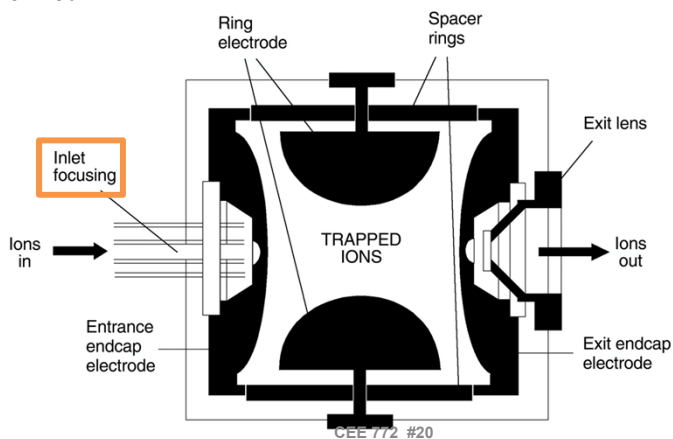
Ion Formation: Chemical Ionization



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Quadrupole Ion Trap (QIT) Configuration

All ions created over a given time period were **trapped** and then **sequentially ejected** from the ion trap into a conventional electron multiplier detector. Thus, all ions were stored while mass analysis was performed.

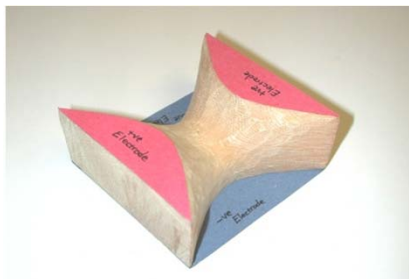


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Ion Trap Theory

- How are ions trapped?

Quadrupole ion traps are dynamic mass analyzers that use an **oscillating electric potential** applied to the ring electrode, called the “fundamental RF”, to **focus ions toward the center of the trap**. This is accomplished by creating a **parabolic potential**, shaped like a “saddle”, inside the trapping volume. The strength of the restoring force linearly increases as the ion trajectory deviates from the central axis, focusing the ion back to the center of the trapping volume.



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Ion Trap Theory

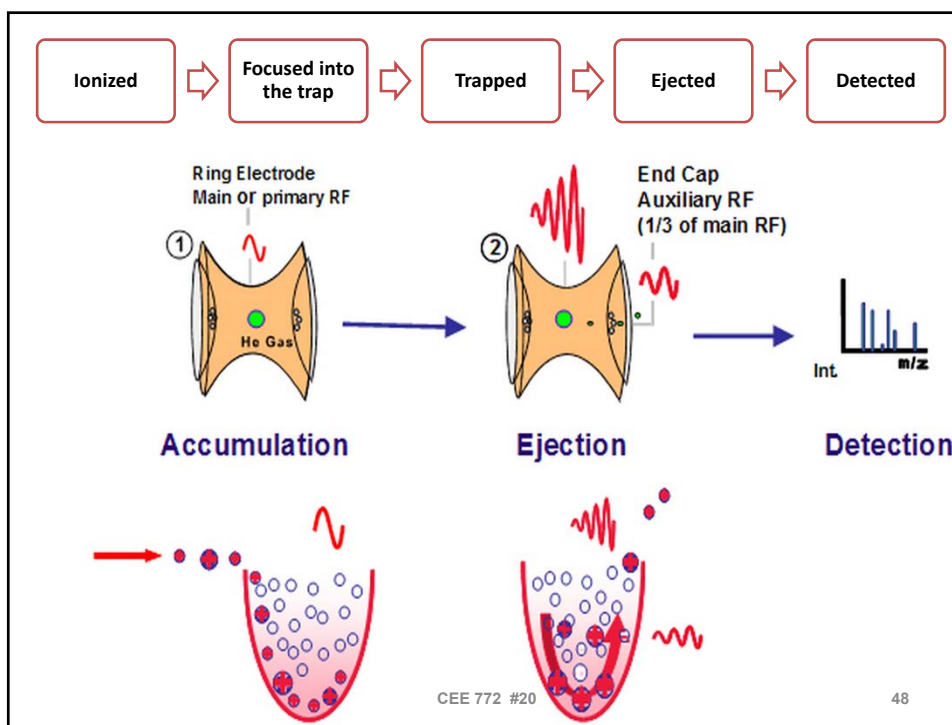
- How are ions **SELECTIVELY AND SEQUENTIALLY** ejected?

Ions are ejected through holes in the **endcap electrode** and are typically detected using an electron multiplier.

Trapped ions of a given m/z oscillate at a frequency known as **the secular frequency**. Resonance conditions are induced by matching **the frequency of a supplementary potential** applied to the endcap electrodes to the secular frequency of the ion. The ion will absorb energy from the applied field and the trajectory will linearly increase toward the endcap electrodes until the ion becomes unstable and is ejected.

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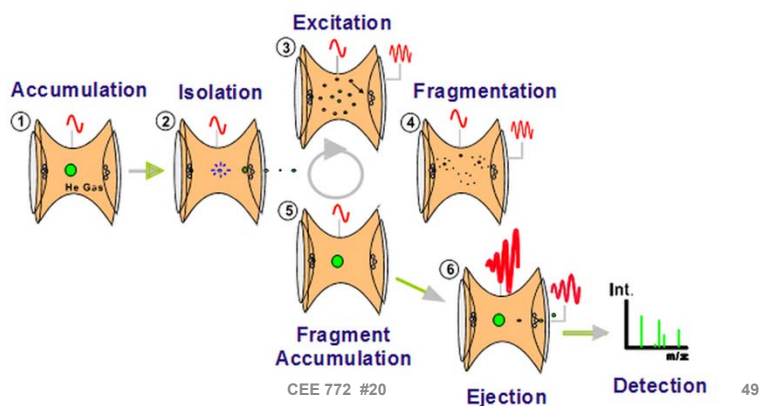
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Tandem MS in QIT

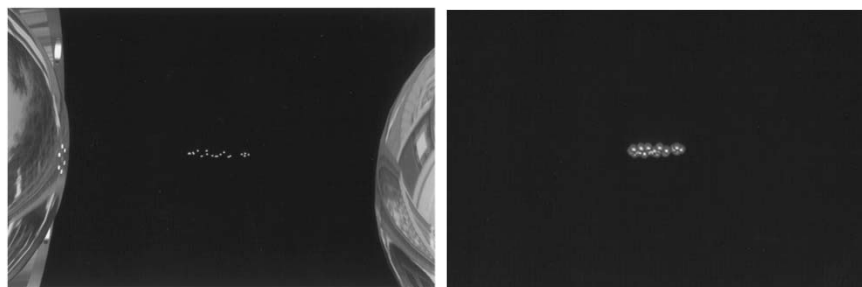
- Tandem mass spectrometry (up to 12 stages, MS¹²)

Structural information is obtained by collision-induced dissociation (CID) with a helium damping gas a mass spectrum is generated by sequentially ejected fragment ions from low m/z to high m/z .



The Damping Gas

The ion trap is typically filled with **helium** to a pressure of ~ 1 mtorr. Collisions with helium **reduce the kinetic energy** of the ions and serve to quickly contract trajectories toward the center of the ion trap, enabling trapping of injected ions. This is called “cooling effect”, where the ion population forms a “packet” near the center of the trap.



Jonscher, Karen R., and John R. Yates III. "The quadrupole ion trap mass spectrometer—a small solution to a big challenge." *Analytical biochemistry* 244.1 (1997): 1-15.

Benefits and Limitations of QIT

- **Benefits**

Compact mass analyzer (roughly the size of a tennis ball)
 Up to 12 stages of tandem mass spectrometry have been performed (multi-stage is unique with QIT)
 Exquisitely sensitive.

- **Limitations**

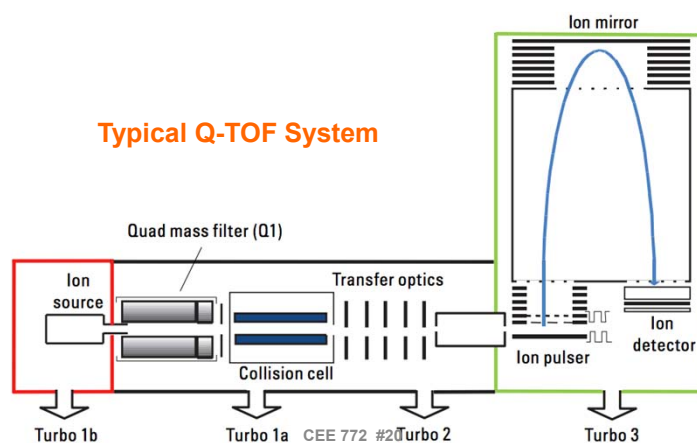
Poor dynamic range
 Subject to "space-charge" effect and ion reaction
 Collision energy not well-defined in CID MS/MS

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Types of Spectrometers

1. Time-of-flight (TOF)

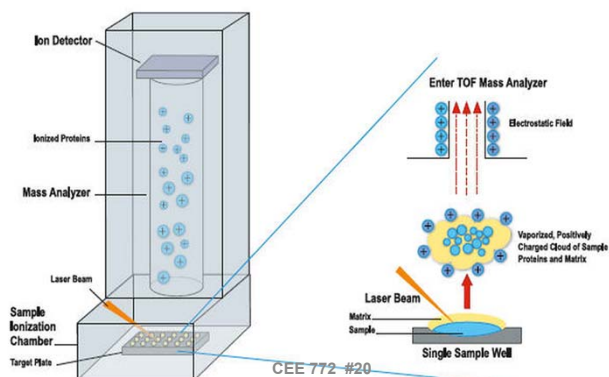


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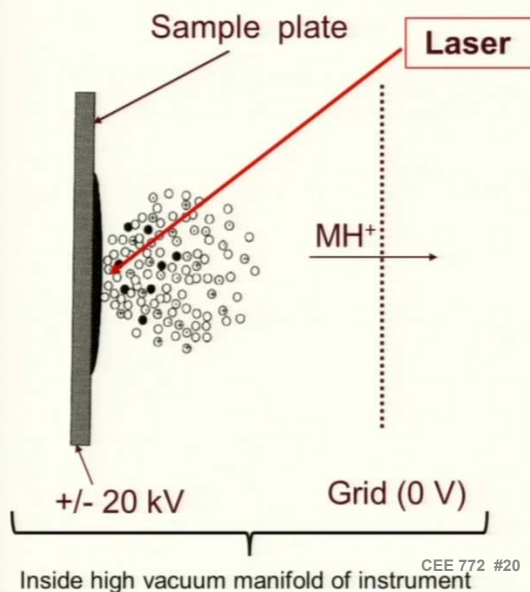
Ion Source: MALDI

- A time of flight mass spectrometer measures the mass-dependent time it takes ions of different masses to move from the ion source to the detector. This requires that the **starting time** (the time at which the ions leave the ion source) is **well-defined**. Therefore, ions are preferably formed by a pulsed ionization method, usually matrix-assisted laser desorption ionization, or **MALDI**.



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MALDI: Matrix Assisted Laser Desorption Ionization



1. Sample is mixed with **matrix** (X) and dried on plate.
2. Laser flash ionizes matrix molecules.
3. Sample molecules (M) are ionized by proton transfer:
 $XH^+ + M \rightarrow MH^+ + X$.

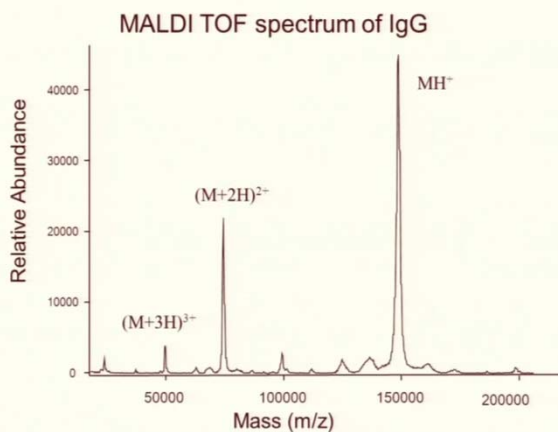
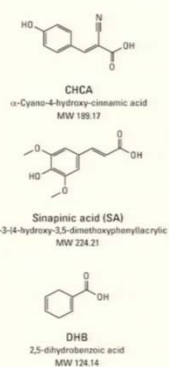


Bruker MALDI-ToF

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MALDI: Matrix Assisted Laser Desorption Ionization

Liquid matrices used for MALDI analysis of biomolecules



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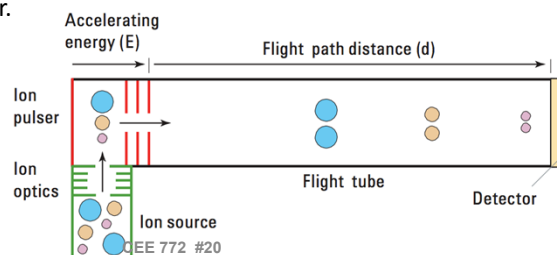
Time-of-flight (TOF) Mass Analyzer

• How does TOF separate ions?

The flight time for each mass is unique. The flight time (t) is determined by the energy (E) to which an ion is accelerated, the distance (d) it has to travel, and its mass (strictly speaking, its mass-to-charge ratio).

$$E = \frac{1}{2}mv^2 \rightarrow v = \sqrt{2E/m}$$

This equation says that for a given kinetic energy, E , smaller masses will have larger velocities, and larger masses will have smaller velocities. Therefore, ions with lower masses arrive at the detector earlier, and higher masses later.



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Time-of-flight (TOF) Mass Analyzer

- **How does TOF separate ions?**

Because: $v = d/t$, there is:

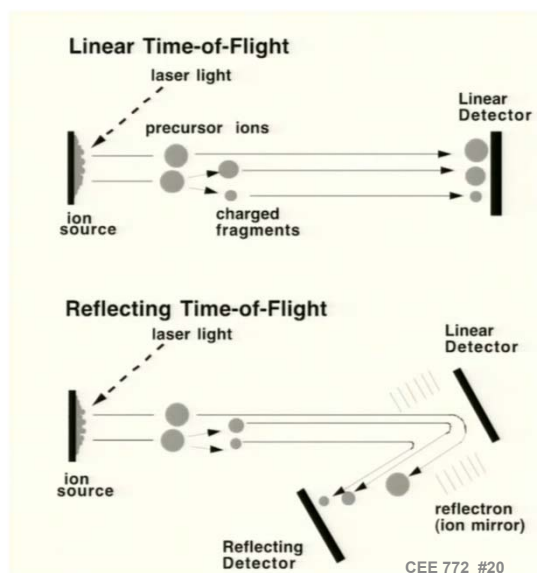
$$m = (2E/d^2)t^2$$

For a given energy (E) and distance (d), the mass is proportional to the square of the flight time of the ion. In design of an TOF mass spectrometer, much effort is devoted to holding the values of **energy (E)** applied to the ions and the **distance (d)** the ion travels constant, so that an **accurate measurement of flight time** will give an accurate mass value.

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Linear vs. Reflecting TOF



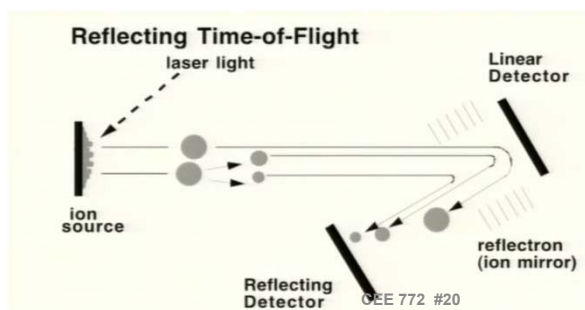
At **higher masses**, resolution is difficult because flight time is much longer. Also at high masses, not all of the ions of the **same m/z** values reach their **ideal TOF velocities**. To fix this problem, often a reflectron is added to the analyzer. The reflectron consists of a series of **ring electrodes of very high voltage** placed at the end of the flight tube. When an ion travels into the reflectron, it is reflected in the opposite

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Reflectron

- A linear-field reflectron allows ions with **greater kinetic energies** to **penetrate deeper** into the reflectron than ions with smaller kinetic energies. The ions that penetrate deeper will take longer time to return to the detector. If a packet of ions of a **given m/z** contains ions with **varying kinetic energies**, then the reflectron will decrease the spread in the ion flight times, and therefore improves the resolution.



Benefits and Limitations

- **Benefits**

- Faster MS analyzer
- Well suited for pulse ionization methods (MALDI)
- High ion transmission
- Highest practical mass range of all MS analyzers

- **Limitations**

- Requires pulsed ionization or ion beam switching
- Limited parent-ion selectivity for tandem mass spectrometry

- [To next lecture](#)