### **Ionization Methods**

- 1. Gas-Phase Ionization Methods
  - A. Electron Impact Ionization
  - **B.** Chemical Ionization
- 2. Desorption Ionization Methods
  - A. Field Desorption Ionization
  - B. Fast Atom Bombardment (FAB) Ionization
  - C. Plasma Desorption Ionization
  - D. Laser Desorption Ionization
- 3. Evaporative Ionization Methods
  - A. Thermospray Mass Spectrometry
  - B. Electrospray Mass Spectrometry

### **Gas-Phase Ionization Methods**

#### A. Electron Impact Ionization

- El was available before other ionization methods were developed
- Must have a stable molecule in the vapor phase (no salts, no proteins or peptides, sugars) Only ~20% of organic molecules
- Stability of Molecular ions: Aromatics> conjugated olefins > alicyclics > sulfides > unbranched hydrocarbons > mercaptans >ketones > amines > esters > ethers > carboxyllic acids > branched hydrocarbons > alcohols
- Must have a lifetime >10<sup>-6</sup> sec

#### Disadvantages of El

- Must be vaporized to 10<sup>-6</sup> torr (not applicable to fragile low vapor, pressure species e.g. salts, sugars, peptides, proteins, polymers)
- El often leads to extensive fragmentation and no molecular ion is observed
- Rearrangements can cause difficulty in explaining structure.

#### **Advantages of EI:**

- Relatively simple to build
- Fragments seen enables library of EI mass spectra containing several 100K spectra (Note there are about 60,000 cpds. common in commerce.)
- Compatible with GC
- Fragments can tell something about structure
- Ion profiles can confirm presence or absence without examining entire spectrum.

### **B.** Chemical Ionization

- The most important "soft ionization" technique to avoid extensive fragmentation in El.
- In CI: Sample molecules are not subjected to bombardment by high energy electrons
- Reagent gas (usually methane, isobutane, ammonia) is introduced into the source and ionized.
- Sample molecules collide with ionized reagent gas molecules (CH<sub>5</sub><sup>+</sup>, C<sub>4</sub>H<sub>9</sub><sup>+</sup>, etc) and undergoes secondary ionization by:
  - 1. Proton transfer producing an [M+1]<sup>+</sup> ion
  - 2. Electrophilic addition producing [M+15]<sup>+</sup>, [M+43]<sup>+</sup> ion
  - 3. [M+18]<sup>+</sup> with NH<sub>4</sub><sup>+</sup>
- CI sometimes give prominent [M-1]<sup>+</sup> because of hydride abstraction
- <u>Main use: detection of molecular ion and hence</u> <u>molecular weight</u>

1. Electron beam generates proton donor (via El of reagent gas)





2. Proton donor protonates sample  $M + XH^+ \longrightarrow MH^+ + X^-$ Parent mass + 1

### The EI and CI mass Spectra of 3,4dimethoxyacetophenone



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### Nobel Prize 2002

"for their development of soft desorption ionisation methods for mass spectrometric analyses of biological macromolecules"

#### Koichi Tanaka (Japan)



#### John B. Fenn (USA)



### Matrix Assisted Laser Desorption/Ionization (MALDI)

- Limitations of Mass Spectrometry
  - Biomolecules and organic macromolecules are fragile
  - Molecular ions or meaningful fragments were limited to only 5-10 kDa at the time

 In 1987, Michael Karas and Franz Hillenkamp successfully demonstrated the use of a matrix to ionize high molecular weight compounds



Michael Karas Franz Hillenkamp





Laser

- 1. Sample (A) is mixed with excess matrix (M) and dried on a MALDI plate.
- 2. Laser flash ionizes matrix molecules.
- 3. Sample molecules are ionized by proton transfer from matrix:

 $MH^+ + A \rightarrow M + AH^+$ .



Koichi Tanaka Nobel Prize 2002



# MALDI

- Method where a laser is used to generate ions of high molecular weight samples, such as proteins and polymers.
- Analyte is embedded in to crystal matrix.
- The presence of an aromatic matrix causes the large molecules to ionize instead of decomposing.
- MALDI involves incorporation of the analyte into a matrix, absorption/desorption of laser radiation, and then ionization of the analyte.
- "MALDI 'theory' or 'models' are still crude and are far away from having predictive power."--- Michael Karas, Chem Rev. 2003, 103, 427-439
- Challenges for MALDI Theories:
  - It is a complex chemical event
  - Happens on a time scale of nanoseconds
- The MALDI technique combined with a MS detector (MALDI-MS) became an indispensable tool in analysis of biomolecules and organic macromolecules.

### **MALDI** Matrix Substances



#### **MALDI Advantages:**

- Gentle Ionization technique
- High molecular weight analyte can be ionized
- Molecule need not to be volatile
- Sub-picomole sensitivity easy to obtain
- Wide array of matrices

### MALDI Disadvantages:

- MALDI matrix cluster ions obscure low m/z species (<600)</li>
- Analyte must have very low vapor pressure
- Pulsed nature of source limits compatibility with many mass analyzers
- Coupling MALDI with chromatography can be difficult
- Analytes that absorb the laser can be problematic
  Fluorescein-labeled peptides

# **Applications of MALDI**

- Peptides and proteins
- Synthetic polymers
- Oligonucleotides
- Oligosaccharides
- Lipids
- Inorganics
- Bacterial identification
- Proteomics

### **Polymer Analysis**



### **Bacterial Identification**

- MALDI-TOF-MS is used for bacterial identification.
- MALDI-TOF-MS can determine mass of proteins of 1-40 kDa.
- Used to diagnose diseases and monitor contamination.
- Identified by:
  - -Biomarkers
  - -Cellular protein content

#### Mass spectral analysis of protein extracts

Biomarkers for Bacteria Protein Extracts			
Organism	Genus Biomarkers (m/z)	Species Biomarkers (m/z)	Strain Biomarkers (m/z)
Bacillus anthracis,	6680, 6837	2385, 3991,	4505
Bacillus anthracis, sterne	6680, 6837	4313 2385, 3991, 4313	2789
Bacillus anthracis, zimbabwe	6680, 6837	2385, 3991, 4313	2850
Bacillus thuringiensis, 4A1	6680, 6837	3932	5916
Bacillus thuringiensis, 4A2	6680, 6837	3932	4871, 7845
Bacillus thuringiensis, 4L2	6680, 6837	3932	2864, 4074, 4548, 5781
Bacillus cereus, 6E1	6680, 6837	5269, 5537, 7365, 9533	N/A
Bacillus subtilis, 3A1	6680ª	3757, 3871, 6304, 9146	N/A
Brucella melitensis, melitensis wild	7379, 8031, 9060	N/A	N/A
Brucella melitensis, REV-1 vaccine	7354, 8030, 9071	N/A	N/A
Brucella melitensis, Suis wild	7378, 8020, 9080	N/A	5844
Brucella melitensis, abortus wild	7379, 8020, 9088	N/A	N/A
Yersinia pestis, India	2460, 3223, 3297, 4361, 4437	N/A	7335
<i>Yersinia pestis,</i> La Paz	3223, 3297, 4361, 4437	N/A	N/A
Yersinia pestis, Nairobi	2453, 3223, 3297, 4361, 4437	N/A	5196, 5638, 7599, 9637
Francisella tularensis, type B	2622, 2770, 3205, 3274, 4120, 4747	N/A	N/A

<sup>a</sup>Doubly charged ion, corrected mass/charge ratio N/A - Data not available

### Electrospray Ionization Mass Spectrometry



John Fenn Nobel Prize 2002 ESI changed the field of mass spectrometry and opened the world of biology to mass spectrometry and enabled:

- Proteomics
- Metabolomics
- Systems biology

## What Is ESI?

- The transfer and ionization of molecules from solution to gas phase by electrospray (An electrical nebulization of liquid that results in the formation of charged micro droplets)
- Vaporization increases the charge density on the surface of the droplets (Electrostatic repulsion increases).
- When the electrostatic repulsion exceeds the surface tension, the droplet undergoes coulombic fission (Formation of daughter droplets)
- The formation of charged ions in the gas phase

# Electrospray Ionization (ESI)







#### **Heated Inlet to MS**

Requirements for Electrospray Ionization:

- Soluble in solvent (water, methanol, acetonitrile)
- Avoid DMSO and DMF
- Pseudomolecular ion is formed MH<sup>+</sup>,[M+Na]<sup>+</sup>, etc.
- Very Gentle
- MW to megadalton range
- Preformed ions Salts, peptides at low pH
- Electrochemical reactions can take place
- Often need a basic site; otherwise add a cation (Na+)

### **Application of ESI**



### **Bovine Serum Albumin**



#### Mass spectrometry and viral analysis

#### Gary Siuzdak<sup>1\*</sup>, Brian Bothner<sup>1</sup>, Mark Yeager<sup>2</sup>, Christophe Brugidou<sup>3</sup>, Claude M Fauquet<sup>3</sup>, Kenway Hoey<sup>4</sup> and Cheng-Ming Chang<sup>2</sup>

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**Background:** Electrospray ionization (ESI) mass spectrometry is a powerful new approach for analyzing biomolecules and biomolecular complexes. Previous studies have provided evidence that non-covalent biomolecular complexes can be observed by ESI mass spectrometry; it is not clear, however, whether the native conformation of the biomolecules is maintained throughout the ionization and analysis process. We set out to address this question using live viruses.

**Results:** Viral ions have been generated in the gas phase using electrospray ionization mass spectrometry. These ions have been collected, following ion filtering through the mass analyzer, and then analyzed by transmission electron microscopy. Transmission electron microscopy revealed that rice yellow mottle virus and tobacco mosaic virus retained their respective spherical and rod-like ultrastructure. The viability of the isolated tobacco mosaic virus was confirmed by inoculation and infection of tobacco plants.

**Conclusions:** These results demonstrate the utility of electrospray for supramolecular complexes with molecular weights of over 40 million Da and offer conclusive evidence that native biomolecular structures can be conserved through the electrospray process.

#### Chemistry & Biology January 1996, 3:45-48

Key words: electron microscopy, electrospray, mass spectrometry, non-covalent, virus

**BIOPHYSICAL CHEMISTRY** 

# Unravelling capsid transformations

The interactions between a virus capsid and its cargo are essential for viral infection as well as in the design of synthetic virus-like particles. Now a combination of analytical techniques has unravelled key steps in the transformation of a model virus and the release of its RNA cargo.

#### Masaki Uchida and Trevor Douglas



# THANK YOU