

# QEL, Spices Board

*In-house Training – 1:  
Analysis of Pesticide Residues*

*Lecture - 1*

*Introduction to instrumentation:  
LC-MS/MS*

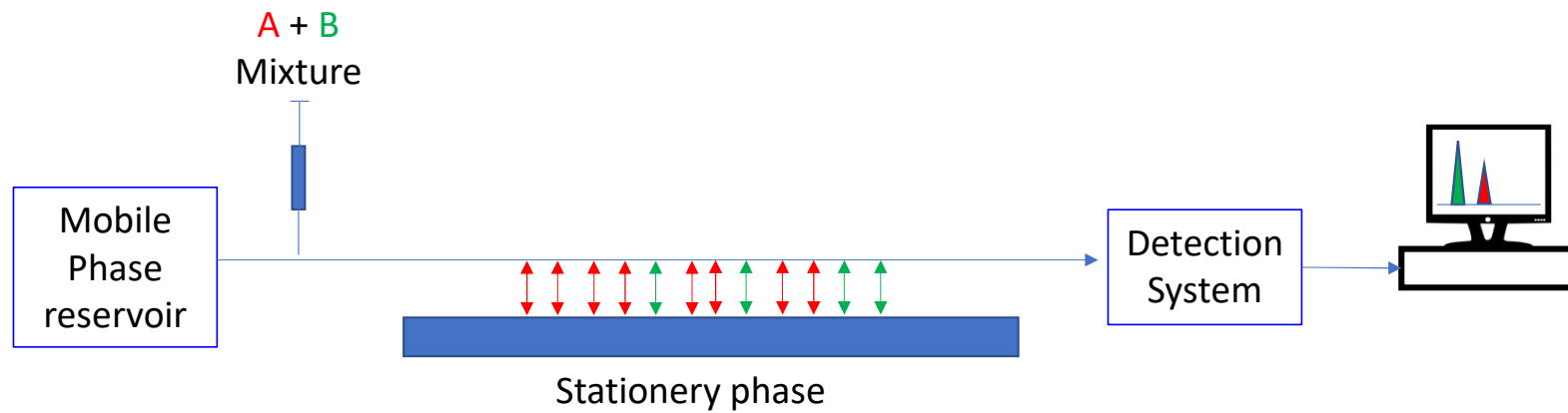
*03-Jul-23*

Dr. Ramesh BN  
Spices Board



# Chromatography

HPLC, GC...



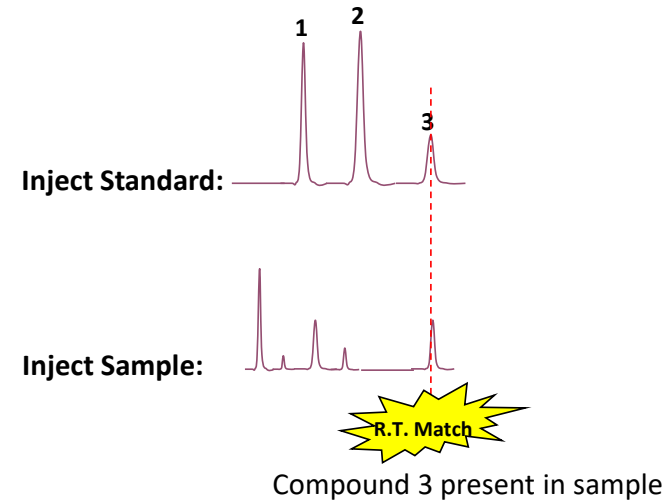
# HPLC Analysis

## Limitations

When **conventional detectors** are used, identification of unknowns by **retention time matching**

Possibility of **errors**

To avoid these errors, we use **mass spectrometry** along with chromatography



### Problems:

- ↳ Two compounds with same Retention Time: **FALSE POSITIVE**
- ↳ Retention Time shift for one compound: **FALSE NEGATIVE**

# Mass Spectrometry

Why?

A mass spectrometer (MS) looks at the **masses** of the analytes being tested

- ↳ Uses **information** derived from these masses for identification
- ↳ High degree of specificity
- ↳ Retention time can be used for additional confirmation

# MS as a detector

## Hyphenated techniques

An MS can function as a detector to a chromatographic system, replacing conventional detectors

- ↳ Gas chromatography (GC-MS), liquid chromatography (LC-MS)
- ↳ Can provide very **specific** and **sensitive** analysis
- ↳ Currently, MS is the default detector in trace analysis work
- ↳ ‘**Tandem**’ MS is possible, which gives even more specificity and sensitivity (MS/MS)

# Mass Spectrometry

## § General principles

# Mass Spectrometry

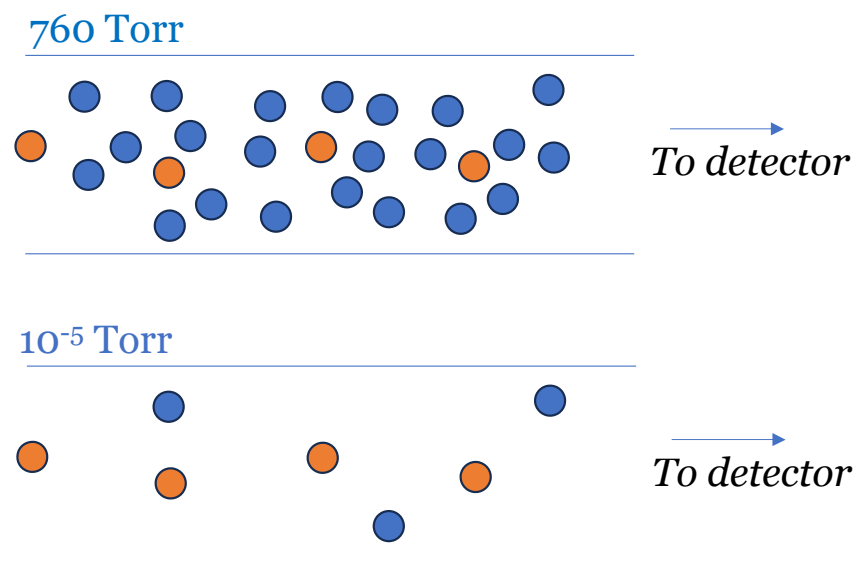
## Principles - 1

A mass spectrometer can 'see' only **charged entities** (ions)

Typically functions under high vacuum

- ↳ The ions need a **stable path through the mass spectrometer**, without colliding with air molecules or with one another

The '**mean free path**': average distance between collisions for a gaseous ion – **must be high**



# Mass Spectrometry

## Principles - 2

In MS the vacuum is achieved in two stages:

- ↳ The **roughing pump** (outside the instrument): from **760 Torr** to  **$10^{-3}$  Torr**
- ↳ The **turbo molecular pump** (inside the instrument): from  **$10^{-3}$  Torr** to  **$10^{-5}$  Torr**





# Mass Spectrometry

## Principles - 3

The connection between LC and MS is called the 'Interface':

- ↳ From HPLC the mobile phase flows out
- ↳ MS cannot handle this – it is a vacuum technique
- ↳ The liquid mobile phase has to be **evaporated** and taken into the MS
- ↳ Loss of analytes should be kept to a minimum
- ↳ In LC-MS, **ionization** also happens in the interface

# Mass Spectrometry

## Principles - 4

MS can operate in two ways.

First way:

- ↳ Molecule of interest is ionized
- ↳ Typically called 'parent' ion.
- ↳ Number of parent ions can be counted for quantification (LC-MS).

Second way:

- ↳ Molecule of interest is ionized, to get the 'parent' ion
- ↳ Parent ion is broken down into multiple 'daughter' ions
- ↳ Selected 'daughter' ions are counted for quantification (LC-MS/MS)

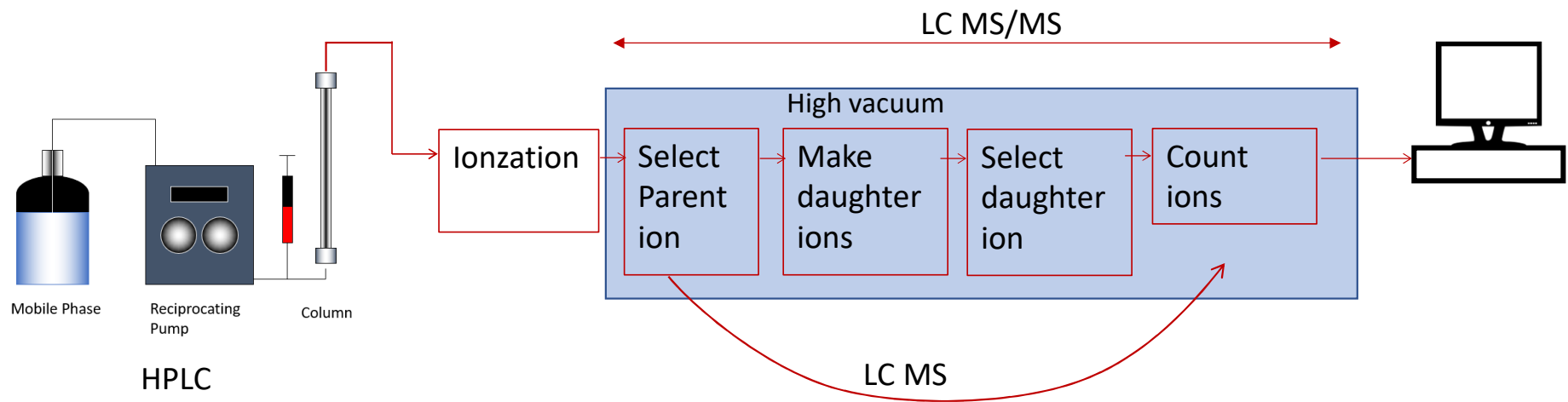
Why is this useful?

# Mass Spectrometry

## § Components and operation

# Mass Spectrometry

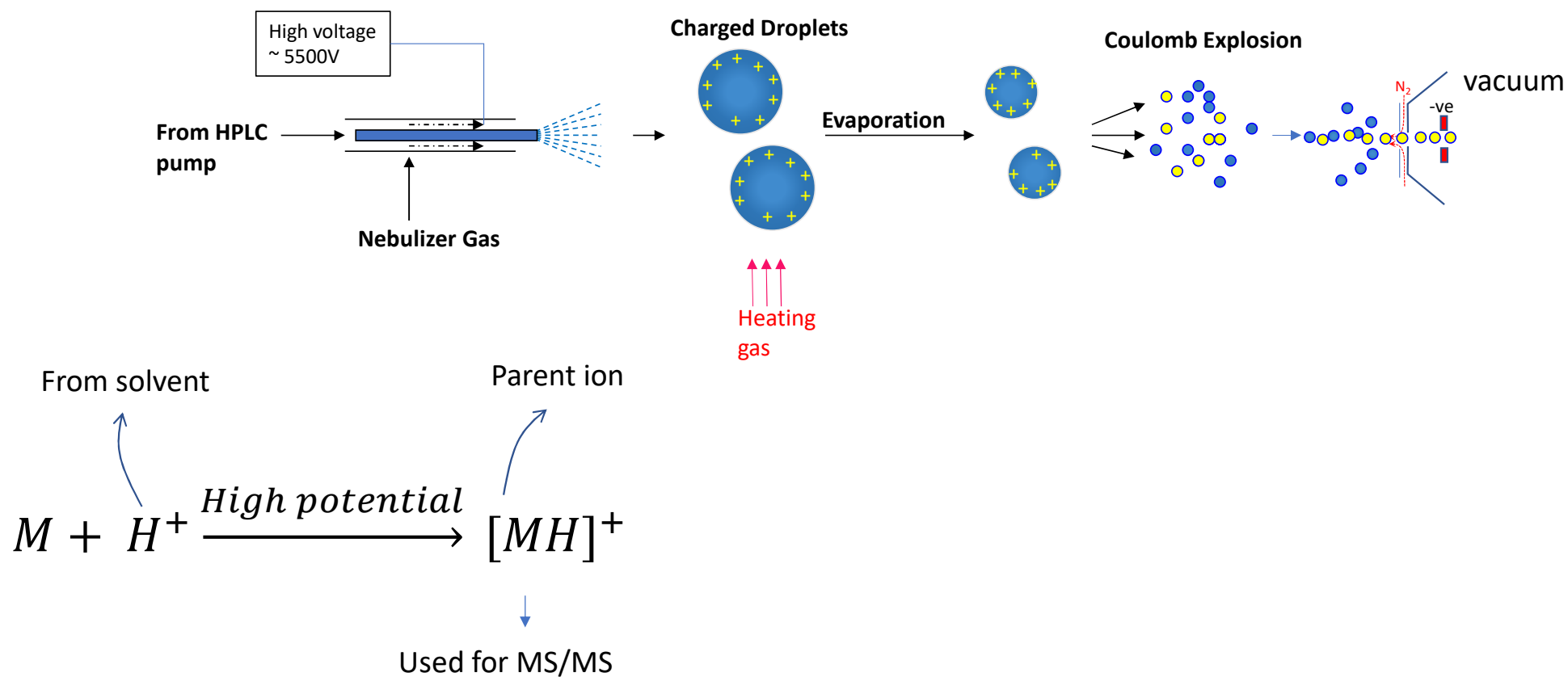
## Components - overview



In LC MS/MS, ionization happens in atmospheric pressure  
Ionization is soft:  $MH^+$  ion is formed  
Standard spectrum libraries not possible

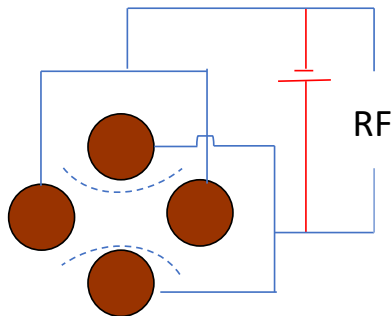
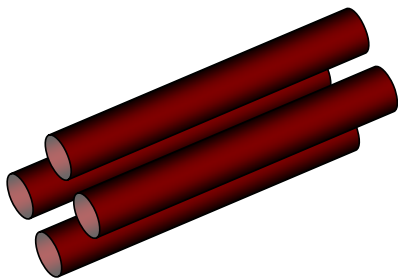
# Mass Spectrometry

## Ionization



# Mass Spectrometry

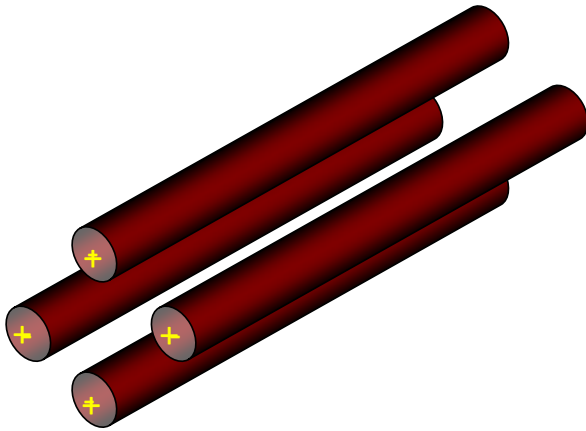
## Quadrupoles (1)



- Four electrodes arranged in a **hyperboloid** fashion
- Two Voltages:
  - **Radio Frequency - RF** (alternating)
  - **Direct Current - DC**
- Applied simultaneously, the two voltages will **trap ions within the quadrupole area**
- **Scanning the voltages** will filter the ions out by mass:
  - For one **RF:DC value**, **only one mass can have a stable trajectory** through the quadrupole
  - When RF:DC is progressively increased (scanned) **Ions pass through in the order of masses**

# Mass Spectrometry

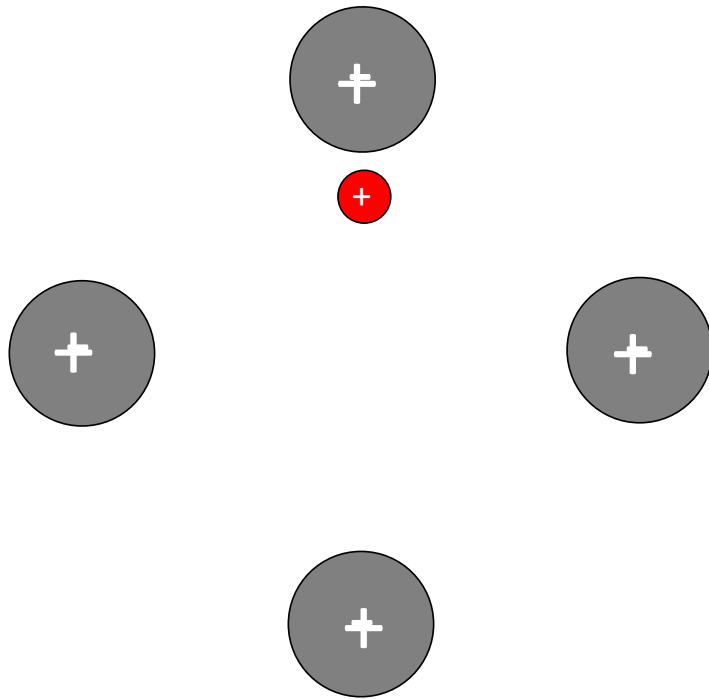
## Quadrupoles (2)



- **Opposite** electrodes of the Quadrupole have the **same** potential
- The potential **alternates** with the same frequency as the RF Voltage

# Mass Spectrometry

## Quadrupoles (3)

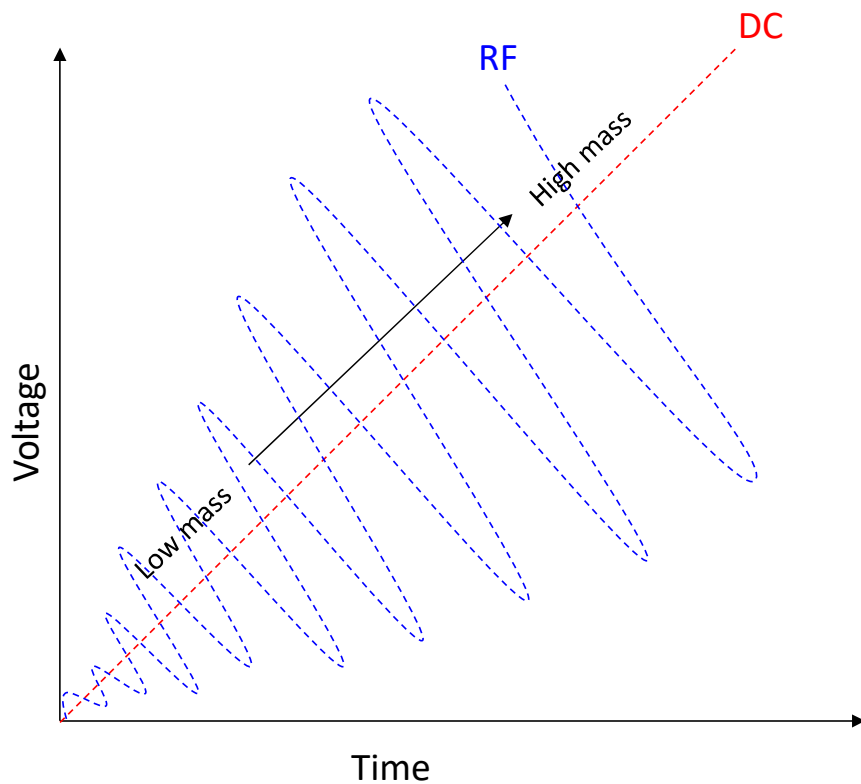


- The ion oscillates in the volume between the quadrupoles
- A **negative potential** at the **exit end** of the quadrupole can be used to pull the ion through the quadrupole



# Mass Spectrometry

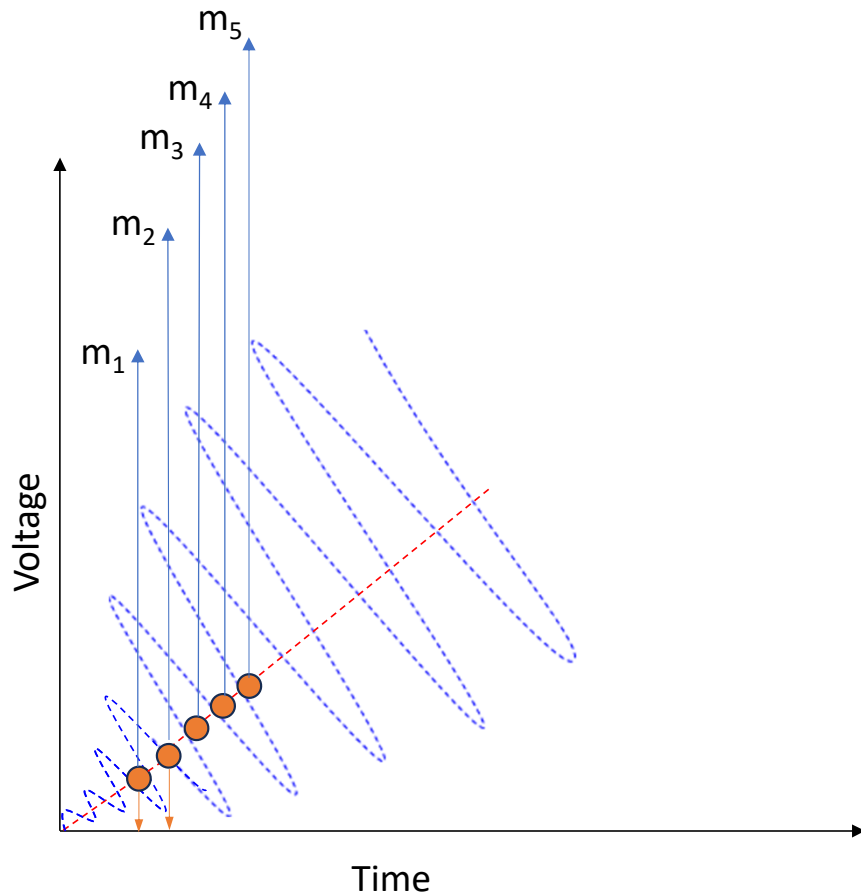
## Mass analysis by the quadrupoles (1)



- When RF and DC are applied together, Quadrupoles function as a mass analyzer.
- The RF-DC voltages can be scanned
  - DC voltage increases with time
  - RF amplitude increases with time, but frequency remains constant
- For one RF-DC combination, only an ion of one particular mass will have a stable trajectory through the quadrupoles.
- Technically, it is the mass/charge ( $m/z$ ) that is affected

# Mass Spectrometry

## Mass analysis by the quadrupoles (2)



RF-DC combinations in the quadrupole can be used to filter ions by mass:

- ↳ Low RF-DC combination, ions of low mass transmitted.
- ↳ High RF-DC combination, ions of high mass is transmitted

# Mass Spectrometry

## Mass analysis by the quadrupoles (3)

Quadrupoles can function in two modes

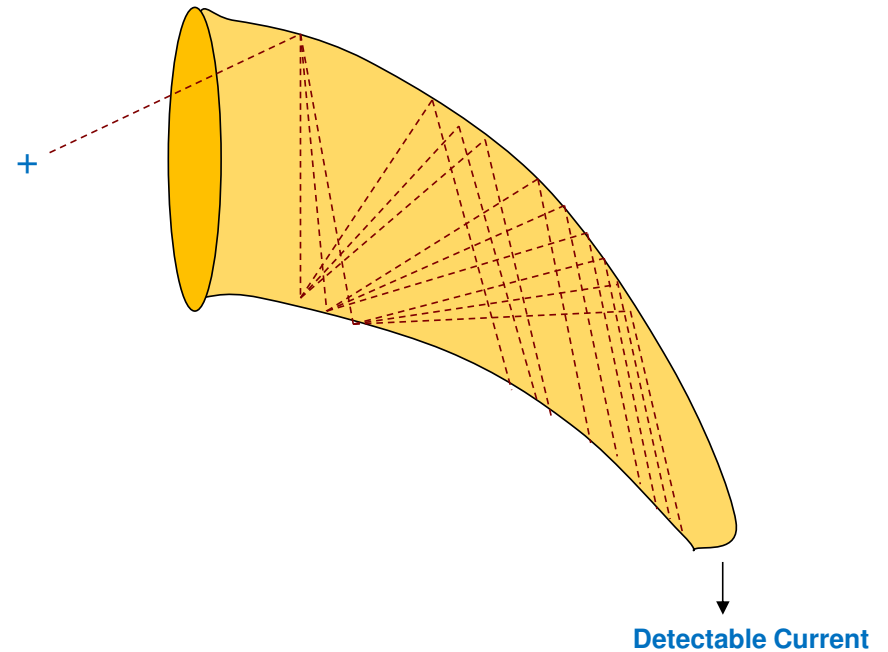
- ↳ **Fixed** RF-DC : Only ions with a fixed mass will be transmitted
- ↳ Switching DC off, i.e. **RF only**:
  - ↳ All ions oscillate within the quadrupole
  - ↳ By keeping a potential at the other end of the quadrupole, **all ions can be transmitted together** through the quadrupole

# Mass Spectrometry

## Detection of ions

### Electron Multiplier

- ↳ The sides of the detector are coated with material that produces electrons on impact by charged particles



# Mass Spectrometry

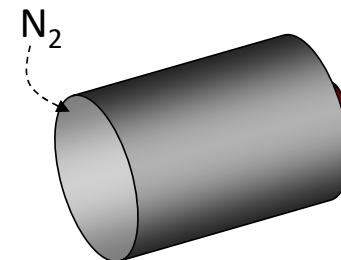
Putting everything together

- Multiple Reaction Monitoring
- Based on the theory that, under the same conditions,
  - An ion from an analyte will always fragment in the same way
  - Fragmentation pattern will be unique
- Highly selective and sensitive
  - We can choose to look at only molecules of interest
  - Quantification in ppb levels

# Mass Spectrometry

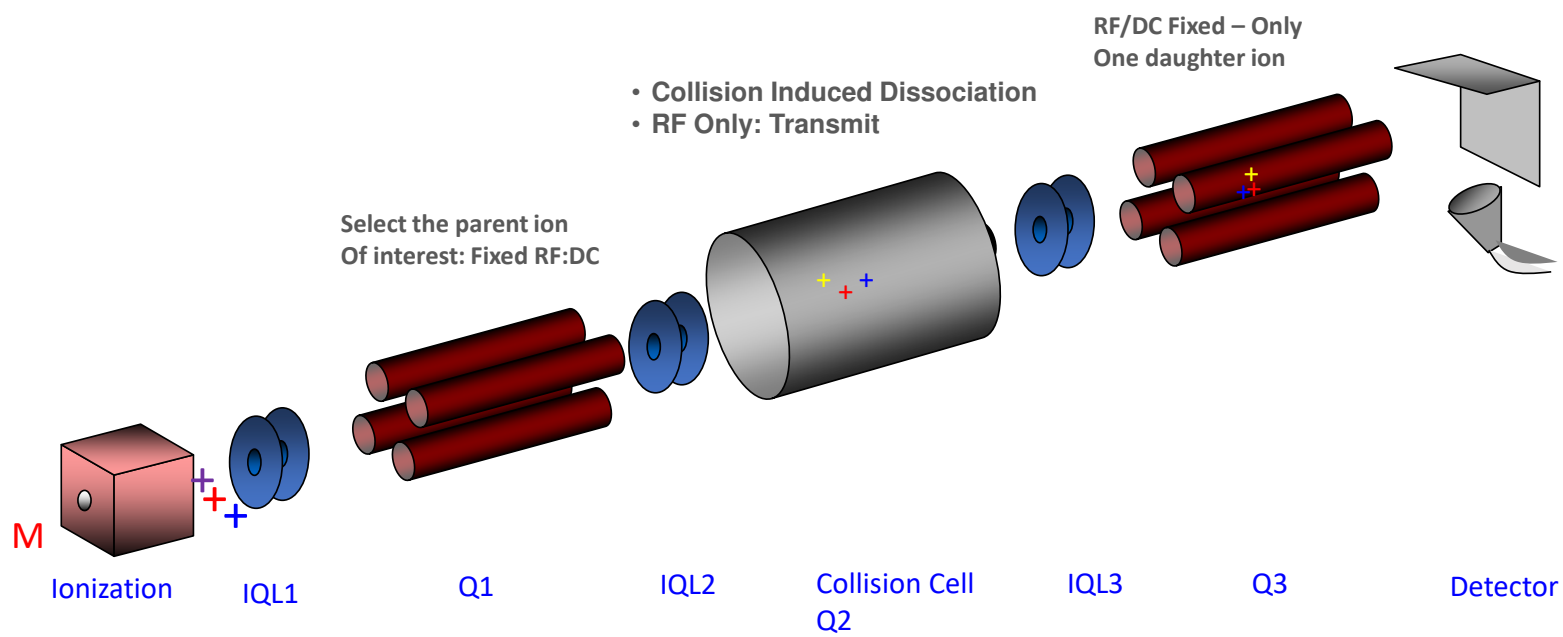
## Putting everything together (2)

- Three Quadrupoles:
  - **First Quadrupole** will select the parent ion  $MH^+$
  - **Second Quadrupole** functions as the **collision cell**:
    - $N_2$  is infused into the collision cell
    - Daughter ions produced by **Collision Induced Dissociation (CID)**
  - **Third Quadrupole** scans the daughter ions
- Linear ion trap:
  - Second quadrupole ejects all ions except the **most intense** ion
  - Can study how this ion **behaves** with time
  - Trapped ion can be fragmented further by CID
- We have AB Sciex Qtrap<sup>®</sup> instrument at QEL Kochi but we are not using it in routine analysis



# Mass Spectrometry

## Putting everything together (3)



# Mass Spectrometry

§ in practice

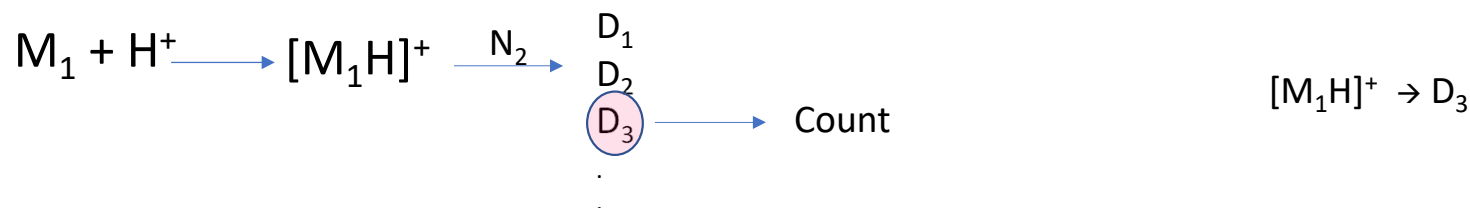


# Multiple Reaction Monitoring

## Quantitative analysis using MS/MS

“Transitions”

### LC MS/MS



Multiple such reactions, or transitions, can be analysed in a single run – hence the name ‘MRM’

# Multiple Reaction Monitoring

## Quantitative analysis using MS/MS (2)





In MRM, we choose 2 transitions per analyte

- ↳ Higher intensity transition is called the **quantifier**, and is used for quantification
- ↳ Lower intensity transition is called the **qualifier**
- ↳ **Quantifier : Qualifier ratio** should match between **standard and sample**

|          | Compound | Parent Ion | Daughter Ion | Type       | Ratio |
|----------|----------|------------|--------------|------------|-------|
| LC MS/MS | Ethion   | 385        | 199          | Quantifier | 28.6  |
|          |          | 385        | 171          | Qualifier  |       |

# MS/MS scan modes

## Not just MRMs

|                         | $M_1$ | Q1                             | Q2                                 | Q3  | Detector  |  |
|-------------------------|-------|--------------------------------|------------------------------------|---|---|--|
| Product ion scan        | $M_2$ | $M_1$<br><i>Fixed m/z</i>      | $d_1, d_2, d_3$                    | $d_1, d_2, d_3$<br><i>Scan</i>  |    | To understand the ways in which an ion can fragment. First step in MS method development |
| Selected ion monitoring | $M_3$ | $M_1$<br><i>Fixed m/z</i>      | $d_1, d_2, d_3$                    | $d_1$<br><i>Fixed m/z</i>   |    | MRM mode,<br>Mainly used for quantitation work   |
| Neutral loss Scan       |       | $M_1, M_2, M_3$<br><i>Scan</i> | $d_1, d_2, d_3$<br>$d_4, d_5, d_5$ | $d_1-x$<br>$d_2-x...$<br><i>Scan</i><br>(only ions that lost an $x$ fragment) |   | To study reaction mechanisms   |
| Precursor ion Scan      |       | $M_1, M_2, M_3$<br><i>Scan</i> | $d_1, d_2, d_3$<br>$d_4, d_5, d_5$ | $d_1$<br><i>Fixed m/z</i>   |  | To find which ions (precursors) produce a given daughter ion                             |

# Topics that remain to be covered in LC-MS/MS

Mass accuracy and resolution, calibration of mass-response, MS tuning

↳ Will be covered in the method validation class

Basic troubleshooting in LC-MS/MS

↳ There will be a separate class on this

# More to learn..

Next session 10<sup>th</sup> June 2023

- ☑ (1) Introduction to instrumentation: LC-MS/MS, 03-Jul-23
- 🕒 (2) Introduction to instrumentation: GC-MS/MS, 10-Jul-23 ←
- 🕒 (3) Pesticide residue analysis - Introduction, 17-Jul-23
- 🕒 (4) Advanced pesticide residue analysis, 24-Jul-23
- 🕒 (5) Method validation: requirements and practice, 31-Jul-23
- 🕒 (6) Introduction to measurement uncertainty calculation, 7-Aug-23

**Dr Ramesh BN**

Scientist C, Spices Board

WhatsApp : +91 8547662471

Mobile : +91 8547662471

Email (O) : [rameshbabu.n@nic.in](mailto:rameshbabu.n@nic.in)

Email (P) : [rameshbn1@gmail.com](mailto:rameshbn1@gmail.com)

Course website : <http://ramesh.-sb.github.io>

**Thank you!**  
§ Questions?