# Mass Spectrometry (MS)

from chapter(s) \_\_\_\_\_ in the recommended text

# A. Introduction

#### **B. Components Of Mass Spectrometers**

molecules on the basis of the ratio of *mass-to-charge*.

primary objective of MS is to determine the *molecular mass* of that entity from the complete molecule *without* fragmentation).

Most mass spectrometers are able to

- create ions in the gas phase
- separate ions on the basis of *m/z* (*ie* an analyzer)
- detect the number of ions of each *m/z*

three basic components to a mass spectrometer an ionization source / an analyzer / a detector.

Electrospray (ESI) is a form of *ionization*.

Quadrupoles are components used for analysis.

*Time-of-flight (TOF)* is a form *analysis*.

Quadrupole ion traps are components used for analysis.

Matrix assisted laser desorption (MALD) is a form of ionization.

Fourier transform (FT) is a form of analysis.

Electron Impact (EI) is a form of *ionization*.

Detectors in MS are usually *electron multipliers*.

Thus, *MALDI-TOF* is a valid description but *ESI-MALDI* is not.

# **C. Primary Ions Formed In Different Ionization Techniques**

In *matrix-assisted lased desorption ionization*, or <u>MALDI</u>, the sample is adsorbed The matrix transfers energy to the sample and mainly *protonates* it to give a cation, *ie*  $[M + H]^+$ .

In *electrospray ionization*, or <u>ESI</u>, the sample in a solvent (*eg* water) is sprayed be protonated by the solvent giving  $[M + 1]^+$  and ions with more than one proton.



Electron impact, removes an electron from molecules to give radical cations

MM as the sample, provided there is no fragmentation, ie  $[M]^{++}$ .

*Resolution* is important in MS when trying to distinguish two materials of similar molecular weights.

Observation of signals when working with tiny amounts of substrate is a question of sensitivity.

It is one of the *most* sensitive forms of MS.

# D. Isotopes In Mass Spectrometry

Element	Isotope	Abundance (%)	Mass number	Exact mass
hydrogen	ЪН	99.99	1	1.00783
carbon	<sup>12</sup> C	98.89	12	12.00000
carbon	<sup>13</sup> C	1.11	13	13.00335
nitrogen	<sup>14</sup> N	99.64	14	14.00307
oxygen	<sup>16</sup> O	99.76	16	15.99492
fluorine	<sup>19</sup> F	100	19	18.99840
phosphorus	<sup>31</sup> P	100	31	30.97376
sulfur	<sup>32</sup> S	95.00	32	31.97207
chlorine	<sup>35</sup> Cl	75.77	35	34.96886
chlorine	<sup>37</sup> Cl	24.23	37	36.96590
bromine	<sup>79</sup> Br	50.69	79	78.91835
bromine	<sup>81</sup> Br	49.31	81	80.91635
iodine	127	100	100	126.904468

CO

 $C_2H_4$ 

C<sub>2</sub>H<sub>5</sub><sup>35</sup>Cl

27.99492

28.03132

64.00801

exact mass

exact mass

exact mass

CH <sub>3</sub> <sup>79</sup> Br	CH <sub>3</sub> <sup>81</sup> Br	C <sub>2</sub> H <sub>5</sub> <sup>37</sup> Cl
93.94184	95.93984	66.00505
exact mass	exact mass	exact mass

A high resolution mass spectrometer can distinguish

containing natural chlorine are separated by 1.99704 atomic mass units (amu's) in a ratio of 3.13:1, and compounds containing natural bromine are separated by 1.99800 amu's in a ratio of 1.03:1.

containing *two* bromines will have 3 molecular ions in a 1:2:1 containing *three* bromines will have 4 molecular ions in a 1:3:3:1

#### Illustrative Interpretation Of Isotopes In MS

the chlorine-containing compound A is number: \_\_2\_\_\_\_\_ the bromine-containing compound B is number: \_\_1\_\_\_\_ the non-halogenated compound C is number: \_\_\_3\_\_\_\_



#### **2** (m/z = 112 and 114):



**3** (m/z = 107):



containing odd numbers of nitrogen atoms (1, 3, 5 etc) always have odd molecular ion m/z values.

#### **E.** Fragmentation

dissociate into smaller cations and radicals

the *most* stable one that is most likely to be observed.

*Electron impact* methods usually show fragmentation, but *ESI* does not.

*ESI* is widely used in contemporary MS, but *EI* instruments are becoming less important.

Fragmentation is usually *undesirable* because observation of the molecular ion is the most important and this *is* useful when complementary methods

observing molecular ions and fragmentation patterns is MS/MS.

*Tandem* mass spectrometry allows observation of molecular ions from peptides and proteins



-Me•

+ СН3



















\_\_\_\_\_

#### The McLafferty Rearrangement



This gives cleavage of the bond between the *and* fragments

All these answers can also be shown as a movement in single electron steps corresponding to radical reactions and using fishhook arrows.

H P +	McLafferty rearrangement	+	H <sub>O</sub>

alkene

radical cation





Molecule on the left can undergo *McLafferty rearrangement* since it has -hydrogen whereas molecule on the right does not.