

Mass Spectrometry (MS)

- An analytical technique for measuring the mass-to-charge ratio (*m/z*) of ions in the gas phase
 - Mass spectrometry is our most valuable analytical tool for determining accurate molecular masses

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Also can give information about structure

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- Proteins can now be sequenced by MS

Mass Spectrometry (MS)

- A technique for measuring and analyzing molecules, that involves introducing enough energy into a (neutral) target molecule to cause its ionization and disintegration.
- The resulting primary ions and their fragments are then analyzed, based on their mass/ charge ratios, to produce a "molecular fingerprint."

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Mass Spectrometry (MS)

- Ionization implies a chemical process induced by physical methods.
- The sample is consumed during the measurement.
- There is no defined stimulation of molecular energy levels through interaction with electromagentic radiation, where you can get the sample back without modification.

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Mass Spectrometry (MS) A mass spectrometer bombards a substance under investigation with an electron beam and quantitatively records the result as a spectrum of positive ion fragments. This record is a Mass Spectrum. A mass spectrum is a presentation of the masses of the positively charged fragments vs. their relative concentration. Separation of the positive charge ion fragment is on the basis of mass. (Mass/Charge)

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Mass Spectrometry

- Molecular weight can be obtained from a very small sample.
- It does not involve the absorption or emission of light.
- A beam of high-energy electrons breaks the molecule apart.
- The masses of the fragments and their relative abundance reveal information about the structure of the molecule.

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MS How It Works

 The physics behind mass spectrometry is that a charged particle passing through a magnetic field is deflected along a circular path on a radius that is proportional to the mass to charge ratio, m/e.



MS How It Works

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- The collection of ions is then focused into a beam and accelerated into the magnetic field and deflected along circular paths according to the masses of the ions.
- By adjusting the magnetic field, the ions can be focused on the detector and recorded.

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- 3 fundamental parts:
 - Ionisation Source,
 - Analyser
 - Detector
- Samples easier to manipulate if ionised
- Separation in analyser according to mass-tocharge ratios (m/z)
- Detection of separated ions and their relative abundance
- Signals sent to data system and formatted in a m/z spectrum
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Separation of Ions

- Only the cations are deflected by the magnetic field.
- Amount of deflection depends on *m/z*.
- The detector signal is proportional to the number of ions hitting it.
- By varying the magnetic field, ions of all masses are collected and counted.

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Mass Terminology

Molecular ion	The ion obtained by the loss of an electron from the molecule
Base peak	The most intense peak in the MS, assigned 100% intensity
M+	Symbol often given to the molecular ion
Radical cation	*ve charged species with an odd number of electrons
Fragment ions	Lighter cations formed by the decomposition of the molecular ion. These often correspond to stable carbcations.
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- Suppose M_1^+ is the parent ion & m_1^+ is the daughter ion and if the reaction $M_1^+ \rightarrow m_1^+$ takes place in the source, then the daughter ion, m_1^+ may travel the whole analyzer region & is recorded as m_1^+ ion.
- On the other hand, if the transition M_1^+ to m_1^+ occurs after the source exist and before arrival at the collector, then m_1^+ is called a metastable ion.

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The symbol m^* should be used to indicate the apparent mass of the product ions giving rise to this peak.

• Traditionally, the peak itself has been called a **metastable peak** and this should preferably be expressed as **metastable ion peak**.

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Resolution & Resolving Power

- Width of peak indicates the resolution of the MS instrument
- The better the resolution or resolving power, the better the instrument and the better the mass accuracy
- · Resolving power is defined as:



 M is the mass number of the observed mass (ΔM) is the difference between two masses that can be separated

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• C ₃ H ₆ C) and C_3H_8O	have no	minal ma	asses of 58 and
60, ar	id can be dist	inguisheo	d by low-r	resolution MS
 C₃H₈C) and $C_2H_4O_2$	have no	minal ma	sses of 60
 distinct 	uich hetweer	them h	/ high_res	olution MS
			/ 111011-163	
aloung	Juisii Detweer	r them by	/ mgn-rea	
aloting	Molecular	Nominal	Precise	
uloung	Molecular Formula	Nominal Mass	Precise Mass	
uloun y	Molecular Formula C ₃ H ₈ O	Nominal Mass 60	Precise Mass 60.05754	
	$\frac{\text{Molecular}}{\text{Formula}}$ C_3H_8O $C_2H_4O_2$	Nominal <u>Mass</u> 60 60	Precise <u>Mass</u> 60.05754 60.02112	

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 virtually 	all	Element	Atomic weight	Isotope	Mass (amu)	Relative A bun dance
common	to	hydrogen	1.0079	$^{1}_{^{2}H}$	1.00783 2.01410	100 0.016
organic		carbon	12.011	¹² C ¹³ C	12.0000 13.0034	100 1.11
compounds	are	nitrogen	14.007	¹⁴ N ¹⁵ N	14.0031 15.0001	100 0.38
mixtures	of	oxygen	15.999	¹⁶ O ¹⁸ O	15.9949 17.9992	100 0.20
isotopes		sulfur	32.066	³² S ³⁴ S	31.9721	100
		chlorine	35.453	³⁵ Cl	34.9689	100
		bromine	79.904	⁷⁹ Br ⁸¹ Br	50.9659 78.9183 80.9163	52.5 100 08.0

Molecules with Heteroatoms

- · Isotopes: present in their usual abundance.
- Hydrocarbons contain 1.1% C-13, so there will be a small M+1 peak.
 - If Br is present, M+2 is equal to M⁺.
 - If CI is present, M+2 is one-third of M⁺.
 - If iodine is present, peak at 127, large gap.

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- If N is present, M⁺ will be an odd number.
- If S is present, M+2 will be 4% of M⁺.

Alkyl Halides

the M + 2 Peak

- Most elements have one major isotope.
- Chlorine has two common isotopes, ³⁵Cl and ³⁷Cl, which occur naturally in a 3:1 ratio.
 - Thus, there are two peaks in a 3:1 ratio for the molecular ion of an alkyl chloride.
 - The larger peak, the M peak, corresponds to the compound containing the ³⁵Cl.
 - The smaller peak, the M + 2 peak, corresponds to the compound containing ³⁷Cl.
 - Thus, when the molecular ion consists of two peaks (M and M + 2) in a 3:1 ratio, a CI atom is present.

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 these fragments may be unstable as well and, in turn, break apart to even smaller fragments www.anilmishra.name













Electron Impact Ionization

- Sample introduced into instrument by heating it until it evaporates
- Gas phase sample is bombarded with electrons coming from rhenium or tungsten filament (energy = 70 eV)
- Molecule is "shattered" into fragments (70 eV >> 5 eV bonds)

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Fragments sent to mass analyzer















Chemical Ionization

- El is not appropriate for some molecules (it causes too much fragmentation)
- Instead, ionize a reagent gas (by EI) then react it with a analyte molecules
- Typically use methane or ammonia for reagent gas

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- CI is easiest to envisage as a two stage process.
- In the first step a reagent gas at relatively high pressure (~ 1 Torr) is subjected to electron impact ionization.
- The products of the EI process at high pressure can be complex mixtures of ions, radicals and neutrals, but some gases produce major products that are favorable for the CI process.

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(FAB) or ions (SIMS) is focused on the liquid or solid sample. The impact of this high energy beam causes the analyte molecules to sputter into the gas phase and ionize in a single step.

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- Ionization method whereby a solution prepared by dissolving sample compounds in a liquid matrix, such as glycerin, is spread on a metallic target.
- The target is then bombarded by neutral atoms (Ar, Xe, Cs) accelerated to 2kV ~ 30kV to generate ions.
- The method is further categorized as solid FAB when the sample is a solid, matrix FAB when a matrix solution is used, and gas-phase FAB when the sample is a gas.

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Electrospray ionization

- When a solution previously separated by charge is sprayed, static droplets with remarkable charge balance disruptions are generated.
- When the solvent in these droplets is evaporated (desolvation) by a heated gas or other method, the sizes become even smaller.
- When the Rayleigh limit is reached, the droplets are automatically destroyed by Coulomb's repulsion force and ions are generated.
- One feature of this method is the ability to generate multiply charged ions by protonation or deprotonation. The method is one of the softest ionization methods available.

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Ion Formation By ESI

Electrospray ionization





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MALDI

Identity of suitable matrix compounds is determined using specific molecular design considerations

- fairly low molecular weight (to allow facile vaporization) large enough (with a high enough vapor pressure) not to
- evaporate during sample preparation or while standing in the spectrometer
- are acidic / act as a proton source to encourage ionization of the analyte
- have strong absorption in the UV so they rapidly and efficiently absorb the laser irradiation
- functionalized with polar groups allowing use in aqueous solutions

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MALDI

- MALDI is based on the bombardment of sample molecules with a laser light to bring about sample ionization.
- The sample is pre-mixed with a highly absorbing matrix compound for the most consistent and reliable results and a low concentration of sample to matrix works best.
- The matrix transforms the laser energy into excitation energy for the sample, which leads to sputtering of analyte and matrix ions from the surface of the mixture.
- In this way energy transfer is efficient and also the analyte molecules are spared excessive direct energy that may otherwise cause decomposition.
- Most commercially available MALDI mass spectrometers now have a pulsed nitrogen laser of wavelength 337 nm.

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MALDI

- The sample to be analysed is dissolved in an appropriate volatile solvent, usually with a trace of trifluoroacetic acid if positive ionisation is being used, an aliquot of this removed and mixed with an equal volume of a solution containing a vast excess of a matrix.
- A range of compounds is suitable for use as matrices:
- Sinapinic acid is a common one for protein analysis
 Alpha-cyano-4-hydroxycinnamic acid is often used for peptide analysis.
 An aliquot of the final solution is applied to the sample target which is allowed to dry prior to insertion into the high vacuum of the mass spectrometer.
- The laser is fired, the energy arriving at the sample/matrix surface optimized, and data accumulated until a m/z spectrum of reasonable intensity has been amassed.
- The time-of-flight analyser separates ions according to their mass(m)-to-charge(z) (m/z) ratios by measuring the time it takes for ions to travel through a field free region known as the flight, or drift, tube. The heavier ions are slower than the lighter ones.

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MALDI

- · Matrix solution is mixed with the analyte (e.g. protein-sample)
 - organic solvent allows hydrophobic molecules to dissolve
 - water allows for hydrophilic molecules to do the same
- Solution is spotted onto a MALDI plate
- solvents vaporize, leaving only the recrystallized matrix
- analyte molecules spread throughout the crystals in cocrystallized MALDI spot

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Interpretation of Mass Spectra

- Most mass spectra are not trivial to decipher due to:
 - interferences
 - complexities of fragmentation
 - (exception: some elemental analysis cases)
- Instead, MS Interpretation is problem solving, 'playing chess', or 'cracking a code'

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Calculation of Molecular Formula

- · After establishing the basic formula with only Carbon/hydrogen,
- Other element can be introduced by substracting the proper hydrocarbon value

Hetero substitution	CH replacement	Hetero substitution	CH replacement
0	CH ₄	Р	C ₂ H ₇
Ν	CH ₂	S	C ₂ H ₈
O+N	C ₂ H ₆	O+S	C ₄
F	CH ₇	1	C ₁₀ H ₇
Si	C_2H_4	Cl,Br	(use isotopes)
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Interpretation of Mass Spectra

- Select a candidate peak for the molecular ion (M⁺)
- Examine spectrum for peak clusters of characteristic isotopic patterns
- Test (M⁺) peak candidate by searching for other peaks correspond to reasonable losses
- Look for characteristic low-mass fragment ions
- · Compare spectrum to reference spectra

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Summary

- The substitute groups like -OH, -OR, -NH2, etc., which lower the ionization potential increase the relative abundance in case of aromatic compounds. Also groups like -NO2, -CN, etc., which increase ionization potential, decrease the relative abundance of the aromatic compounds.
- Absence of molecular ion peak in the mass spectrum means that the compound under examination is highly branched or tertiary alcohols. Primary and secondary alcohols give very small molecular ion pages peaks.
- In case of Chloro or Bromo compounds, isotope peaks are also formed along with the molecular ion peak. In case of Bromo compounds, M^* & (M^{+} +2) peaks are formed in the intensity ratio 1:3.

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Summary

- Fragment Ion Peak
 When potential and energy is given to the molecular ion during electron impact , further cleavage takes place and ions of lower mass number (fragment ions) are produced which gives the fragment peak.
 Many of the peaks in the mass spectrum are due to fragment ions. Fragment peaks in the spectrum give valuable information regarding molecular structure, because fragmentation is specific to the structure of the molecule.
- The greater the fragmentation of the molecular ion or parent ion, the greater is the loss in its intensity. The probability of fragmentation decreases in the following order: •
 - ALCOHOLS > BRANCHE HYDROCARBONS > CARBOXYLIC A CLOS > ETHERS > ESTERS > AMINES > KETONS >THIOLS > STRAIGHT CHAIN HYDROCARBONS > SULPHIDES > ALLCYCLIC COMPOUNDS > CONJUGATED DIENES-AROMATIC COMPOUNDS.
- Fragmentation is initiated by electron impact. Only a small part of the driving force for fragmentation is energy transferred as the result of the impact. •

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