



# Brief Concept of LC-MS

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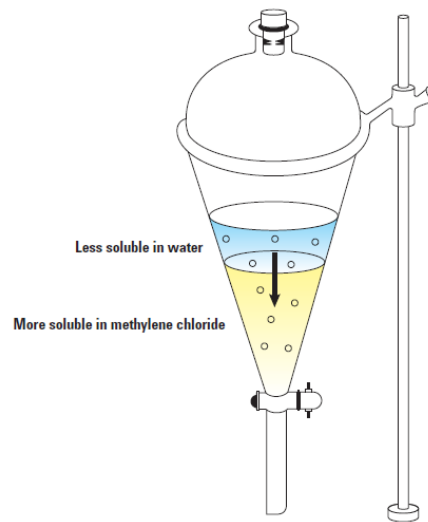
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- How good is your sample preparation
- Quick understand LC
- How to make a good separation
- Understand a short brief of Mass Spectrometer definition
- Connected to Mass Spectrometer
- Type of Mass Spectrometer
- How to choose appropriated Mass Spectrometer

# Liquid-Liquid Extraction

- separating analytes from interferences by partitioning the sample between two immiscible liquids
- One phase is aqueous, another is organic
- More hydrophilic goes to aqueous while as more hydrophobic will be found mainly in organic phase



# LLE: How to choose a solvent

- A low solubility in water (<10%).
- Volatility for easy removal and concentration after extraction.
- Compatibility with the HPLC or GC detection technique to be used for analysis (avoid solvents that are strongly UV-absorbing or that may cause GC detection problems, such as chlorinated solvents in conjunction with electron capture detector).
- Polarity and hydrogen-bonding properties that enhance recovery of the analytes in the organic phase.
- High purity to minimize sample contamination

# LLE: Disadvantage

- Emulsion formation
- Analytes strongly adsorbed to particulates
- Analytes bound to high molecular weight compounds (e.g. protein-drug interactions)
- Mutual solubility of the two phases
- No automate
- High organic consumption
- Not good for complex extraction

# Solid-supported Liquid-Liquid Extraction

- Overcome LLE disadvantage with greater benefits
- Diatomaceous earth particle serves as a stationary phase for aqueous phase
- Aqueous-base sample will added to dry sorbent and dispersed through solid support
- Next, a small volume of immiscible organic solvent is added (required gentle pressure or mild vacuum) to allow partitioning



- Greater reproducibility and recoveries compared to LLE techniques
- Prevents emulsification often associated with LLE
- Reduced solvent requirements compared to LLE
- Can be completely automated unlike LLE
- Improved cleanliness of sample extract compared to protein precipitation techniques
- Improved sensitivity compared to protein precipitation techniques

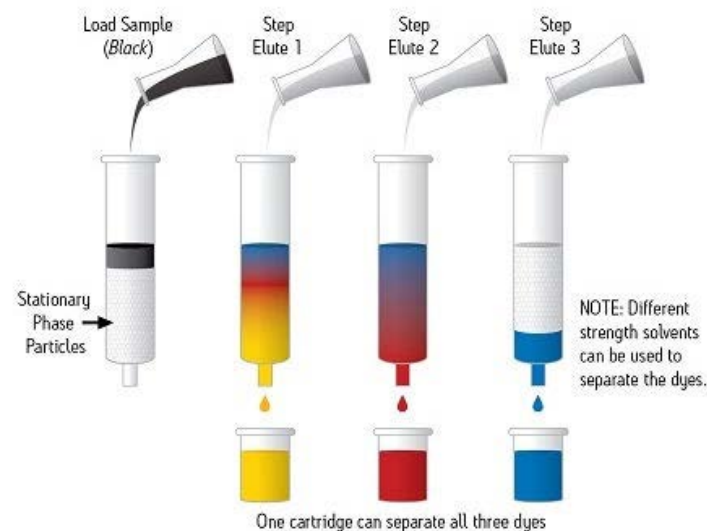
- Can be used only aqua-base sample
- Not significant greater than LLE in term of recovery
- Deal with manifold (req. more complicate method development)
- Sometimes need pre-buffered (Commercially available)



# Solid Phase Extraction

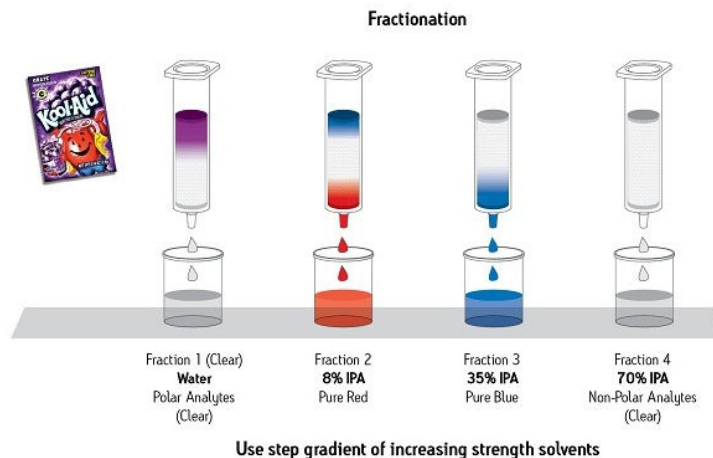
SPE is important chromatographic preparation based on chemically different components in sample.

- Each of components can be eluted by appropriate solvent
- Can be used for gas phase by trapping on sorbent using reactive chemical or specified with some materials
- Known as “Liquid-Solid Extraction”



# SPE Benefits

- Simplification of complex sample matrices along with compound purification
- Reduced Ion Suppression in Mass Spectrometry Technique (Desalting)
- Capability to Fractionate Sample Matrix to Analyze Compounds by Class
- Enrichment of Very Low Level Compounds



# SPE: Disadvantages

- Mix Mechanism can be taken place inside cartridge
- Irreversible adsorption of some analytes on SPE cartridges
- More complex method development is required
- Sometimes need evaporating step

- **Q**uick-**E**asy-**C**heap-**E**ffectiveness-**R**ugged-**S**afe (“catcher”) is now commonly used in pesticide from FOOD analysis both LC and GC
- 2-Processes; extraction followed by clean-up
- Known as “ LL and SPE”



# QuEChERS : Extraction Step

Magnesium Sulfate aids the extraction and remove residue water from organic solvent and unwanted contaminants

- Considerations

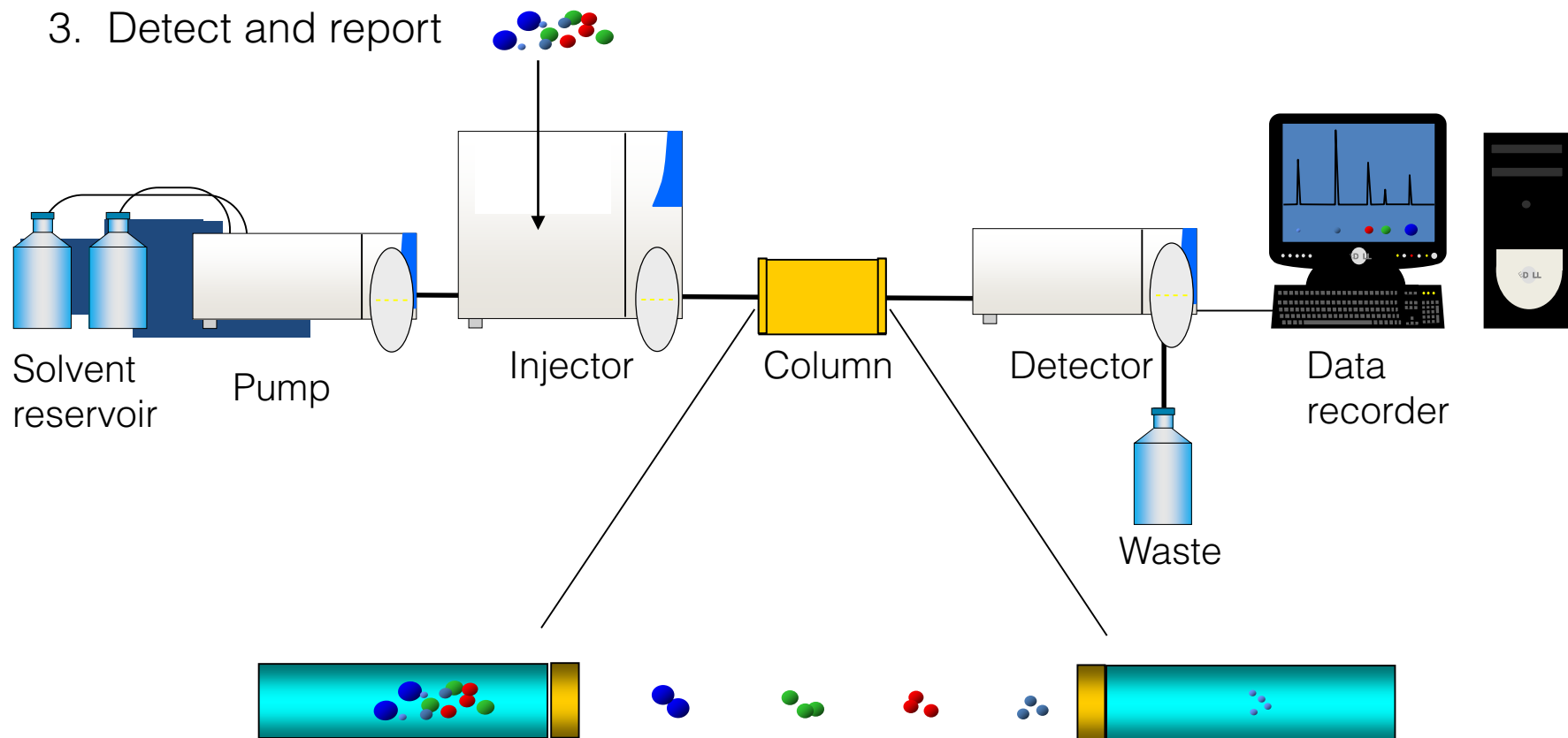
- Base sensitive compound -> with sodium acetate
- Non-base sensitive compound -> with sodium citrate or sodium chloride



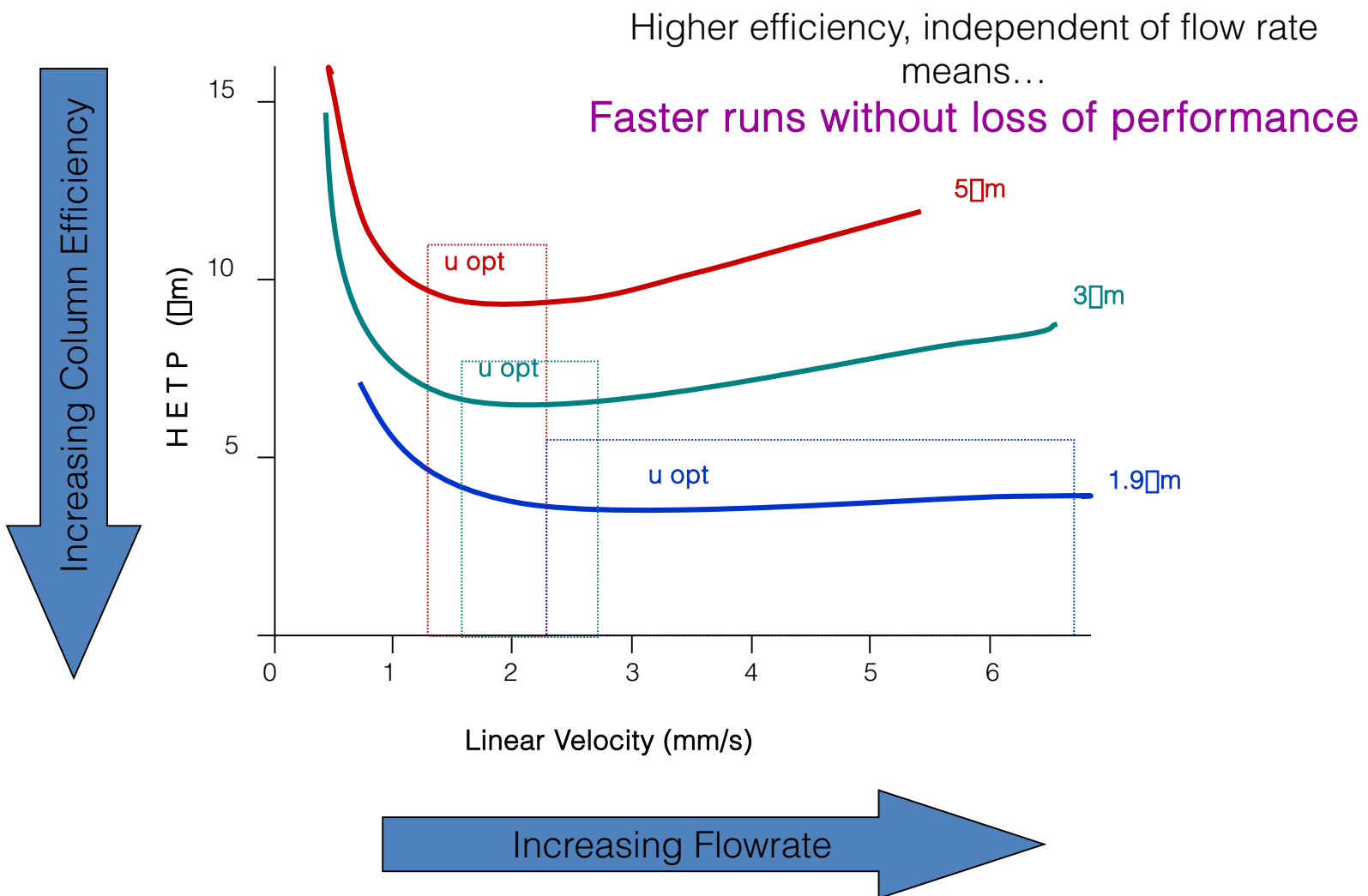
- Determine the properties of sample matrixes
  - General
  - Fatty
  - Pigmented
  - Highly Pigmented
- Adsorbents
  - C18: REMOVE Low fat interference
  - PSA (Primary-Secondary Amine): REMOVE Sugars and organics acid
  - GBC (Graphitized Black Carbon): REMOVE pigmented, chlorophyll, carotenoid etc.

# Quick Understand LC

1. Inject sample mixture
2. Separate into individual components
3. Detect and report



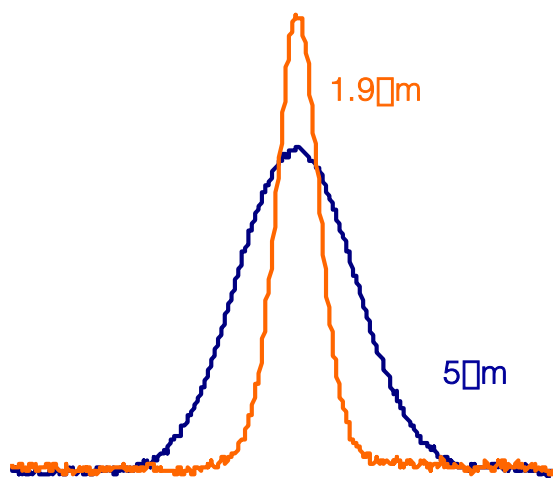
# What is the Small Particle Advantage ?





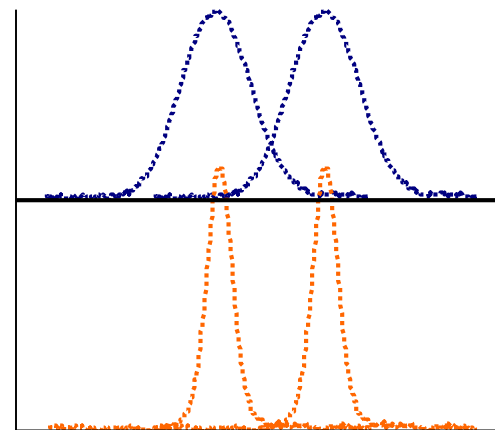
# Efficiency is the key!!!

## Small Particle Advantage



N = 142,000 plates/m  
(189% higher)

N = 75,000 plates /m

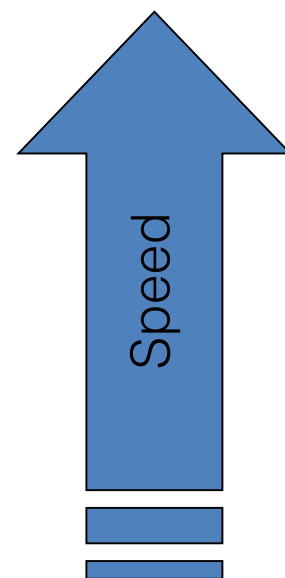
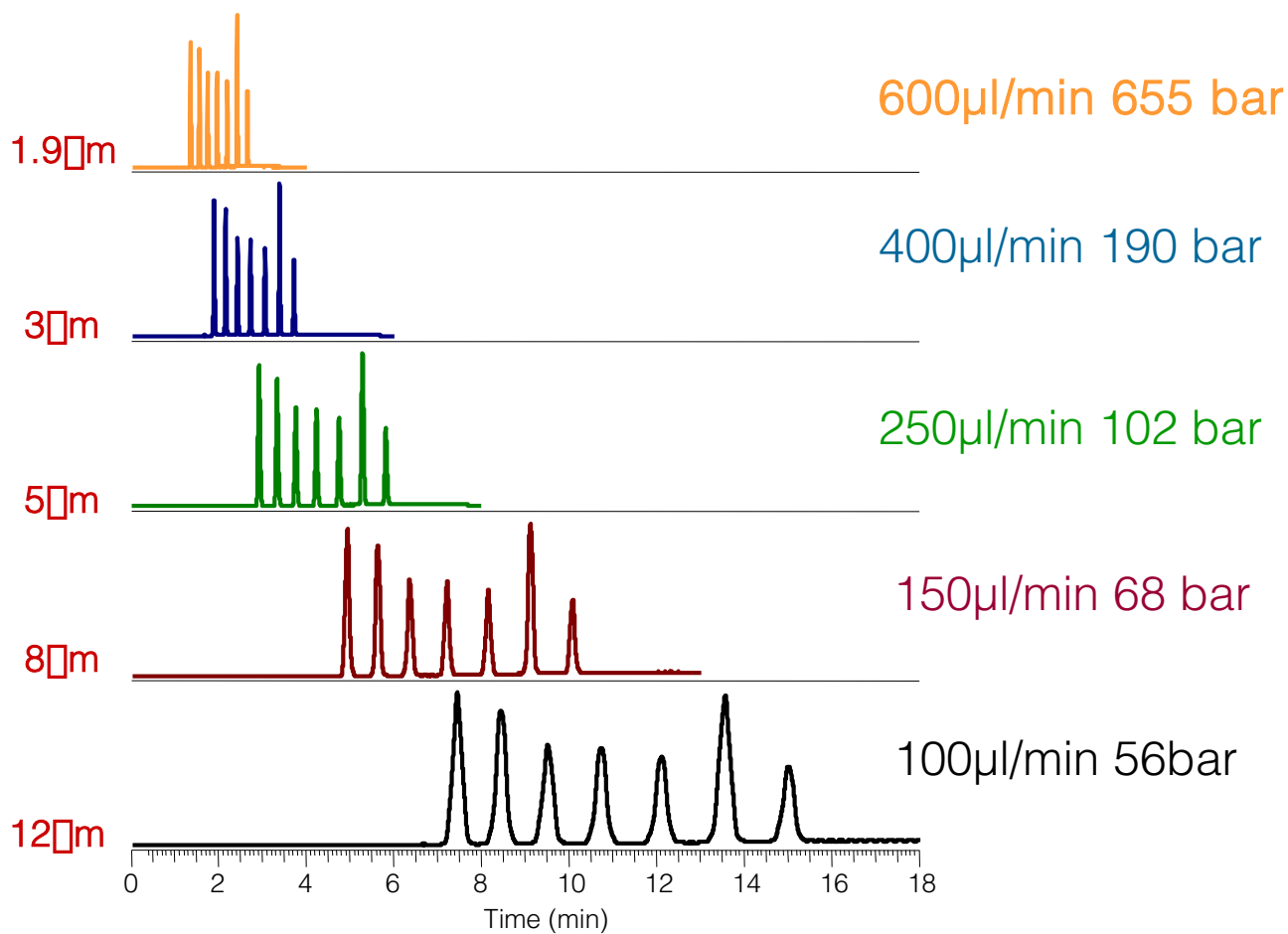


$$R_s = \frac{1}{4} \frac{(\alpha - 1)}{\alpha} \underbrace{\sqrt{N}}_{\text{Efficiency}} \frac{k}{1+k}$$

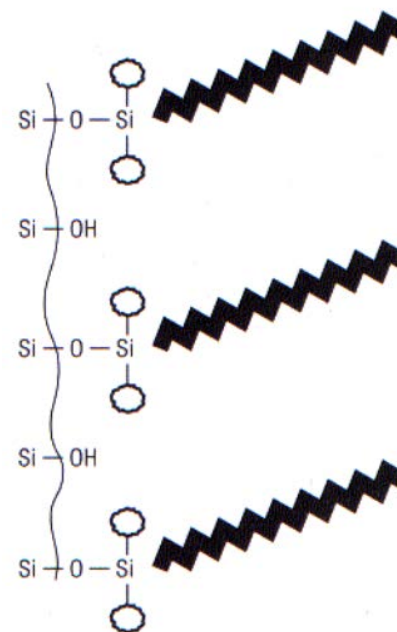
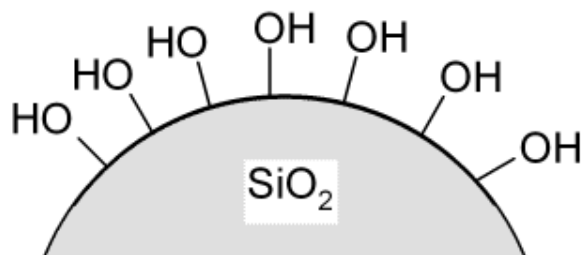
Selectivity      Efficiency      Retention

- Higher resolution – narrower peaks
- Higher sensitivity – taller peaks
- Higher peak capacity (more peaks / unit time) – narrower peaks

*Speeding up analysis with 1.9  $\mu\text{m}$  Hypersil GOLD*



# Wait a minute.....what's column chemistry????!!!

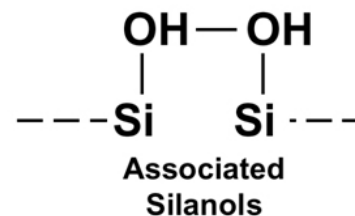
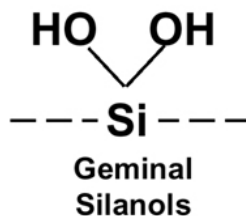
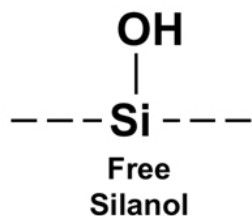


- Diol
- C1, C4, C8, C18
- Aminopropyl
- Nitrile
- Phenyl
- Pentafluorophenyl
- Cation Exchanger
- Anion Exchanger
- etc.

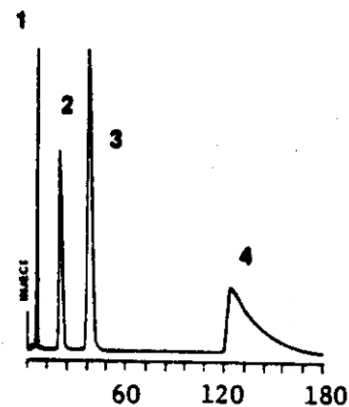
Chemistry is variant. Is all C18 the same ?

NO !

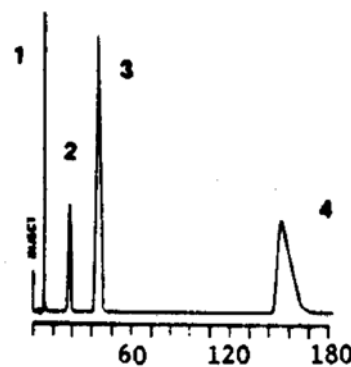
# Un-reacted: Interaction with Basic Molecules



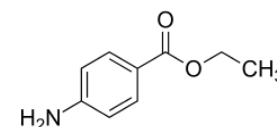
pKa



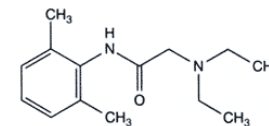
No  
endcapping



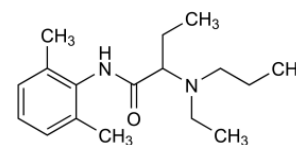
Endcapping



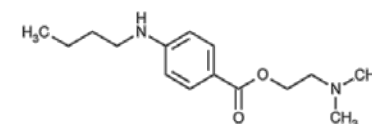
2.8



7.9

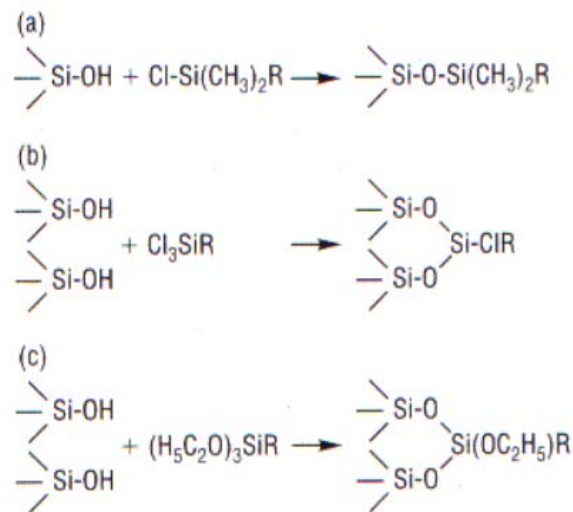
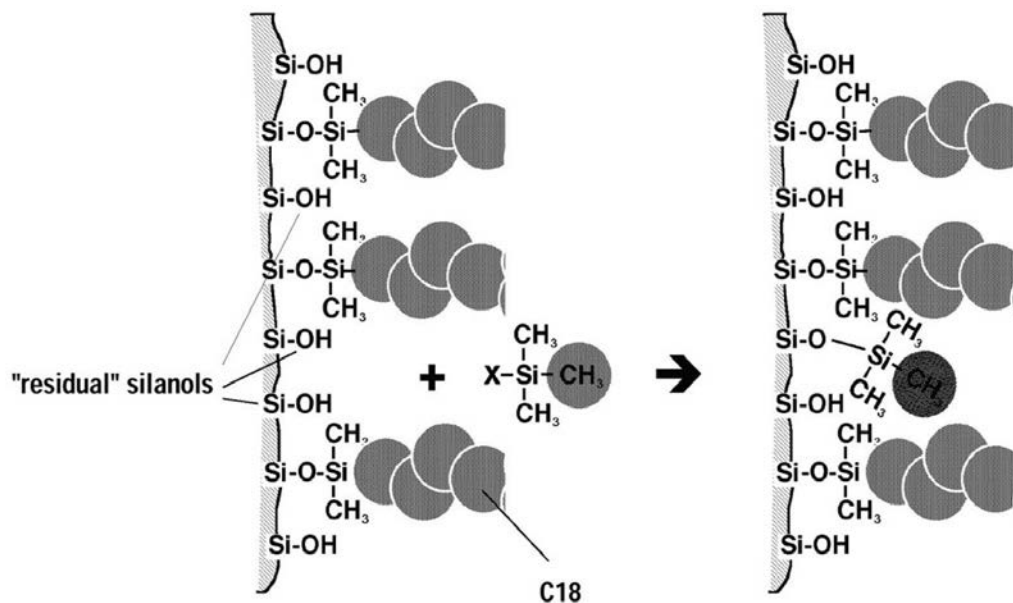


7.9



8.5

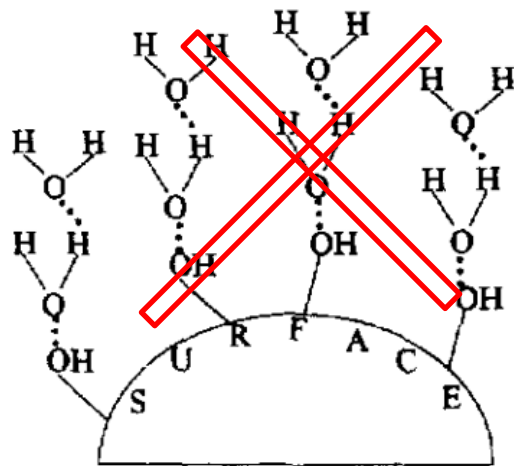
# Endcapping : Prevent peak Tailing & interaction with alkaline



- Polar; amide, urea, ether
- Hydrophilic
- Trimethylsilyl
- etc.

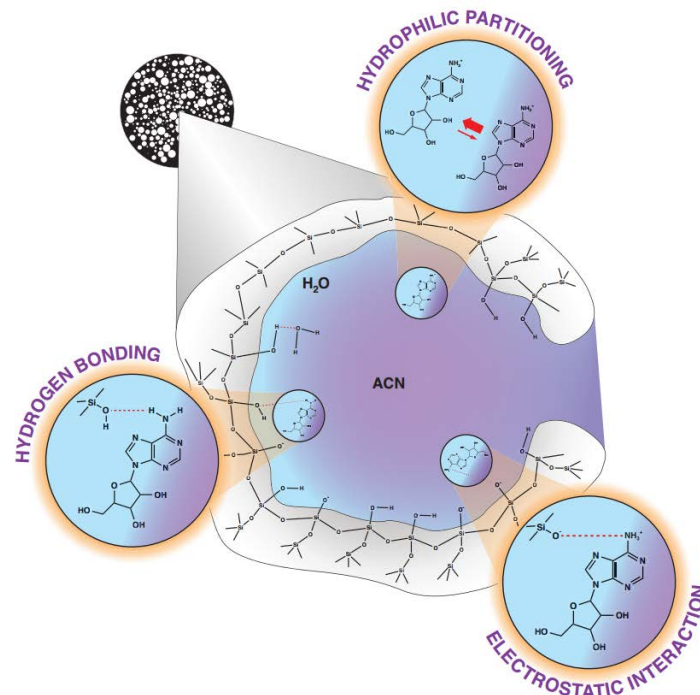
- Dimethyl silane
- Chloro silane
- Trifunction alkoxy silan
- etc.

aQ



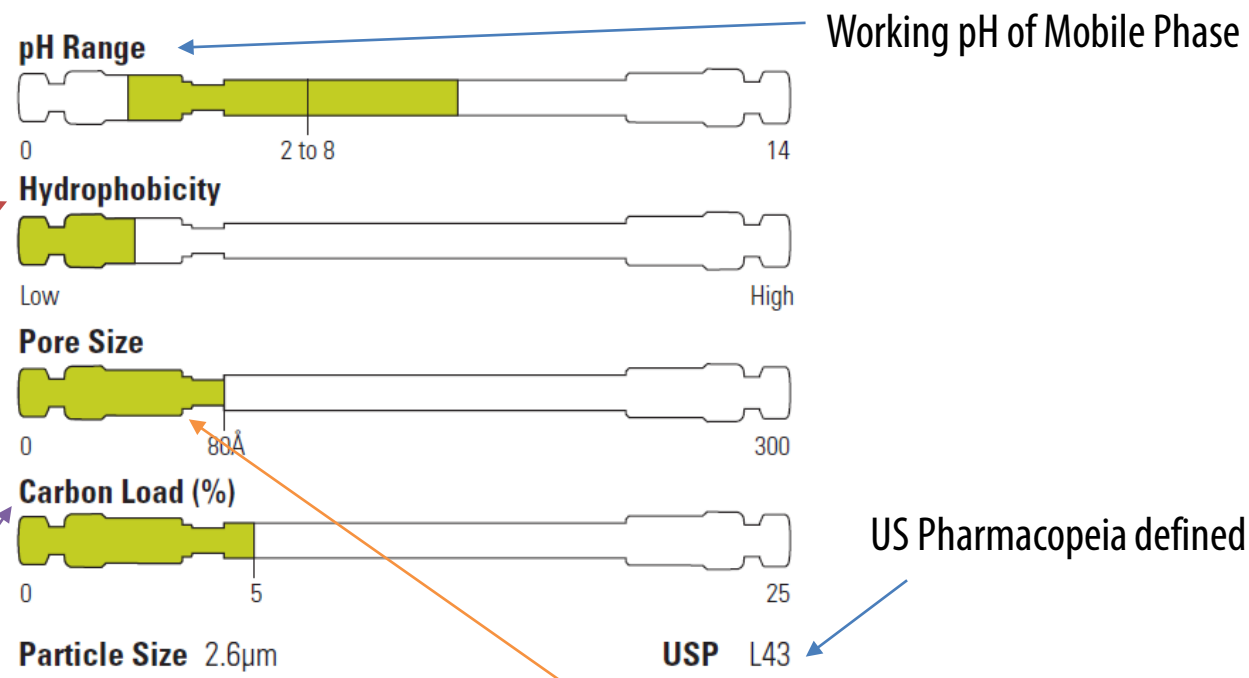
Stable in 100 % Aqueous with polar endcapping,  
Enhance retention of polar compounds

HILIC



Retain highly polar and hydrophilic compound, no endcapping → can't use with more than 50% aqueous

# Column Properties



Correlated to % carbon, reflected to polarity of column

% carbon content, higher is not always better resolution. Higher is more hydrophobic surface that resistance to high pH

60-90 for small molecule  
90-120 for both small molecule and peptide  
120-300 for peptide or protein

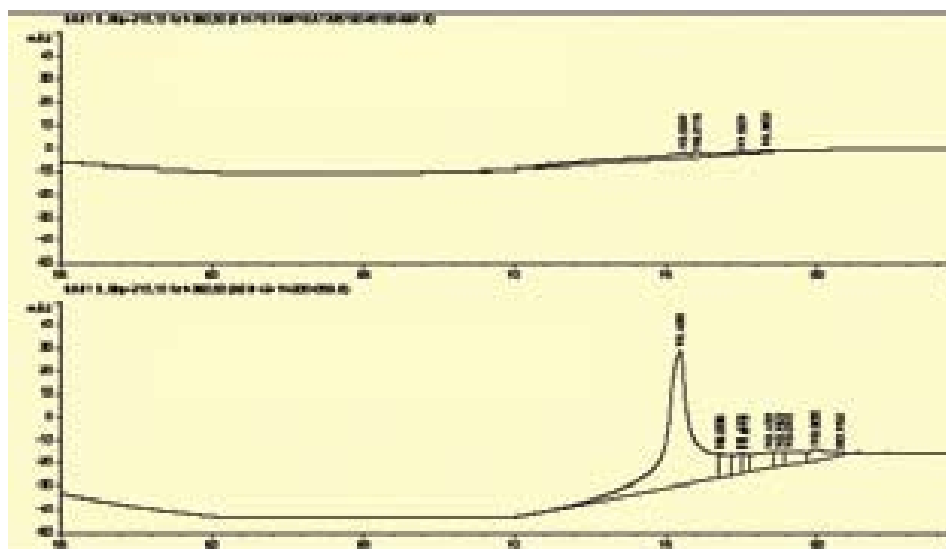


# How to make a good separation

- Test solubility with mobile phase
- Single analyte should started with Isocratic
- More than 2 analytes should started with Gradient
- 0.1% acid help ion pairing separation and enhance ionization step
- DO NOT use phosphate buffer, NaOH, HCl or other non-volatiles buffer in LC-MS system

# Acetonitrile Effect

- Very good used with water to facilitated the best separation
- Caused a baseline-shift
- Absorb at 210 nm



# Ion Pairing

- Common Ion Pairing  $\text{NH}_4^+$ ,  $\text{Na}^+$ ,  $\text{Cl}^-$ ,  $\text{H}^+$
- Increase separation efficiency but affect  $m/z$  in Mass Spectrometer
- Competitive ion species
- Can be illuminated in fragmentation processes

ppm, ppb, ppt or  $\mu\text{g/mL}$ ,  $\text{ng/mL}$ ,  $\text{pg/mL}$

- Density of matter is NOT EQUAL
- Effect Quantitation

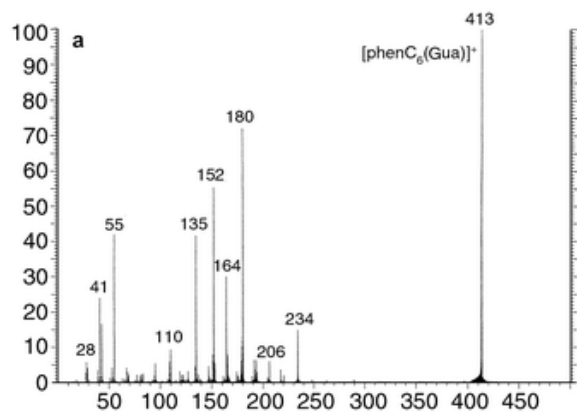
$$d=m/v$$

d=density (gram/mL)

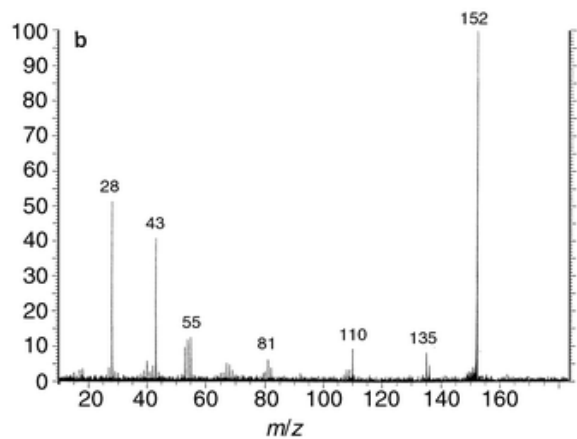
m=mass (gram)

v=volume (mL)

# What is a Mass Spectrometer?



“A device to measure the mass-to-charge ratio of individual molecules that have been converted to ions”

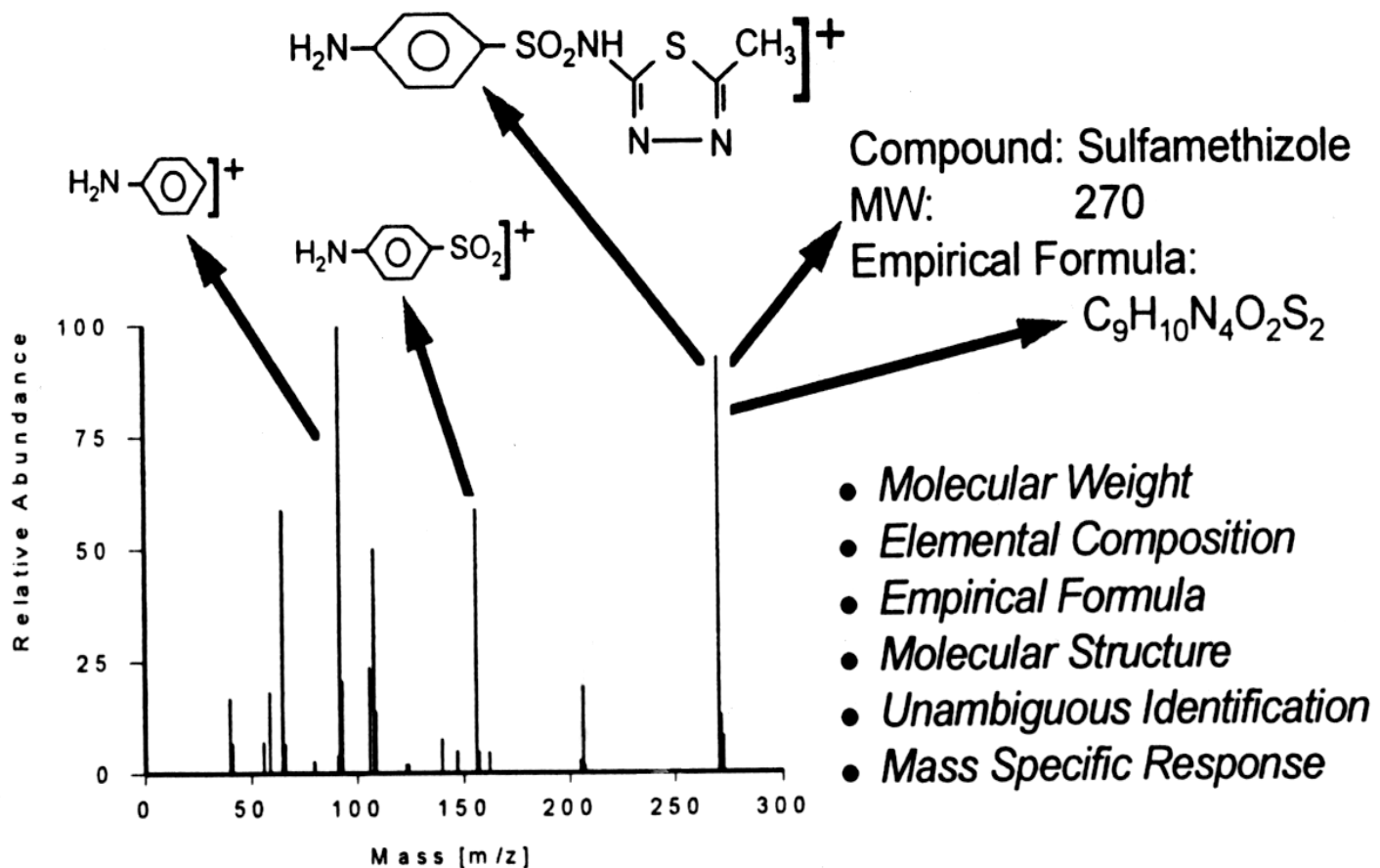


➤ Mass Spectrum: A plot of mass to charge ( $m/z$ ) vs. relative or absolute intensity

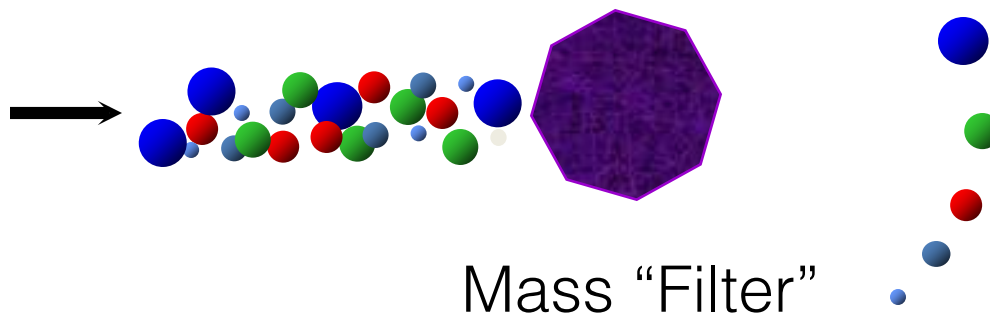
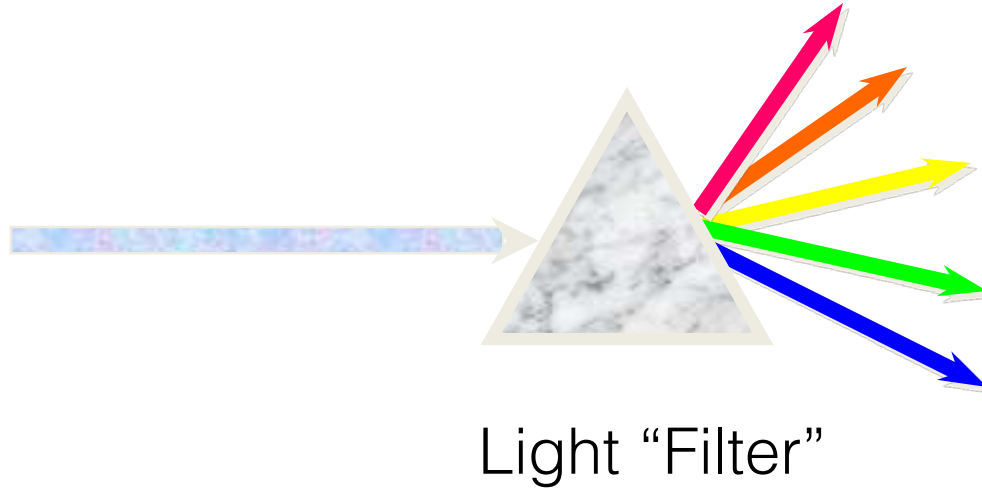
# Welcome to the world of Mass Spectrometer

- ▶ All mass analyzers determine the mass of an **ion**
- ▶ All mass analyzers determine the **mass-to-charge ratio**
- ▶ All mass analyzers measure **gas-phase ions**
- ▶ All mass analyzers must operate at very low pressure (a **vacuum**)

# Information Rich Data

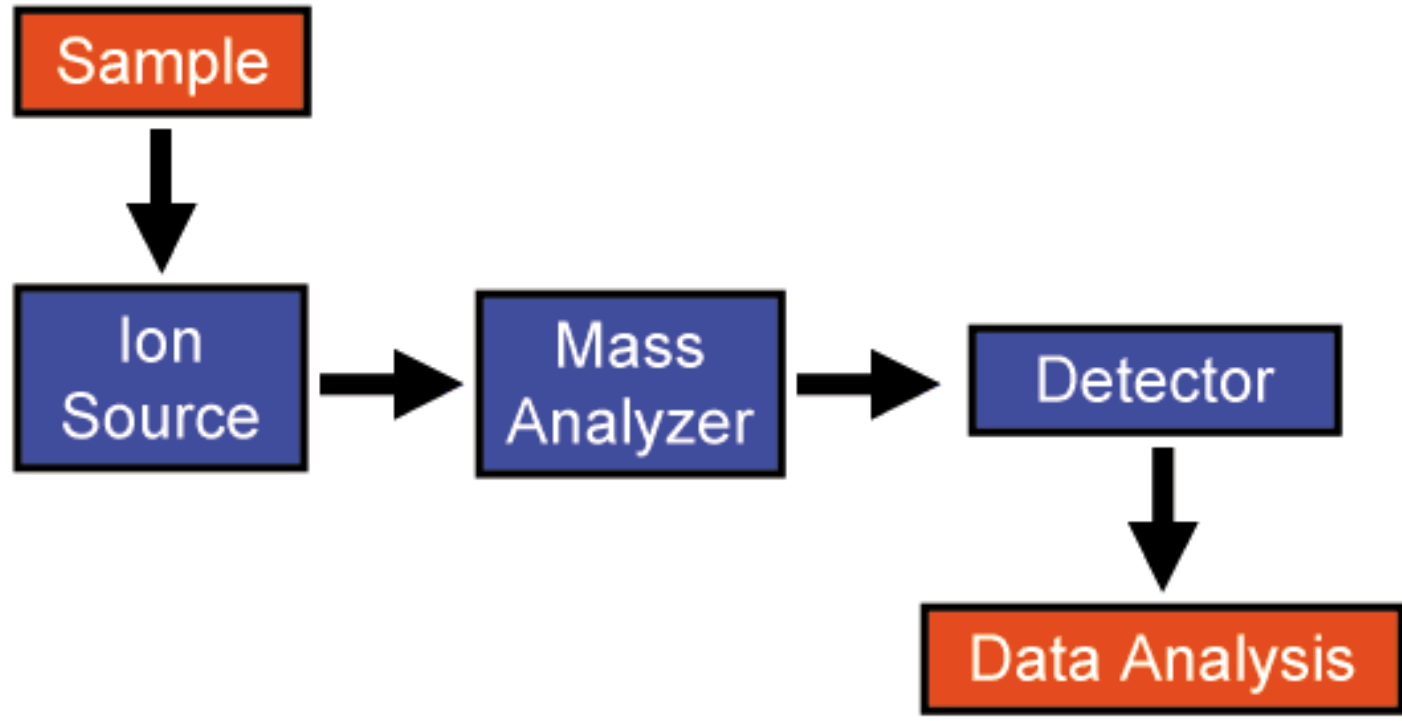


# Spectrum Formation

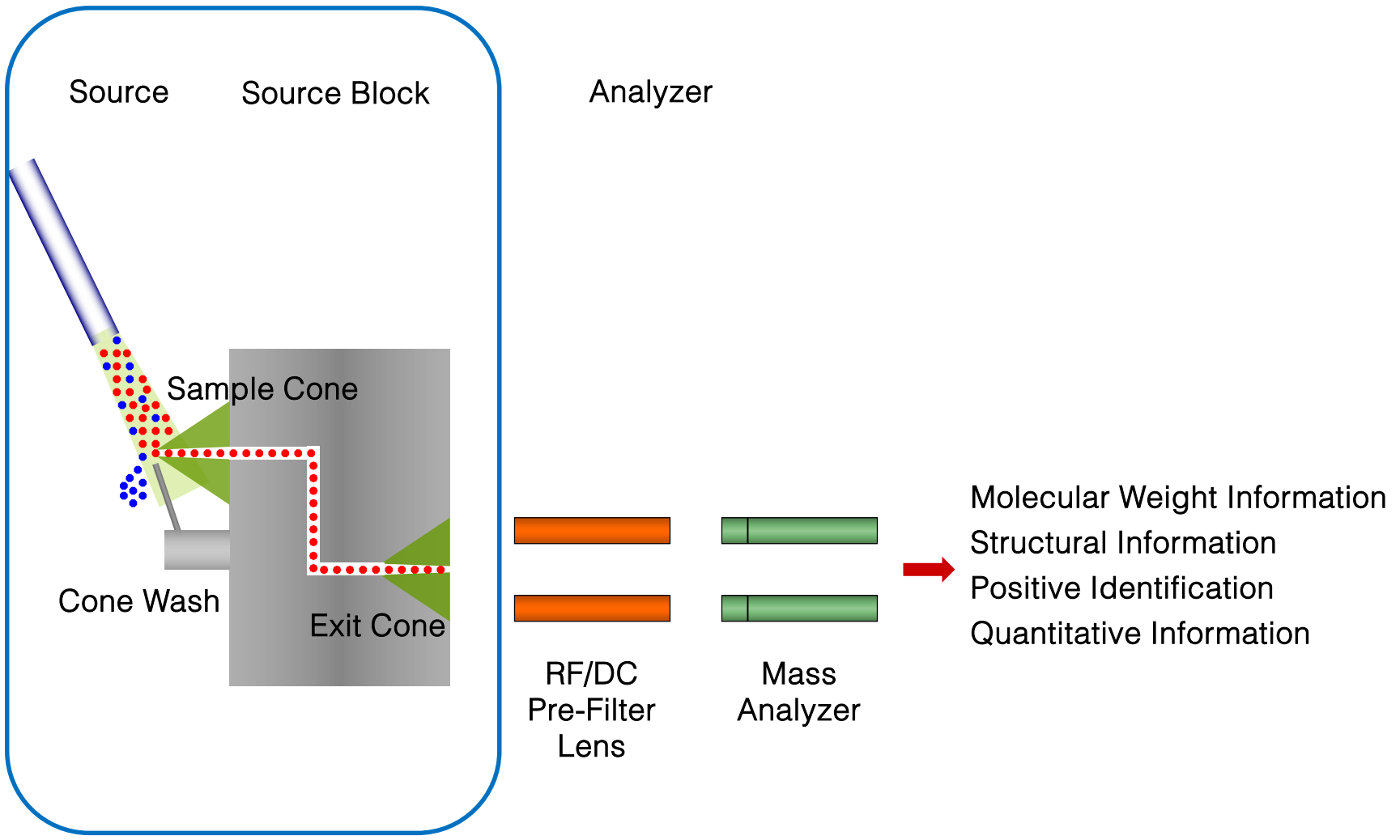




# Mass Spectrometer Diagram



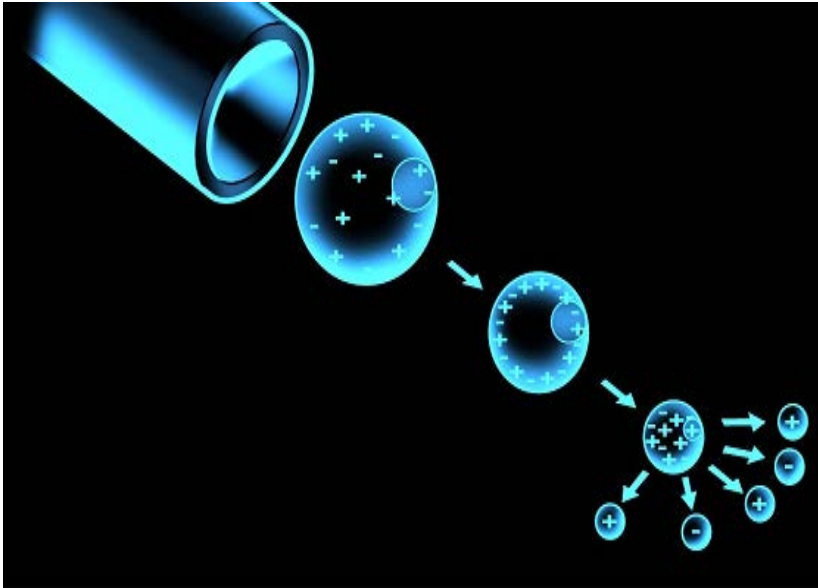
# Ion Source & Path



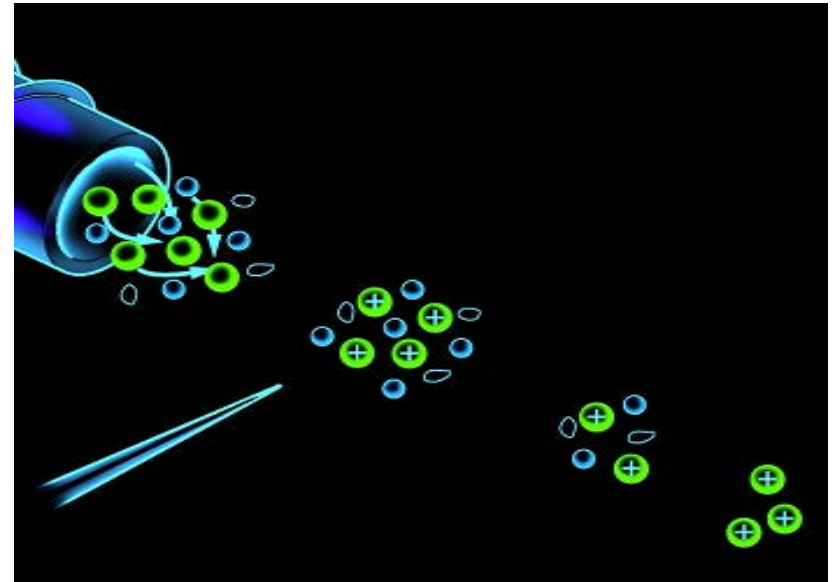
# Ion Source

- API; APCI, ESI, APPI
- Fast Atom Bombardment
- Matrix Assisted Laser Desorption/Ionization (MALDI)
- Ion Attachment
- Field Desorption
- Induced Couple Plasma
- Direct Analysis of Real Time (DART)

# Atmospheric Pressure Ionization



Electrospray Ionization



Atmospheric Pressure  
Chemical ionization

# Chemistry Considerations ESI or APCI

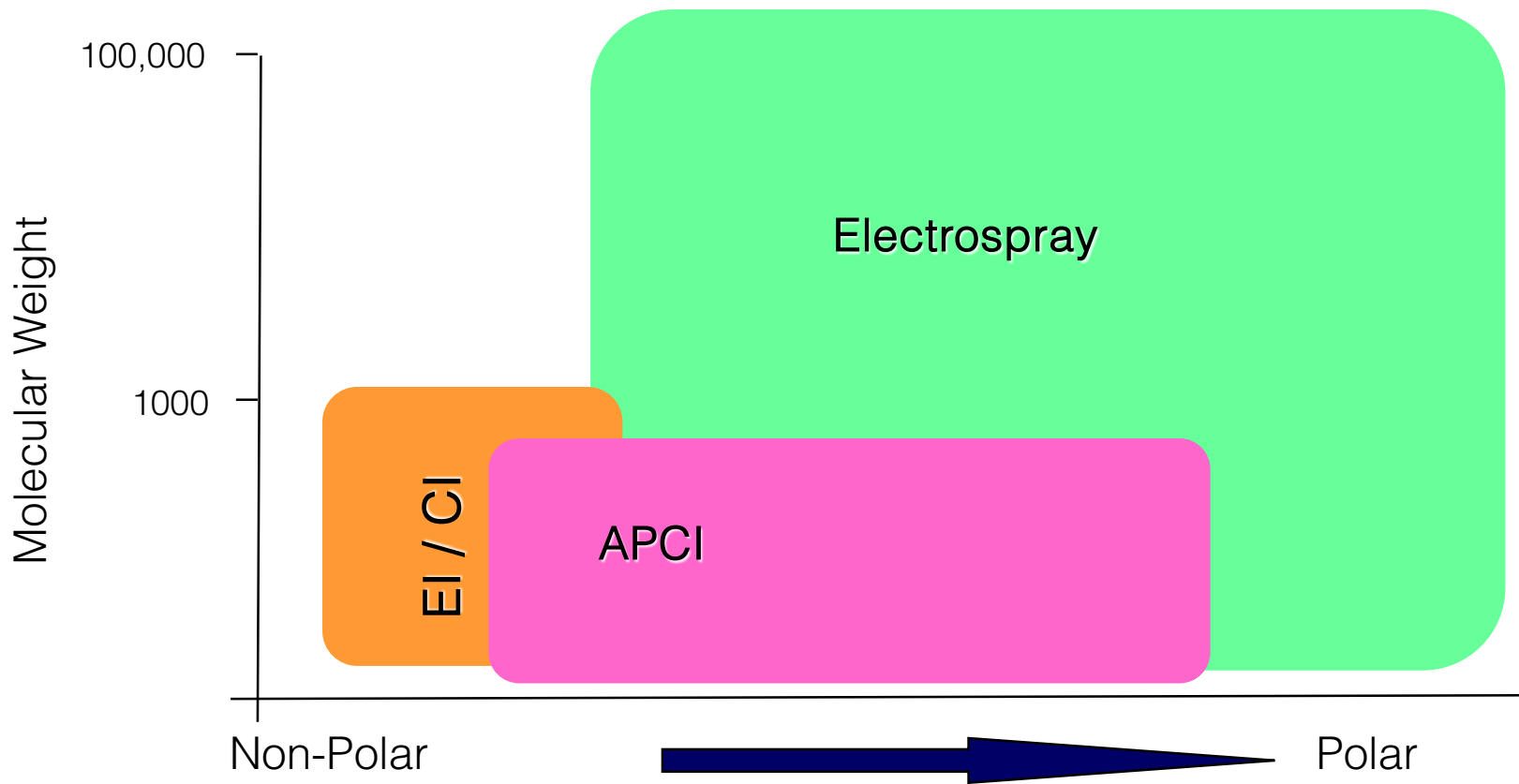
## ESI:

- Ions formed by solution chemistry
- Good for thermally labile analytes
- Good for polar analytes
- Good for large molecules (Proteins / Peptides)

## APCI:

- Ions formed by gas phase chemistry
- Good for volatile / thermally stable
- Good for non-polar analytes
- Good for small molecules (Steroids)

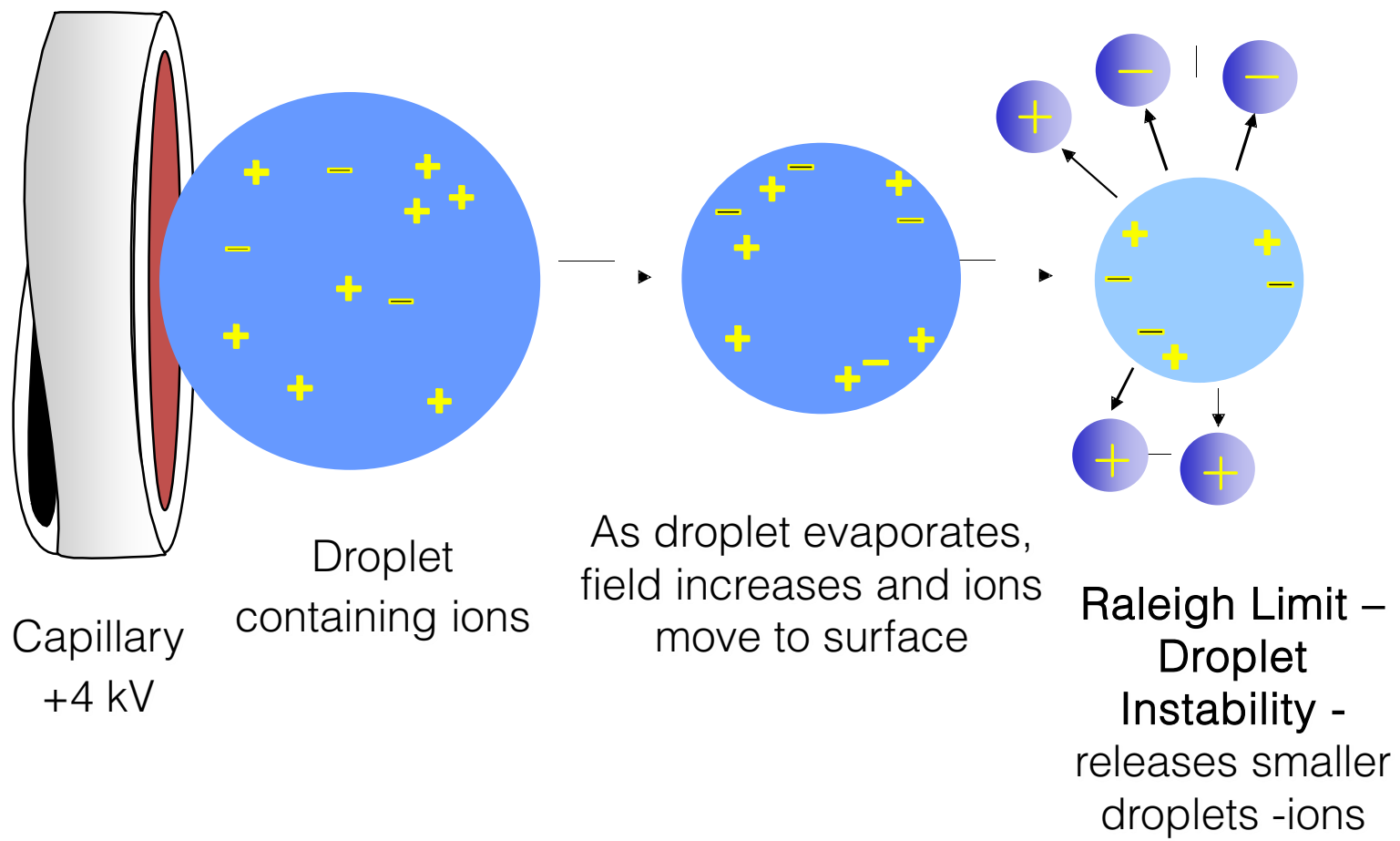
# Which Ionization Mode ?



# Step in Electrospray

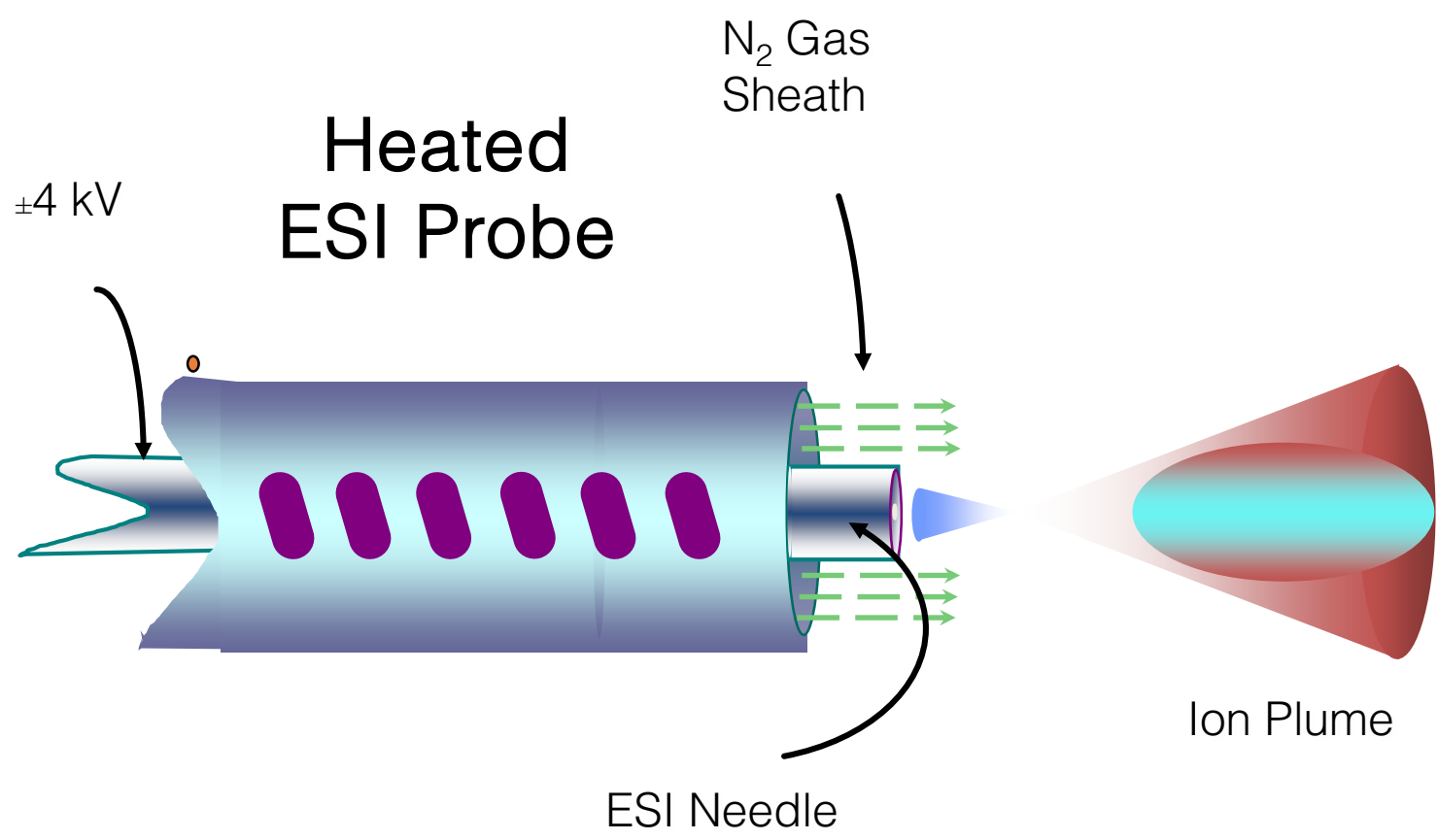
- Production of charged droplets from a capillary tip
  - Under influence of strong electric field
- Reduction of droplet size
  - Rapid solvent evaporation
  - Repeated coulombic explosions (fission)
- Transfer of ions from surface of small droplets to gas phase
  - No heat of evaporation

# Ion Evaporation Theory



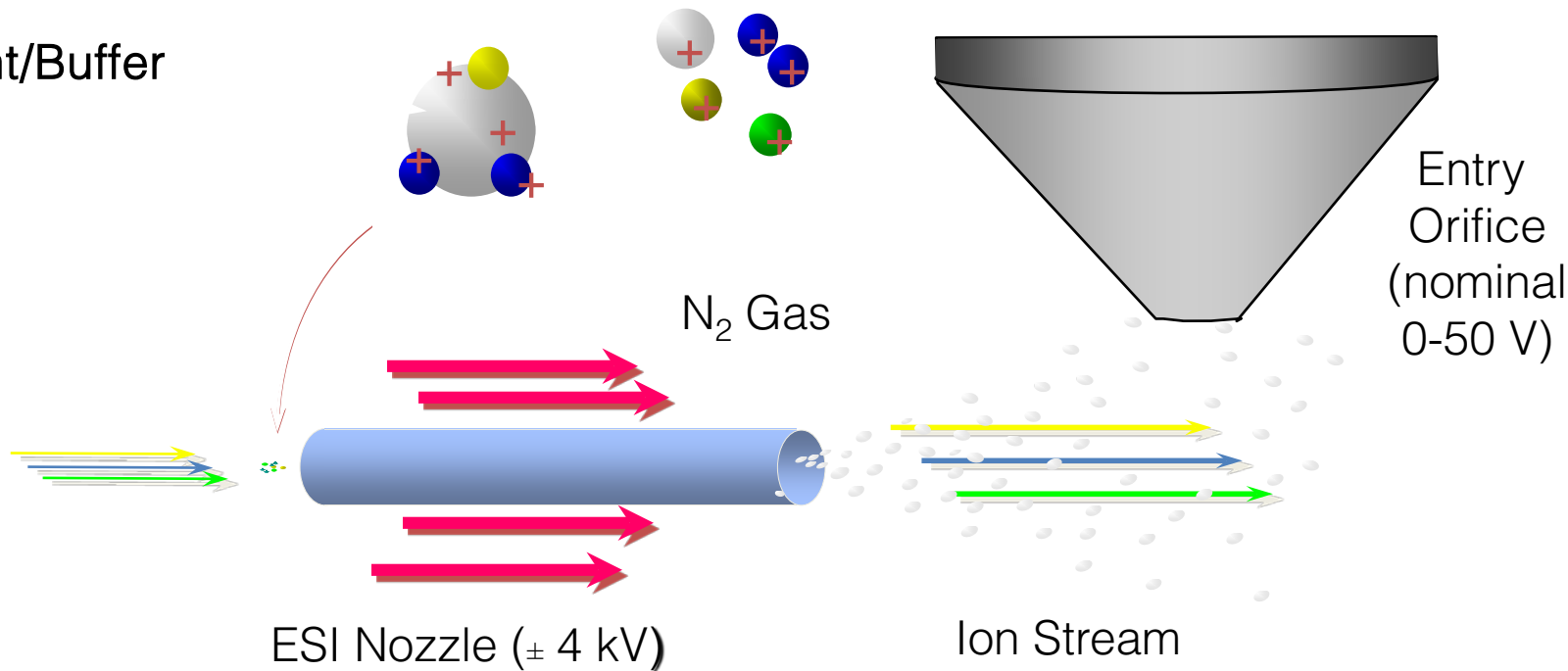


# Electrospray Nozzle Detail



# ESI/MS Ion Injection (Desolvating the Spray)

Solvent/Buffer

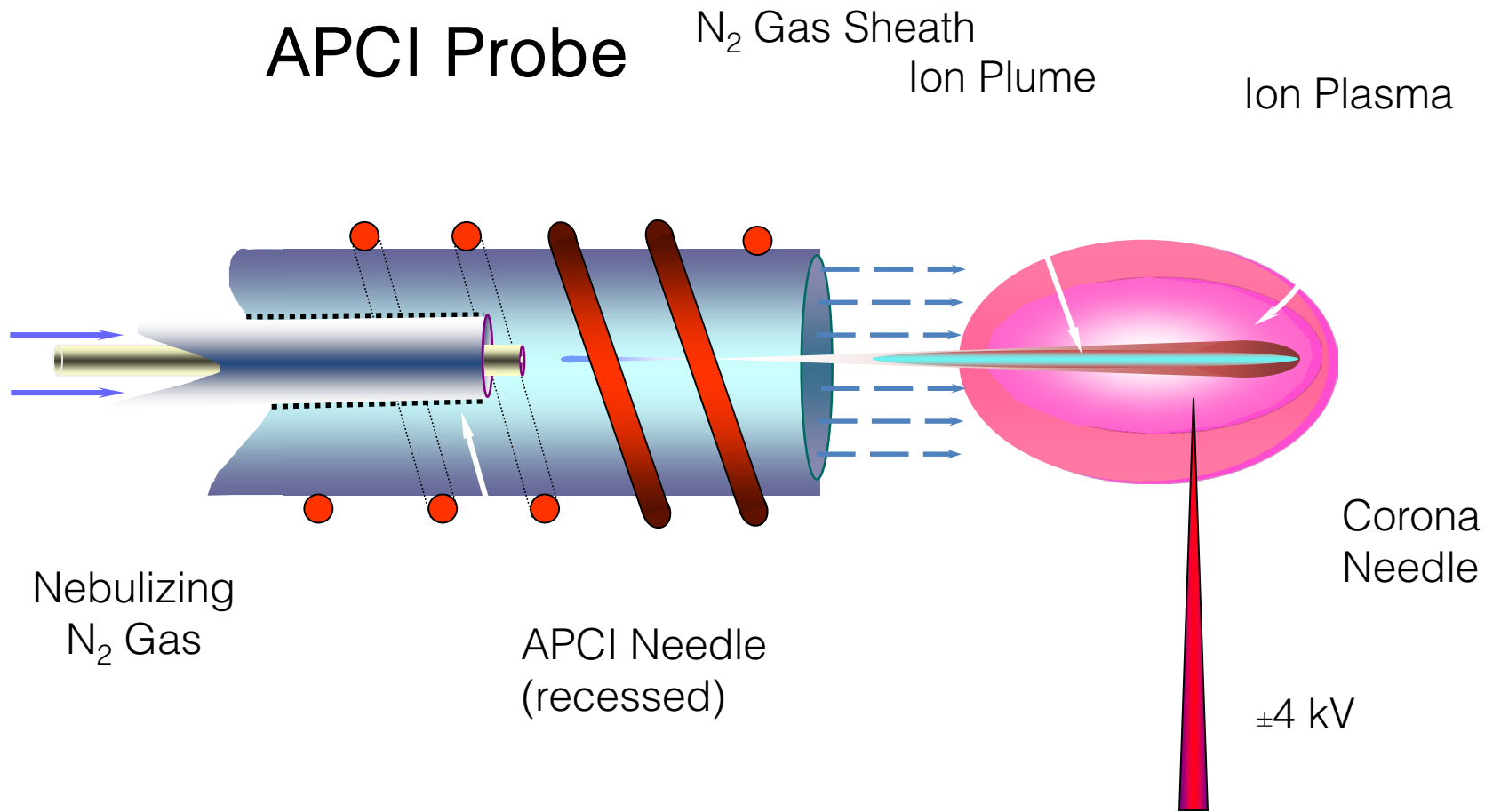


# Electrospray Ionization

- Low and high molecular weights
- Singly and multiply charged species
- Very soft ionization
- Mobile phase should have a polar component

# APCI Nozzle Detail

## Heated APCI Probe



# Atmospheric Pressure Chemical Ionization (APCI)

- Low molecular weight ( $<1000$ )
- Singly charged species only
- Thermal fragmentation may occur
- Mobile phase can be non-polar (normal phase)



How to Choose a  
proper Mass  
Spectrometer ?

# Typical Mass Accuracy

Type of MS	Mass accuracy	Utility for
Quadrupole	0.1 $\mu$	Identify
Traps	0.1 $\mu$	Identify
TOF	0.0001 $\mu$	Empirical formula/ composition
Sector	0.0001 $\mu$	Empirical formula/ composition
FT-MS	0.0001 $\mu$	Empirical formula/ composition

# Periodic Table of Elements

## Average Mass

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18
1 <b>H</b> Hydrogen 1.00794																	2 <b>He</b> Helium 4.002602
2 <b>Li</b> Lithium 6.941	4 <b>Be</b> Beryllium 9.012182																
3 <b>Na</b> Sodium 22.98976928	12 <b>Mg</b> Magnesium 24.3050																
4 <b>K</b> Potassium 39.0983	20 <b>Ca</b> Calcium 40.078	21 <b>Sc</b> Scandium 44.955912	22 <b>Ti</b> Titanium 47.887	23 <b>V</b> Vanadium 50.9415	24 <b>Cr</b> Chromium 51.9961	25 <b>Mn</b> Manganese 54.938045	26 <b>Fe</b> Iron 55.845	27 <b>Co</b> Cobalt 58.933195	28 <b>Ni</b> Nickel 58.6934	29 <b>Cu</b> Copper 63.546	30 <b>Zn</b> Zinc 65.38	31 <b>Ga</b> Gallium 69.723	32 <b>Ge</b> Germanium 72.64	33 <b>As</b> Arsenic 74.92160	34 <b>Se</b> Selenium 78.96	35 <b>Br</b> Bromine 79.904	36 <b>Kr</b> Krypton 83.798
5 <b>Rb</b> Rubidium 85.4678	38 <b>Sr</b> Strontium 87.62	39 <b>Y</b> Yttrium 88.90585	40 <b>Zr</b> Zirconium 91.224	41 <b>Nb</b> Niobium 92.90638	42 <b>Mo</b> Molybdenum 95.96	43 <b>Tc</b> Technetium (97.9072)	44 <b>Ru</b> Ruthenium 101.07	45 <b>Rh</b> Rhodium 102.90550	46 <b>Pd</b> Palladium 106.42	47 <b>Ag</b> Silver 107.8682	48 <b>Cd</b> Cadmium 112.411	49 <b>In</b> Indium 114.818	50 <b>Sn</b> Tin 118.710	51 <b>Sb</b> Antimony 121.760	52 <b>Te</b> Tellurium 127.60	53 <b>I</b> Iodine 126.90447	54 <b>Xe</b> Xenon 131.293
6 <b>Cs</b> Caesium 132.9054519	56 <b>Ba</b> Barium 137.327	57-71	72 <b>Hf</b> Hafnium 178.49	73 <b>Ta</b> Tantalum 180.94788	74 <b>W</b> Tungsten 183.84	75 <b>Re</b> Rhenium 186.207	76 <b>Os</b> Osmium 190.23	77 <b>Ir</b> Iridium 192.217	78 <b>Pt</b> Platinum 195.084	79 <b>Au</b> Gold 196.966569	80 <b>Hg</b> Mercury 200.59	81 <b>Tl</b> Thallium 204.3833	82 <b>Pb</b> Lead 207.2	83 <b>Bi</b> Bismuth 208.98040	84 <b>Po</b> Polonium (209)	85 <b>At</b> Astatine (209)	86 <b>Rn</b> Radon (222)
7 <b>Fr</b> Francium (223)	88 <b>Ra</b> Radium (226)	89-103	104 <b>Rf</b> Rutherfordium (261)	105 <b>Db</b> Dubnium (262)	106 <b>Sg</b> Seaborgium (266)	107 <b>Bh</b> Bohrium (264)	108 <b>Hs</b> Hassium (277)	109 <b>Mt</b> Meitnerium (268)	110 <b>Ds</b> Darmstadtium (271)	111 <b>Rg</b> Roentgenium (272)	112 <b>Uub</b> Ununbium (285)	113 <b>Uut</b> Ununtrium (284)	114 <b>Uuq</b> Ununquadium (289)	115 <b>Uup</b> Ununpentium (288)	116 <b>Uuh</b> Ununhexium (292)	117 <b>Uus</b> Ununseptium	118 <b>Uuo</b> Ununoctium (294)

- C** Solid
- Hg** Liquid
- H** Gas
- Rf** Unknown

Metals					Nonmetals		
Alkali metals	Alkaline earth metals	Lanthanoids	Transition metals	Poor metals	Other nonmetals	Noble gases	
		Actinoids					

For elements with no stable isotopes, the mass number of the isotope with the longest half-life is in parentheses.

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57 <b>La</b> Lanthanum 138.90547	58 <b>Ce</b> Cerium 140.116	59 <b>Pr</b> Praseodymium 140.90765	60 <b>Nd</b> Neodymium 144.242	61 <b>Pm</b> Promethium (145)	62 <b>Sm</b> Samarium 150.36	63 <b>Eu</b> Europium 151.964	64 <b>Gd</b> Gadolinium 157.25	65 <b>Tb</b> Terbium 158.92535	66 <b>Dy</b> Dysprosium 162.500	67 <b>Ho</b> Holmium 164.93032	68 <b>Er</b> Erbium 167.259	69 <b>Tm</b> Thulium 168.93421	70 <b>Yb</b> Ytterbium 173.054	71 <b>Lu</b> Lutetium 174.968
89 <b>Ac</b> Actinium (227)	90 <b>Th</b> Thorium 232.03806	91 <b>Pa</b> Protactinium 231.03588	92 <b>U</b> Uranium 238.02891	93 <b>Np</b> Neptunium (237)	94 <b>Pu</b> Plutonium (244)	95 <b>Am</b> Americium (243)	96 <b>Cm</b> Curium (247)	97 <b>Bk</b> Berkelium (247)	98 <b>Cf</b> Californium (251)	99 <b>Es</b> Einsteinium (252)	100 <b>Fm</b> Fermium (257)	101 <b>Md</b> Mendelevium (258)	102 <b>No</b> Nobelium (259)	103 <b>Lr</b> Lawrencium (260)



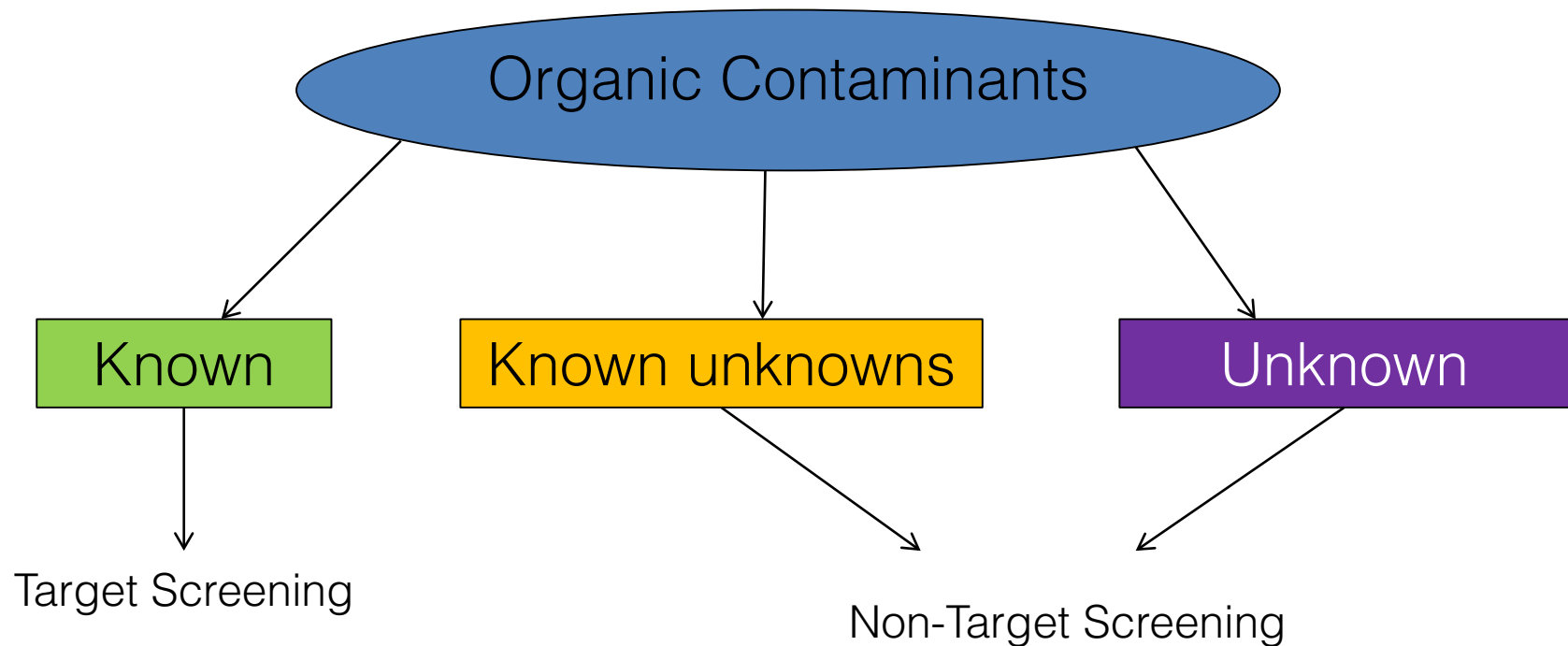
# How's About Mass Accuracy

- Average Mass = summing the [average atomic masses](#) of the constituent elements, H<sub>2</sub>O;  $1.00794 + 1.00794 + 15.9994 = 18.01528$ .
- Exact Mass = summing the masses of the individual isotopes of the molecule, H<sub>2</sub>O;  $1.0078 + 1.0078 + 15.9994 = 18.0106$ .

## *The Others Stories;*

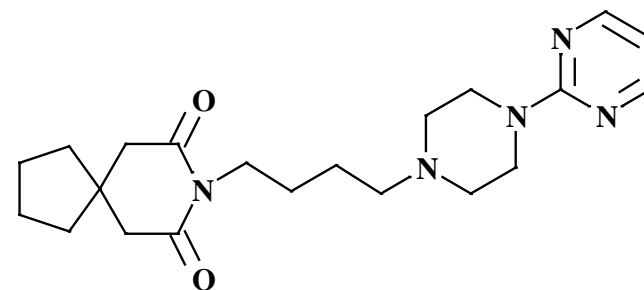
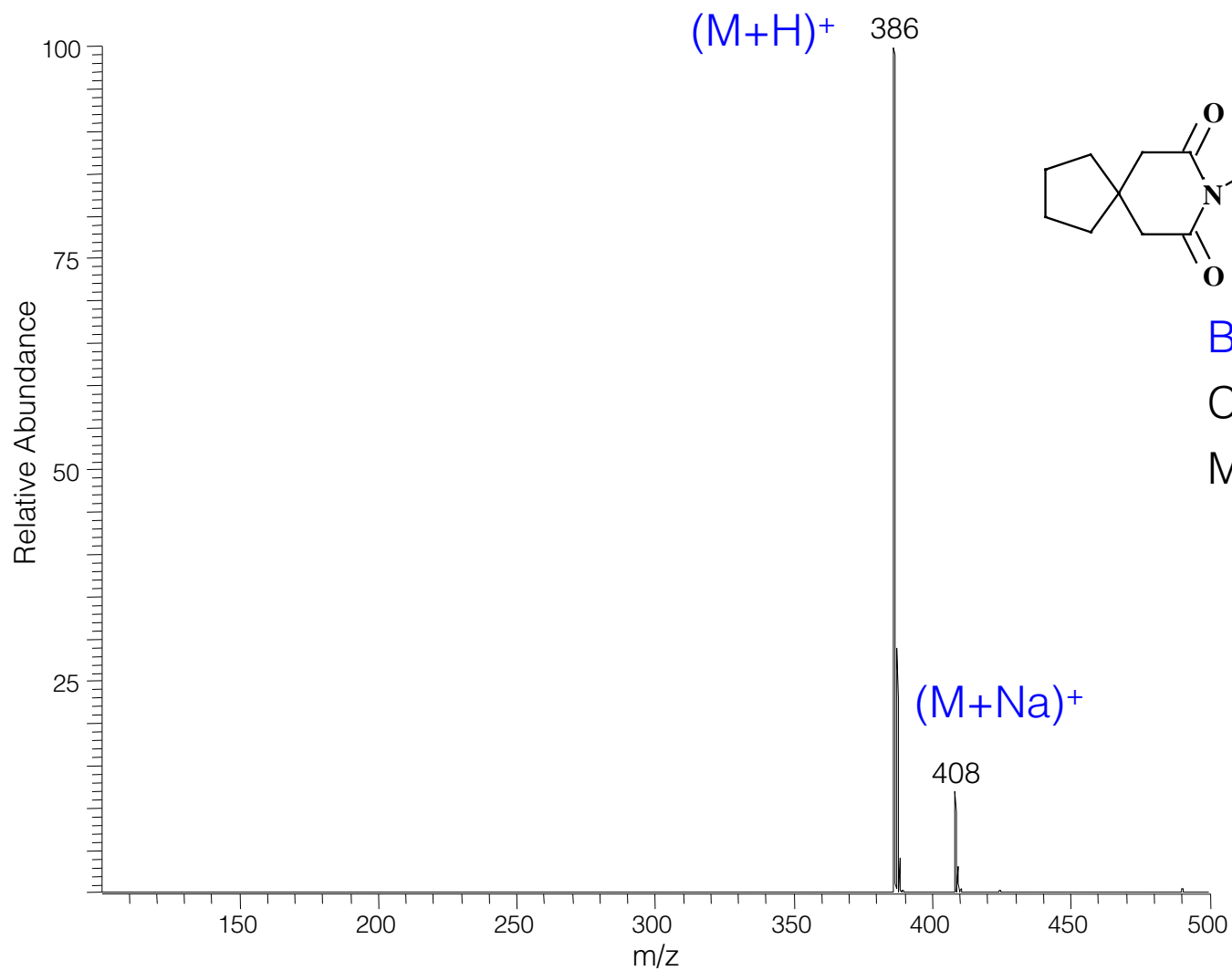
- Isotopomer (Isotopic Isomer) = same type of isotope but difference in position, CH<sub>3</sub>CHDCH<sub>3</sub> vs CH<sub>3</sub>CH<sub>2</sub>CH<sub>2</sub>D
- Isotopologues = difference in isotope in the molecules, H<sub>2</sub>O  
HOD
- Monoisotopic = sum of masses in molecule. Using of most abundance or stable isotope.

# Strategies for screening



Rapid and sensitive screening methods able to assign positive hits undoubtedly to particular organic compounds

# Full-Scan MS of Buspirone

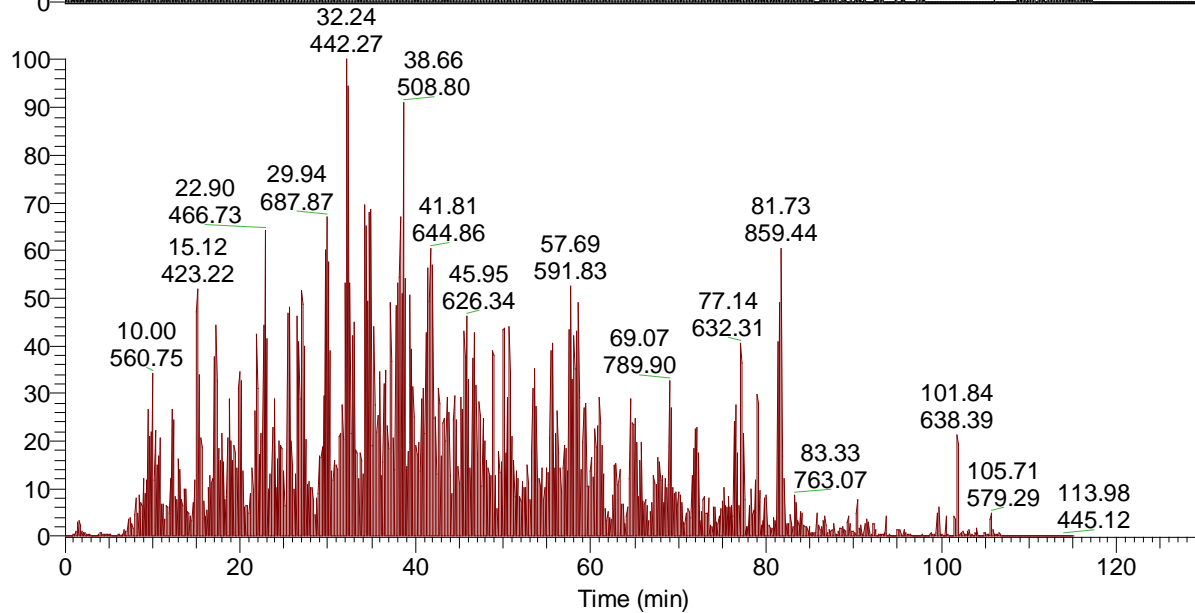
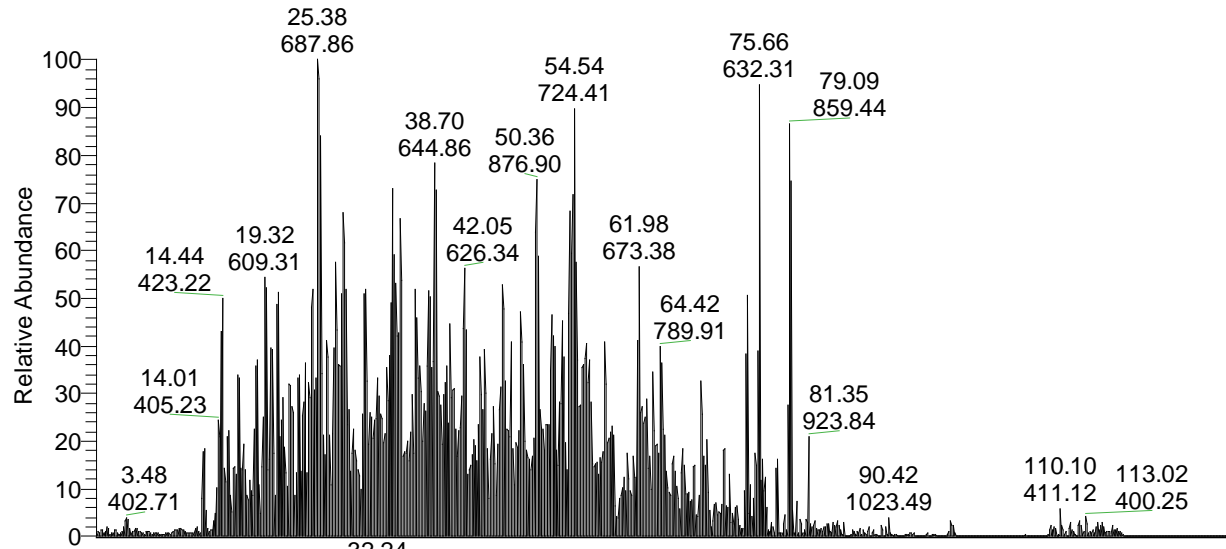


Buspirone

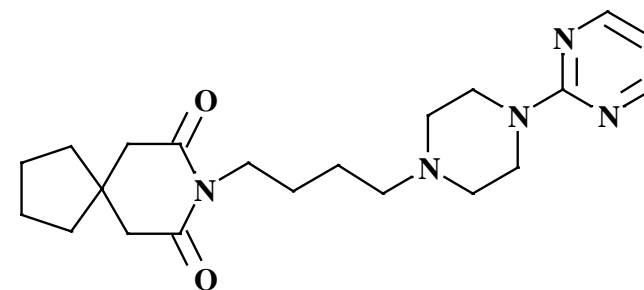
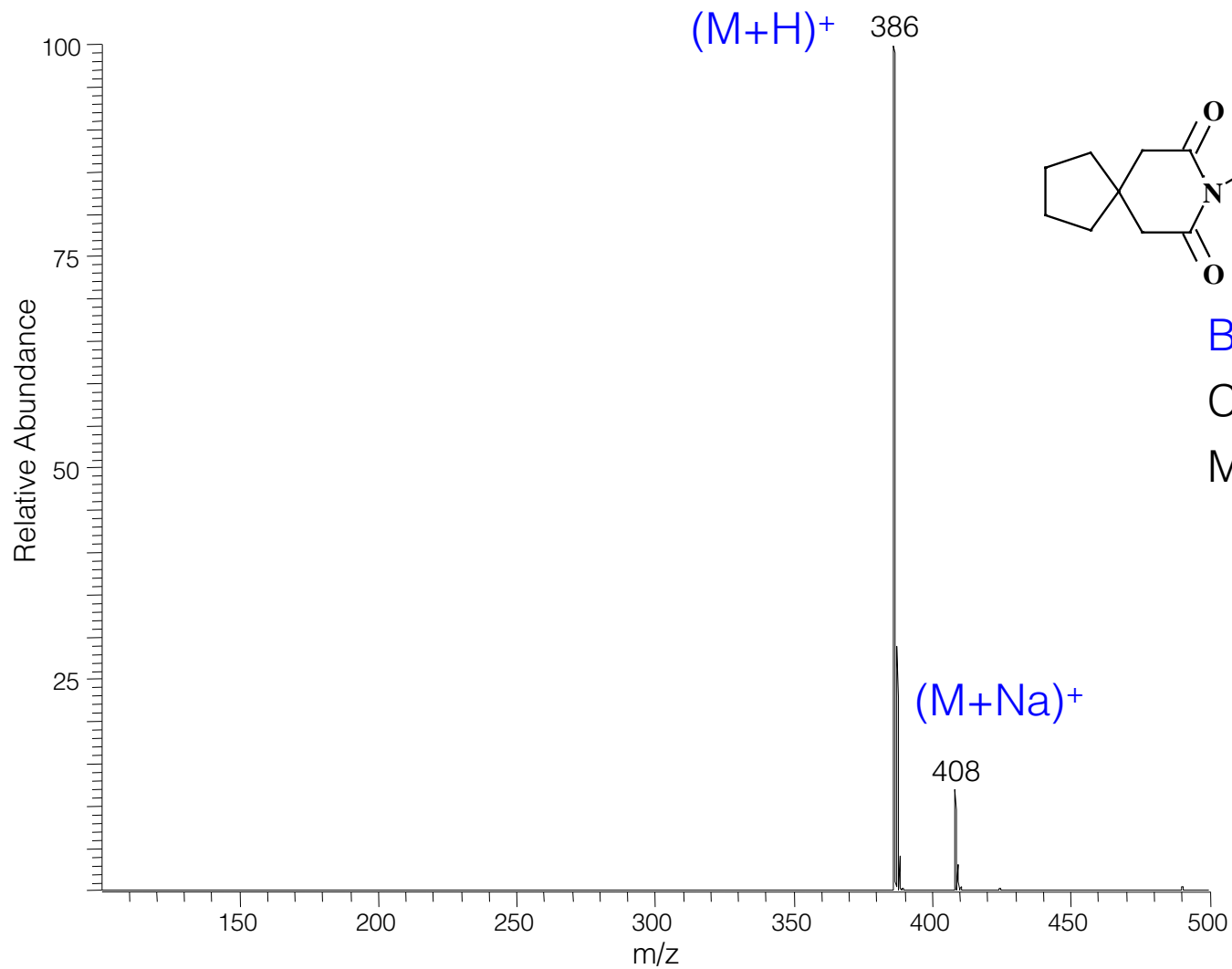
$C_{21}H_{31}N_5O_2$

MW = 385

# Real-life Full-Scan



# SIM MS of Buspirone

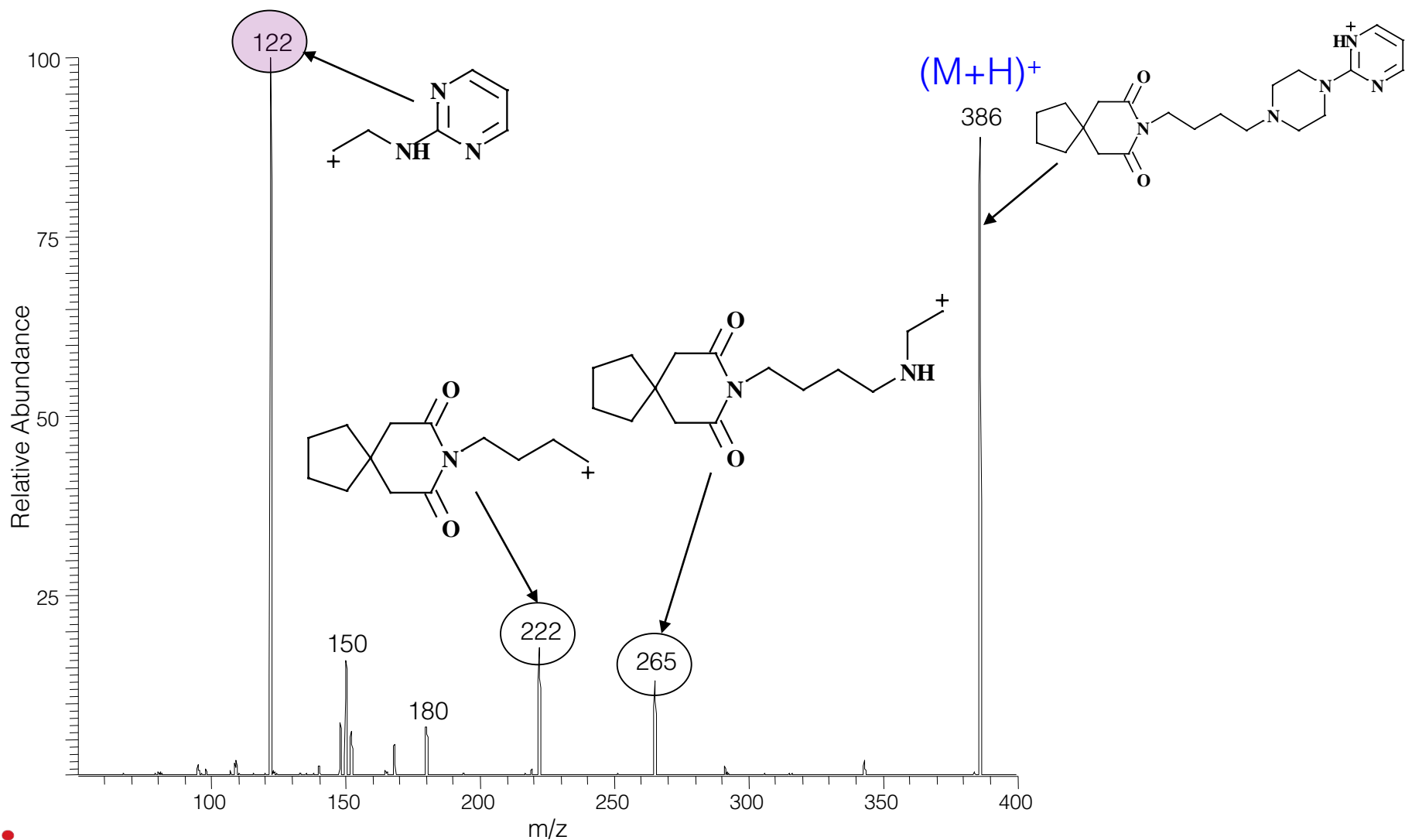


Buspirone

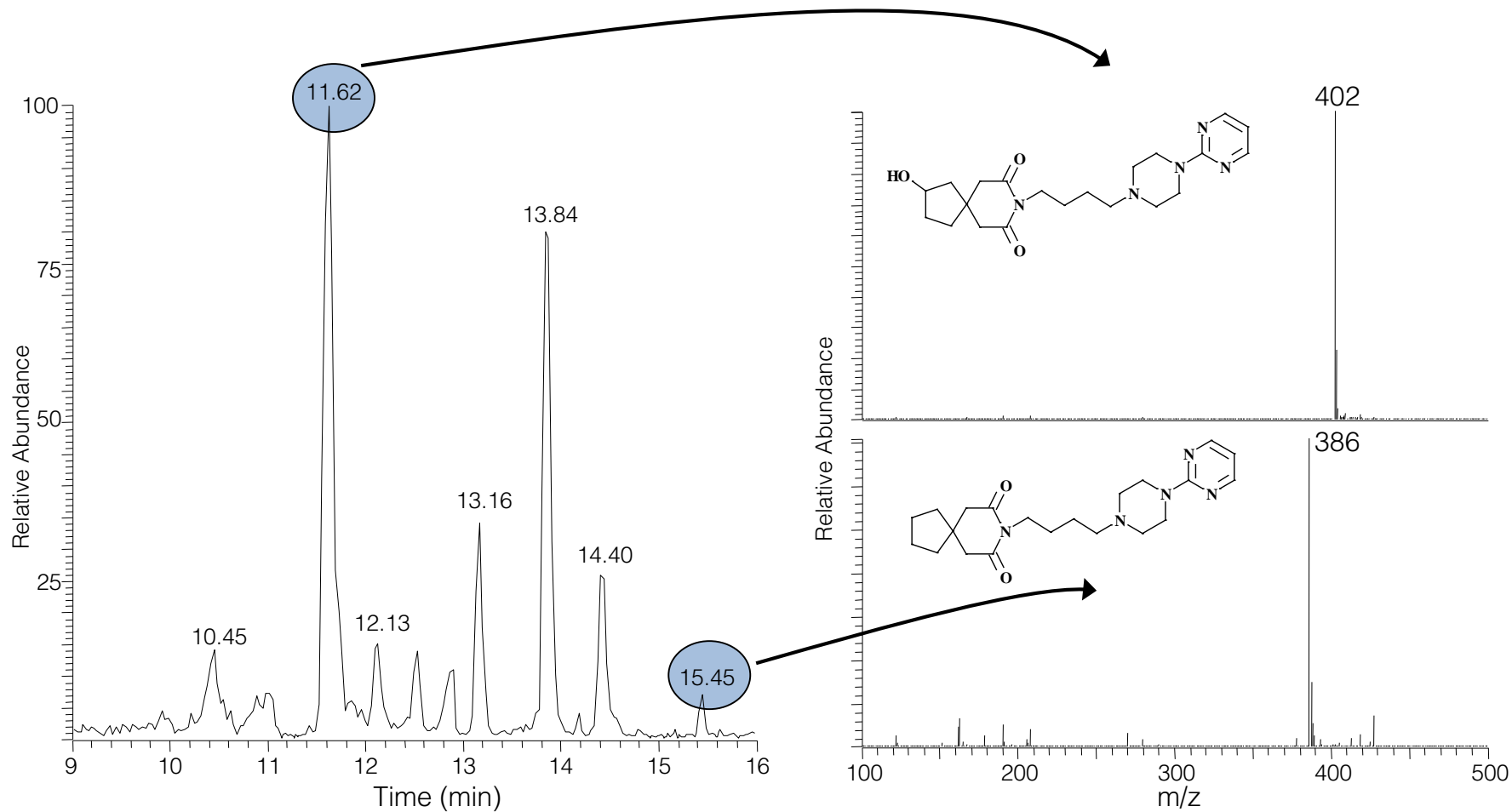
$C_{21}H_{31}N_5O_2$

MW = 385

# Product Ion Spectrum of Buspirone

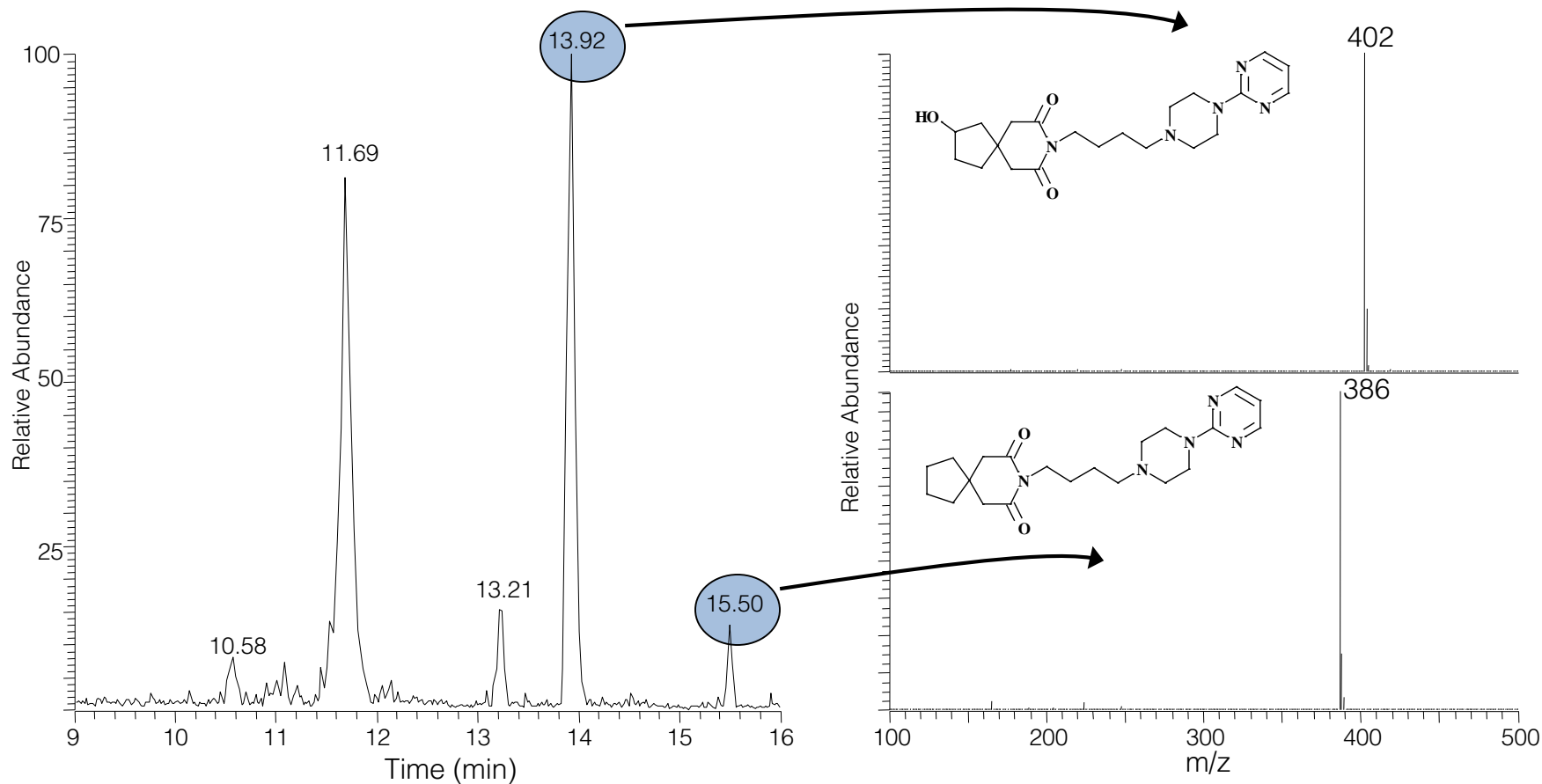


# Precursor Ion Scan Mode for Buspirone Metabolites



Precursor Ion Scan: Q3 set to m/z 122

# Neutral Loss Scan of Buspirone Metabolites



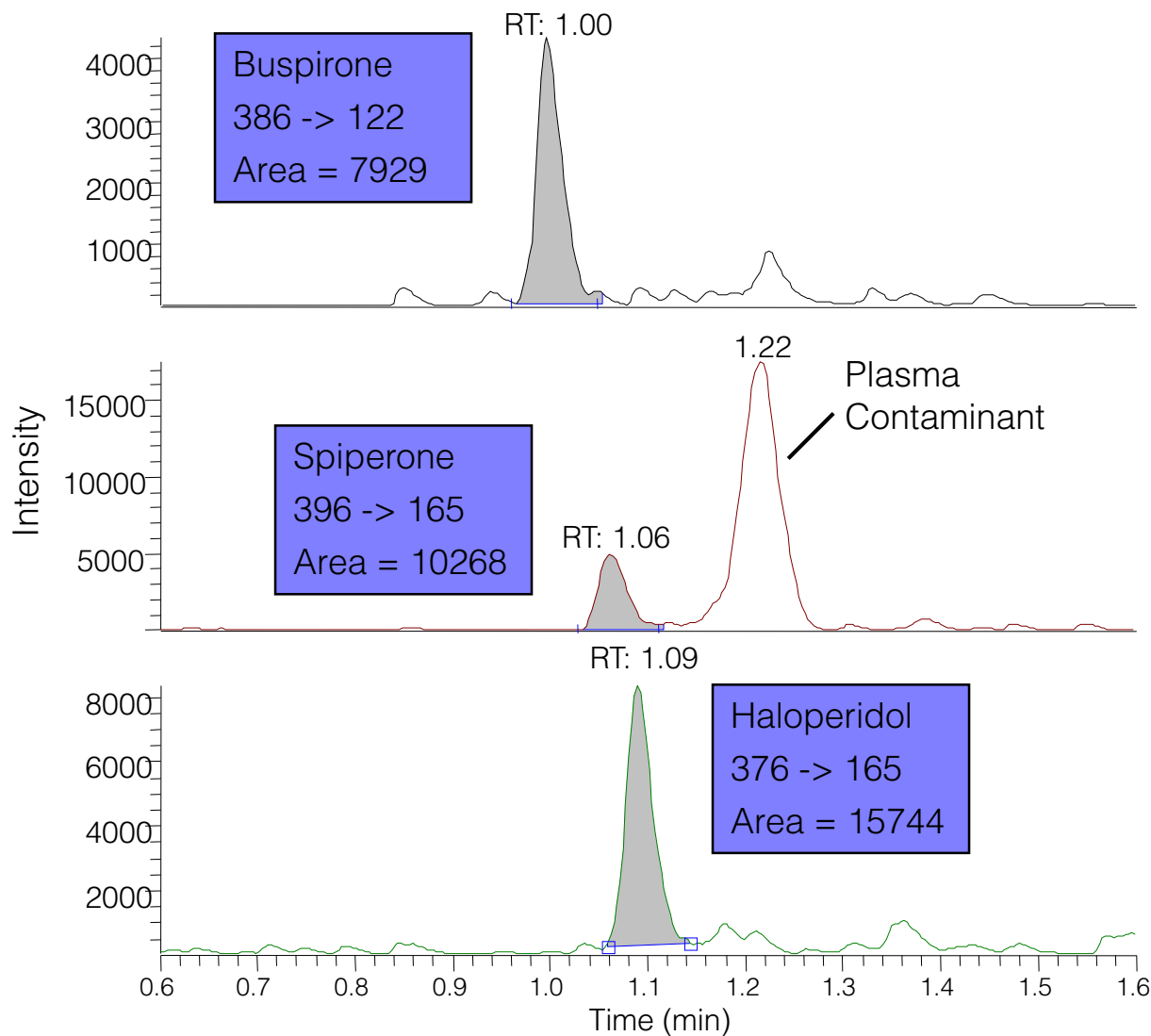
Neutral Loss Scan: Q1/Q3 difference set to 121 Da



# Scan Modes

Scan Mode	Purpose
Full-Scan	MW Info.
SIM	Quantitation
Product	Structural Info.
SRM	Targeted Quantitation
Neutral Loss	Analyte Screening
Precursor	Analyte Screening

# Quantitation Using SRM Mode on the TSQ Quantum



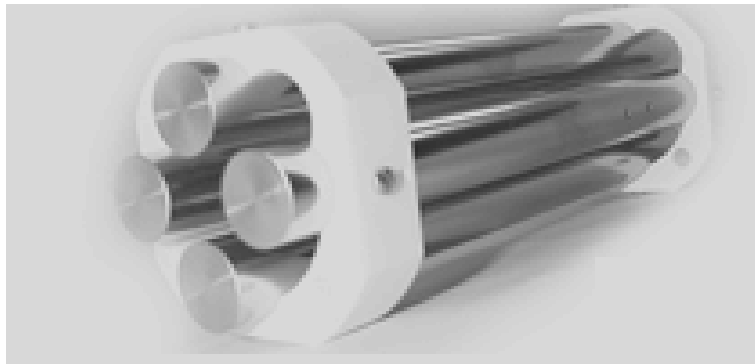
- LC/ESI-MS/MS, SRM Mode
- 10 pg/mL Buspirone, Spiperone & Haloperidol in Bovine Plasma
- 100 fg on column
- Ballistic Gradient Method @ 1 mL/min (no split)
- Total acquisition time = 1.6 min

# Targeted Screening Ideal

- High isolation power for higher discrimination
- High precision for accurate mass identification
- Low detection limit
- High mass stability for a long lasting mass calibration
- Highly selective ion monitoring (H-SRM)
- Low dwell time and no cross-talk for no-false interpretation
- Fast polarity switching

# Mass Analyzer - Quadrupole

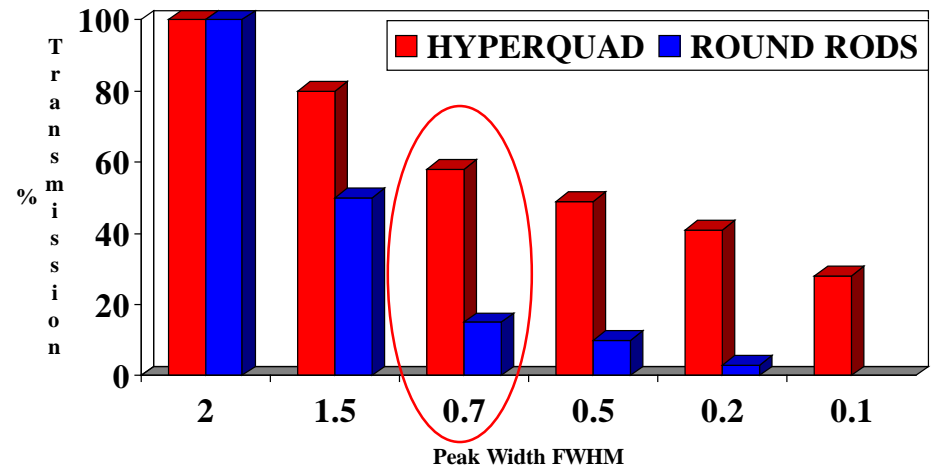
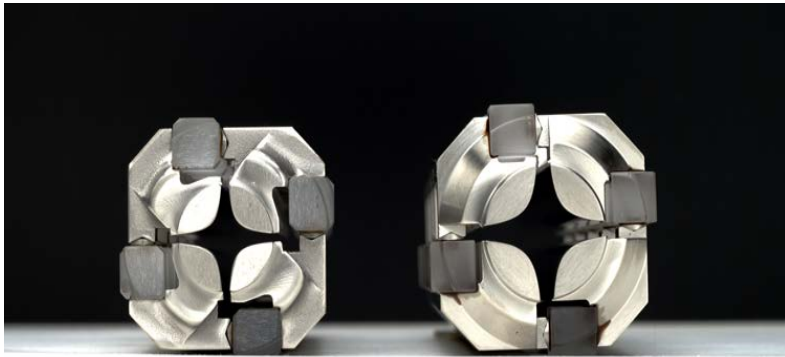
- Quadrupole, comprise of
- Quadrupole pre-filter; Hyperbolic, Round, Square



Controlled by RF and DC voltage !  
Ion transmitting @ 1 m/z in given  
time

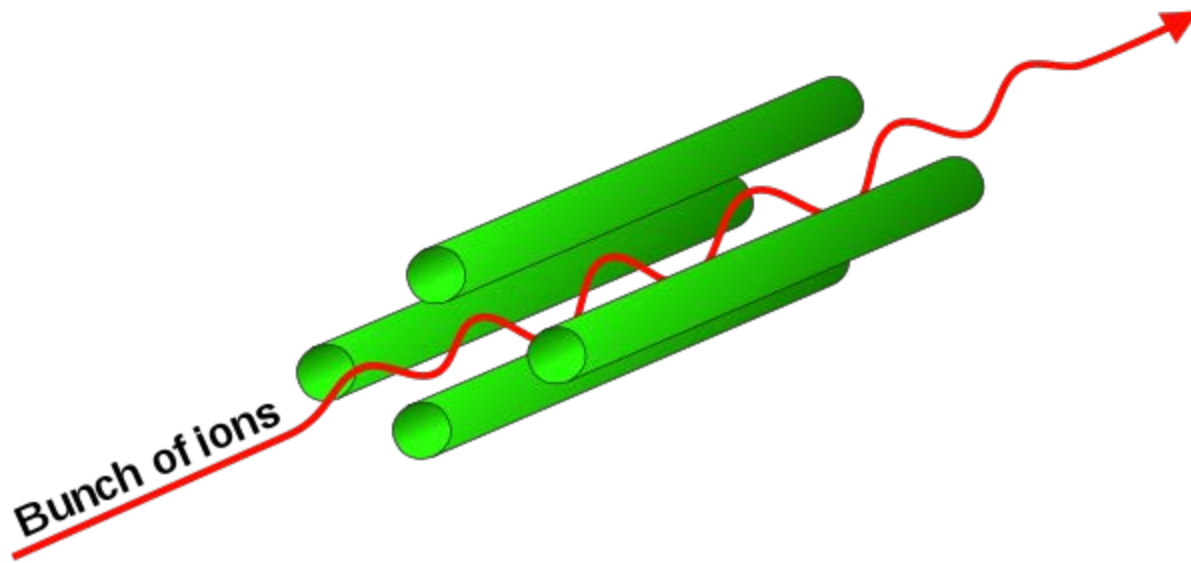
# Advantages Of HyperQuads

- ▶ Forms Pure Quadrupolar Fields (improves peak shape)
- ▶ Reduces Fringing Effects (sensitivity enhancement)
- ▶ Significantly Improves Resolution
- ▶ Improves Transmission (sensitivity enhancement)



# Type of Quadrupole Instruments

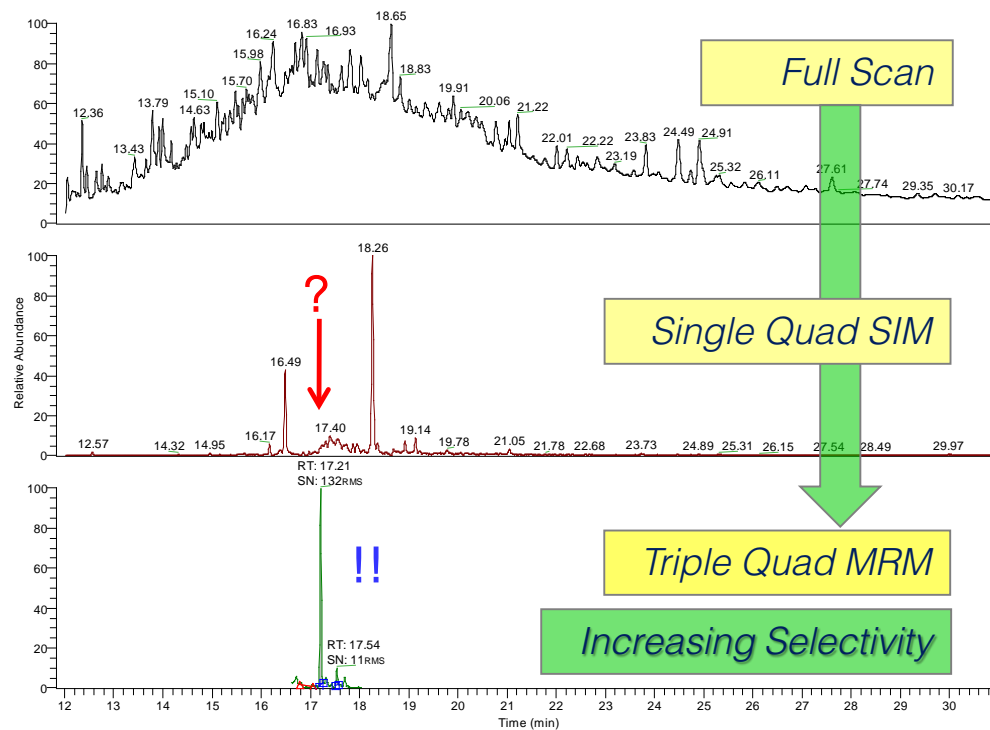
- Single Quadrupole
- Triple Quadrupole



# Applications Driver

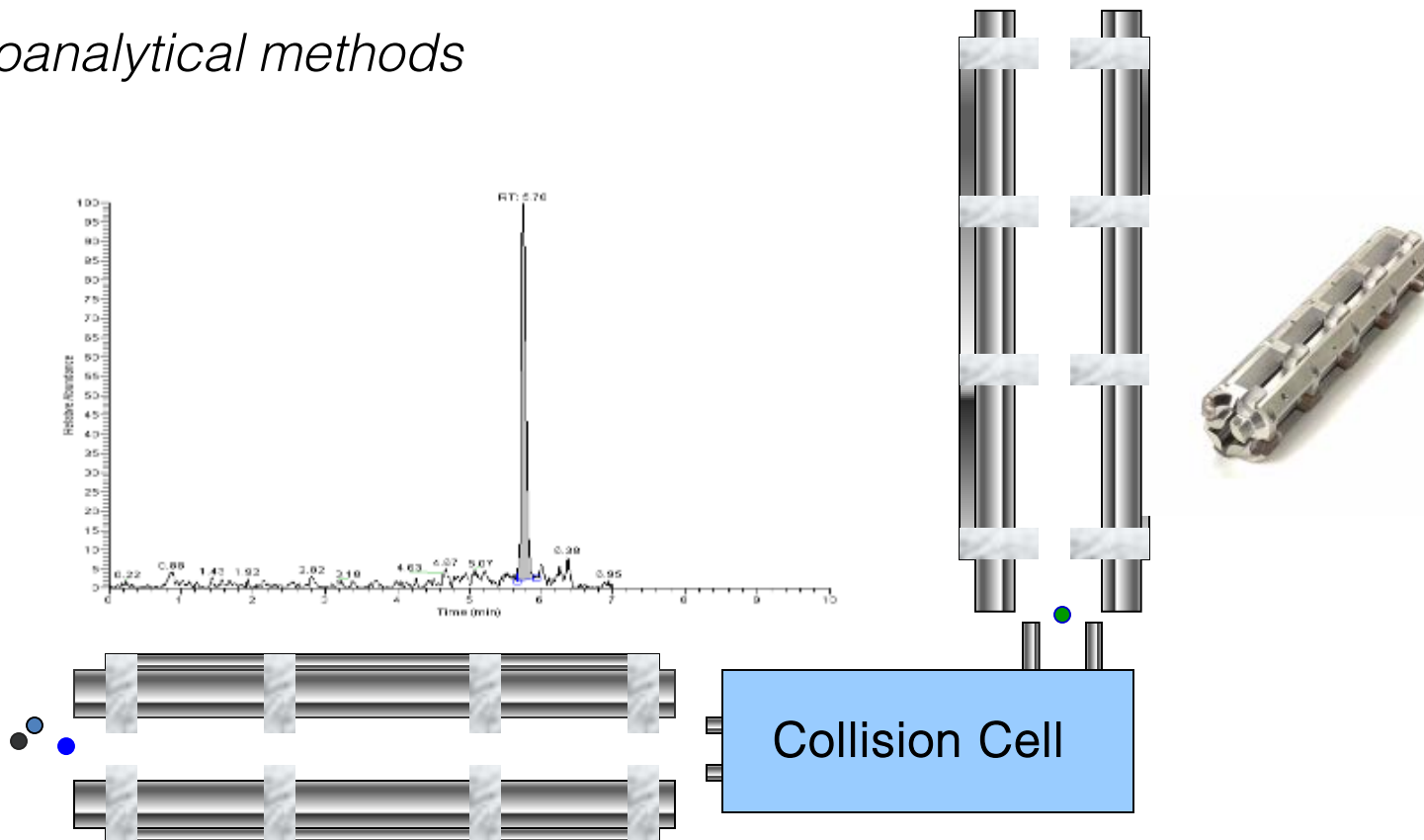
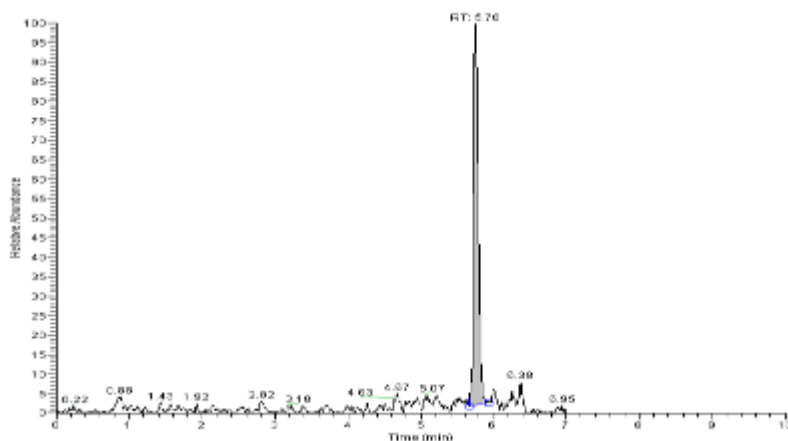
- High Sensitivity in Matrix Samples
  - Lower levels for increased number of analytes
  - Shorten expensive sample prep
  - Small sample volumes with reduced clean-up

## Food Safety Anabolic steroids in meat production



# H-SRM Operation – More Method Robustness

✓ *Robust bioanalytical methods*

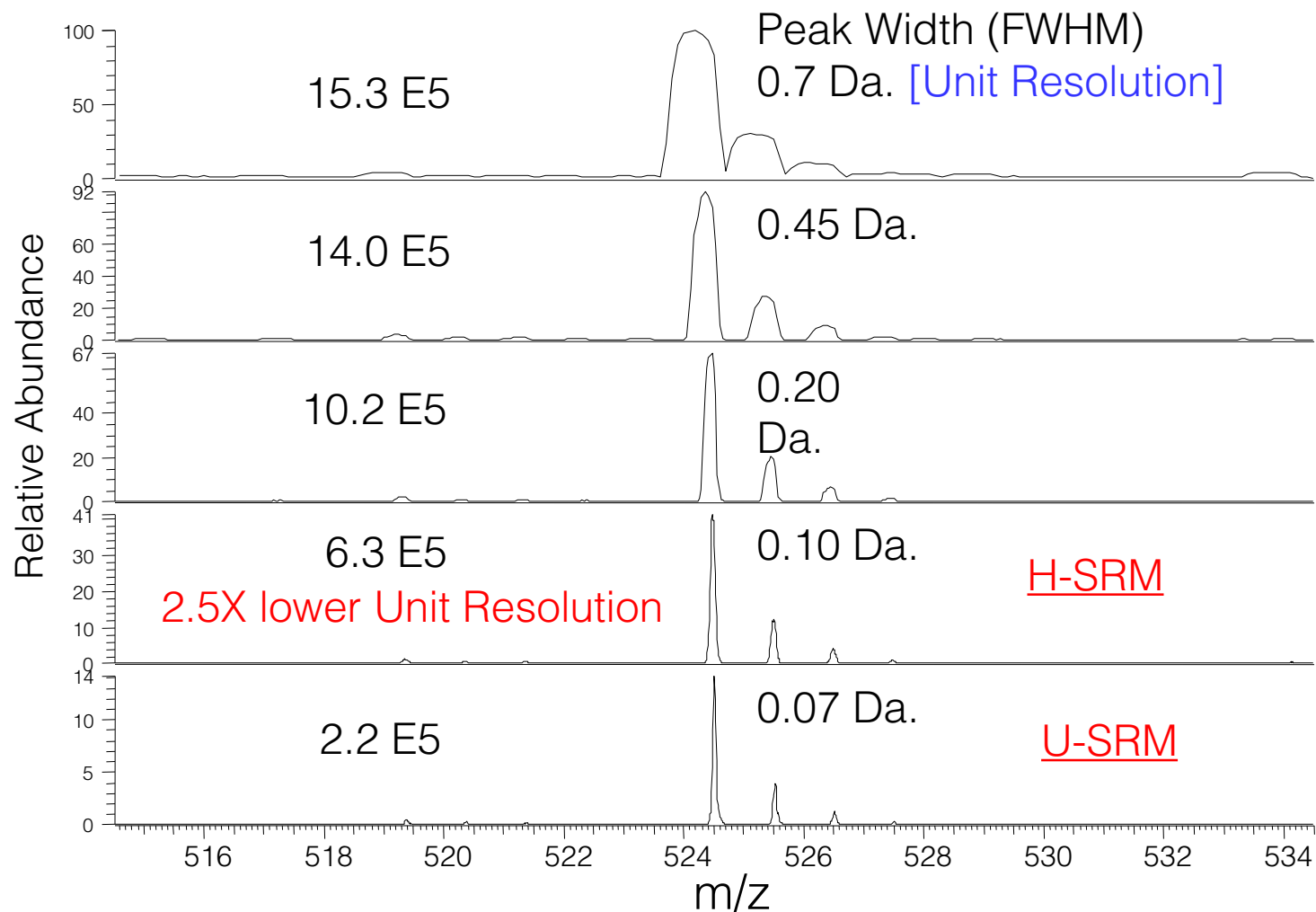


● = only pre-cursor ion transmitted



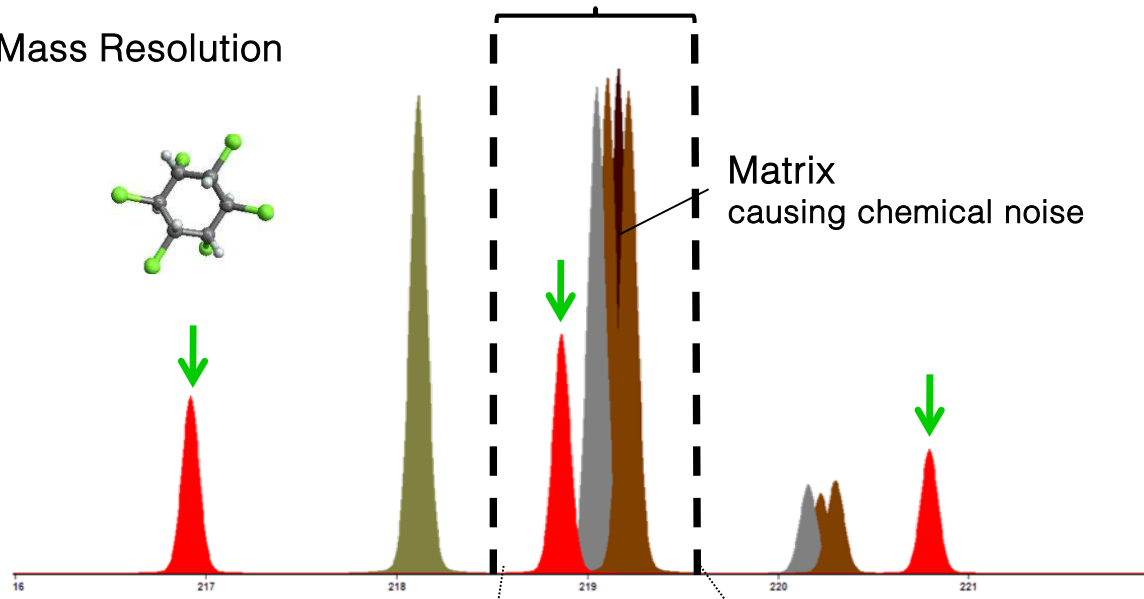


# Transmission Efficiency on TSQ Quantum



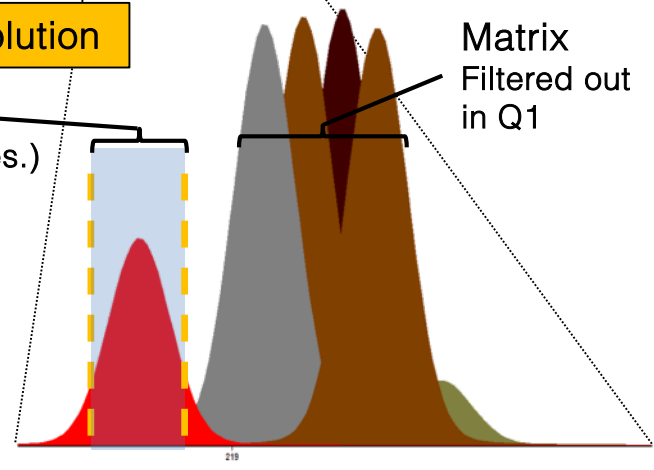
# Mass Selectivity - SRM w. Enhanced Mass Resolution

Nominal Mass Resolution Window



Enhanced Mass Resolution

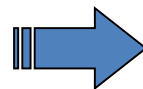
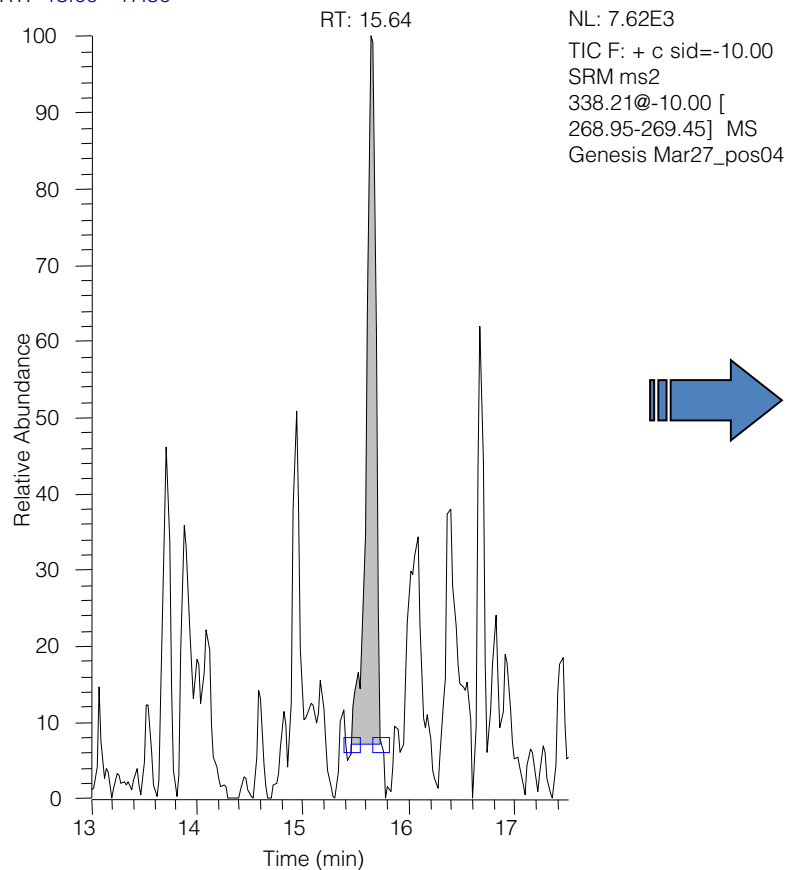
HCH Precursor Ion  
Q1= 218.86 m/z (0.1 Da res.)



# SRM vs H-SRM for analysis of Bitertanol

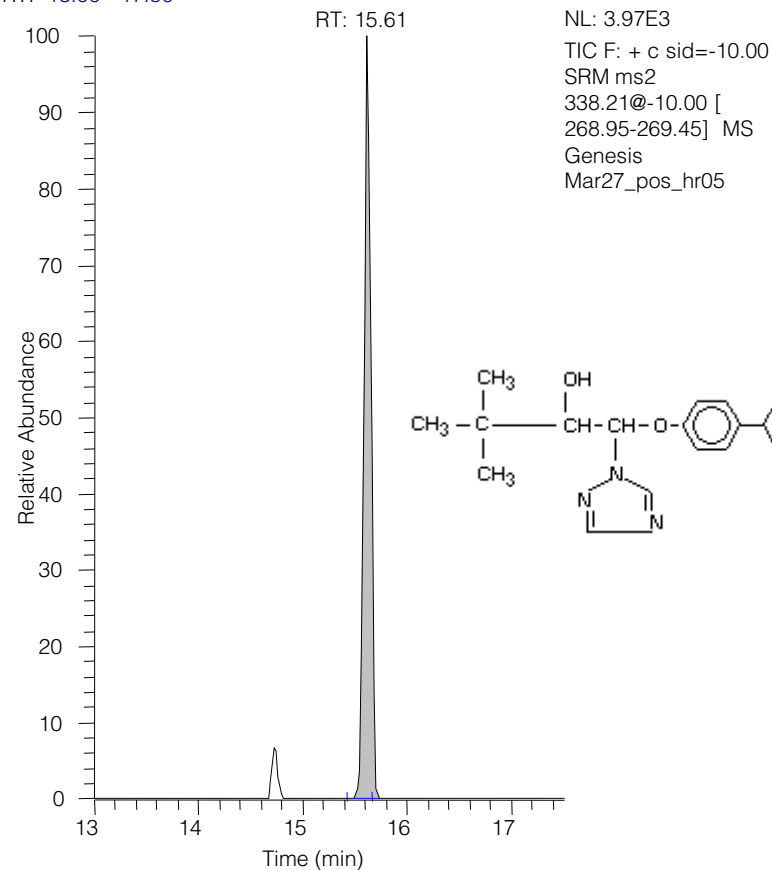
## SRM Mode

RT: 13.00 - 17.50



## H-SRM Mode

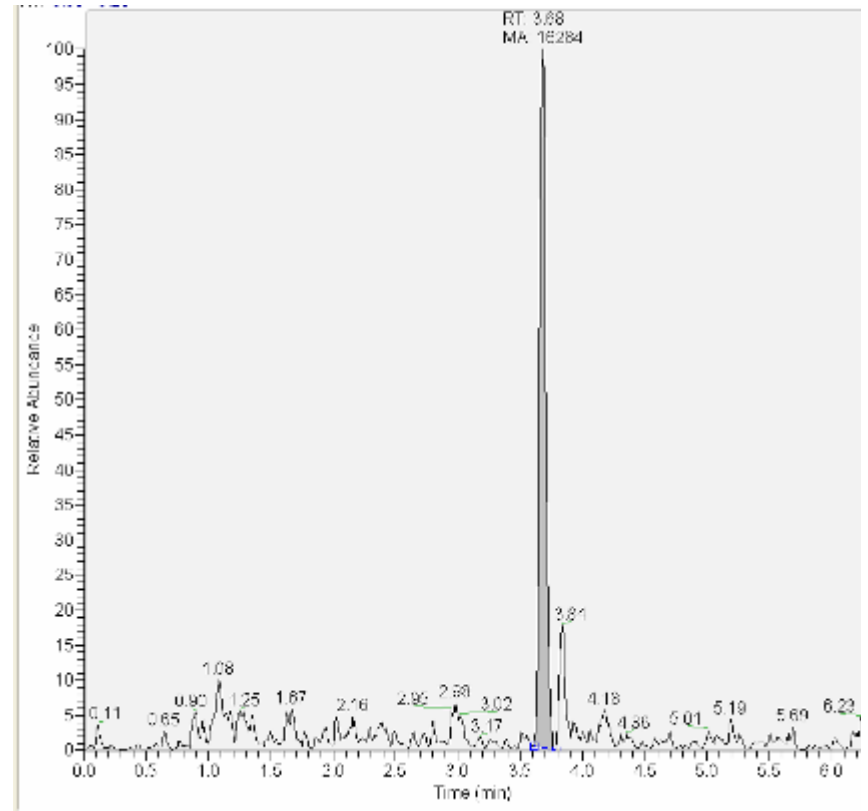
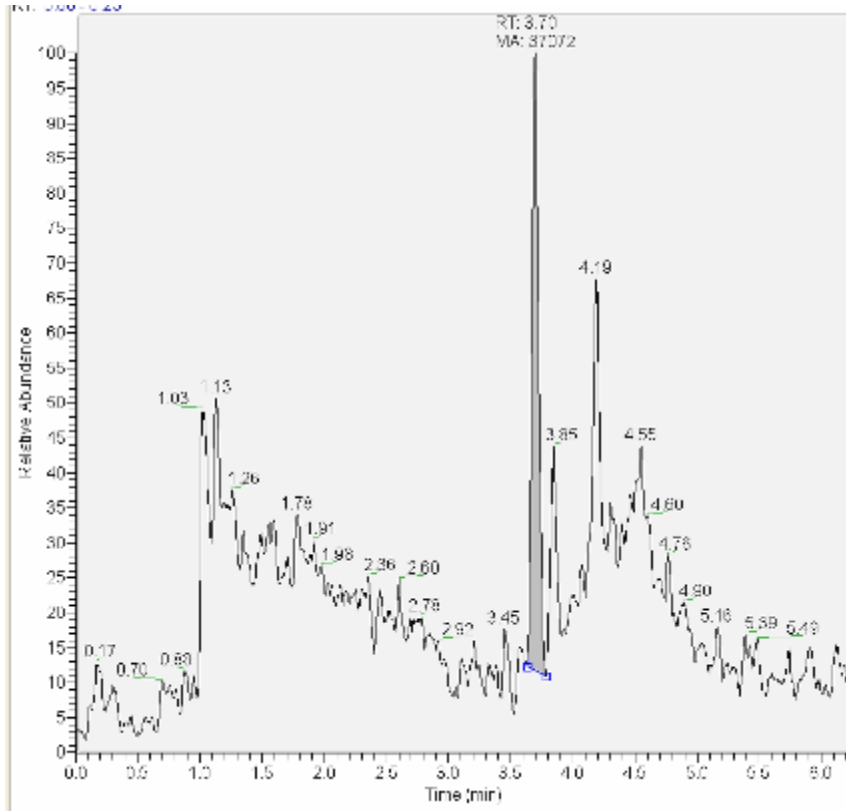
RT: 13.00 - 17.50



# High resolution Comparison

0.7mDa

0.1mDa



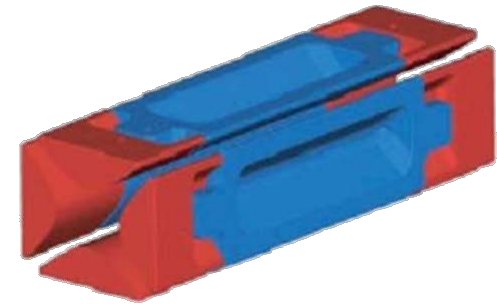
0.5ng/ml hydrolysed Amino Acid standard

# Non-Targeted Screening or Newborn Ideal

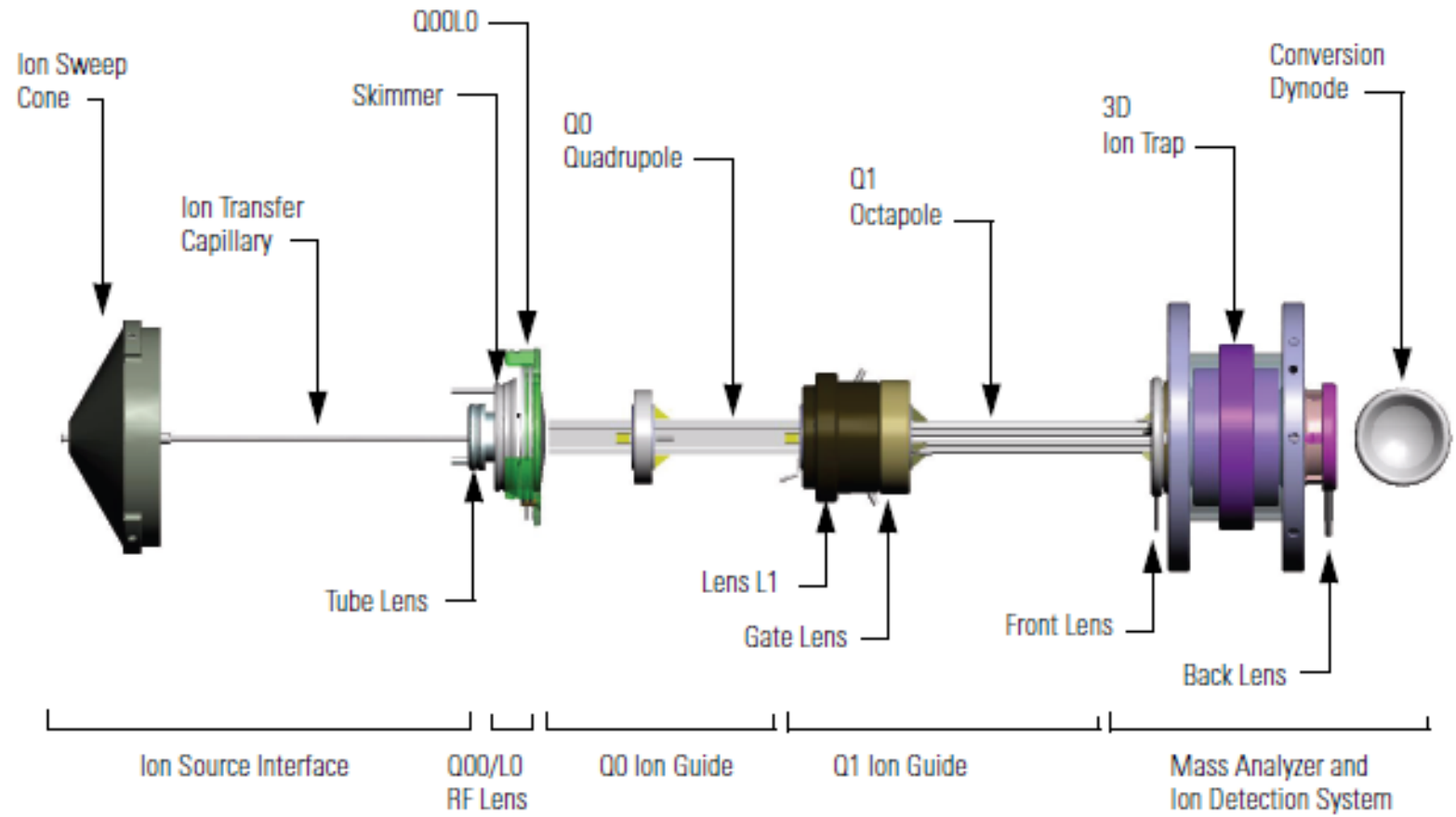
- High isolation power for higher discrimination
- High precision for accurate mass identification
- High resolution for more identification
- High mass stability for a long lasting mass calibration
- $MS^n$
- Library availability for easy interpretations

# Type of Ion Trap Instruments

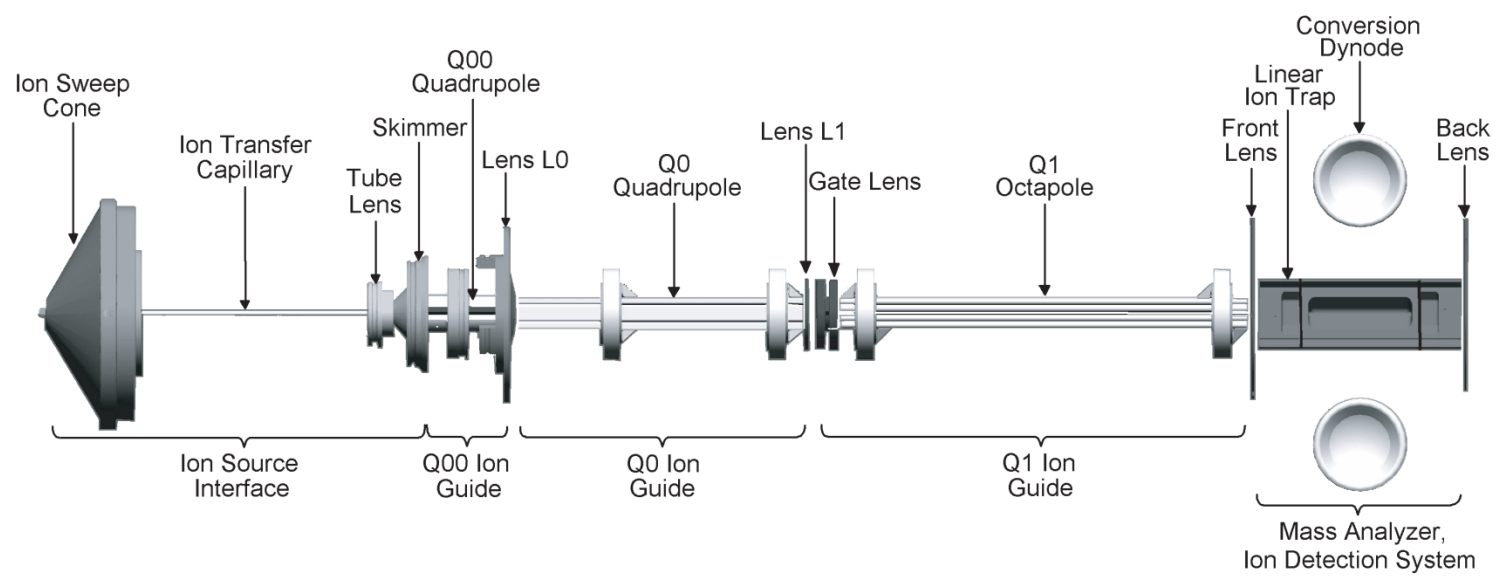
- 3D Ion Trap
- 2D Ion Trap



# Linear Ion Trap (3D) Mass Spectrometer

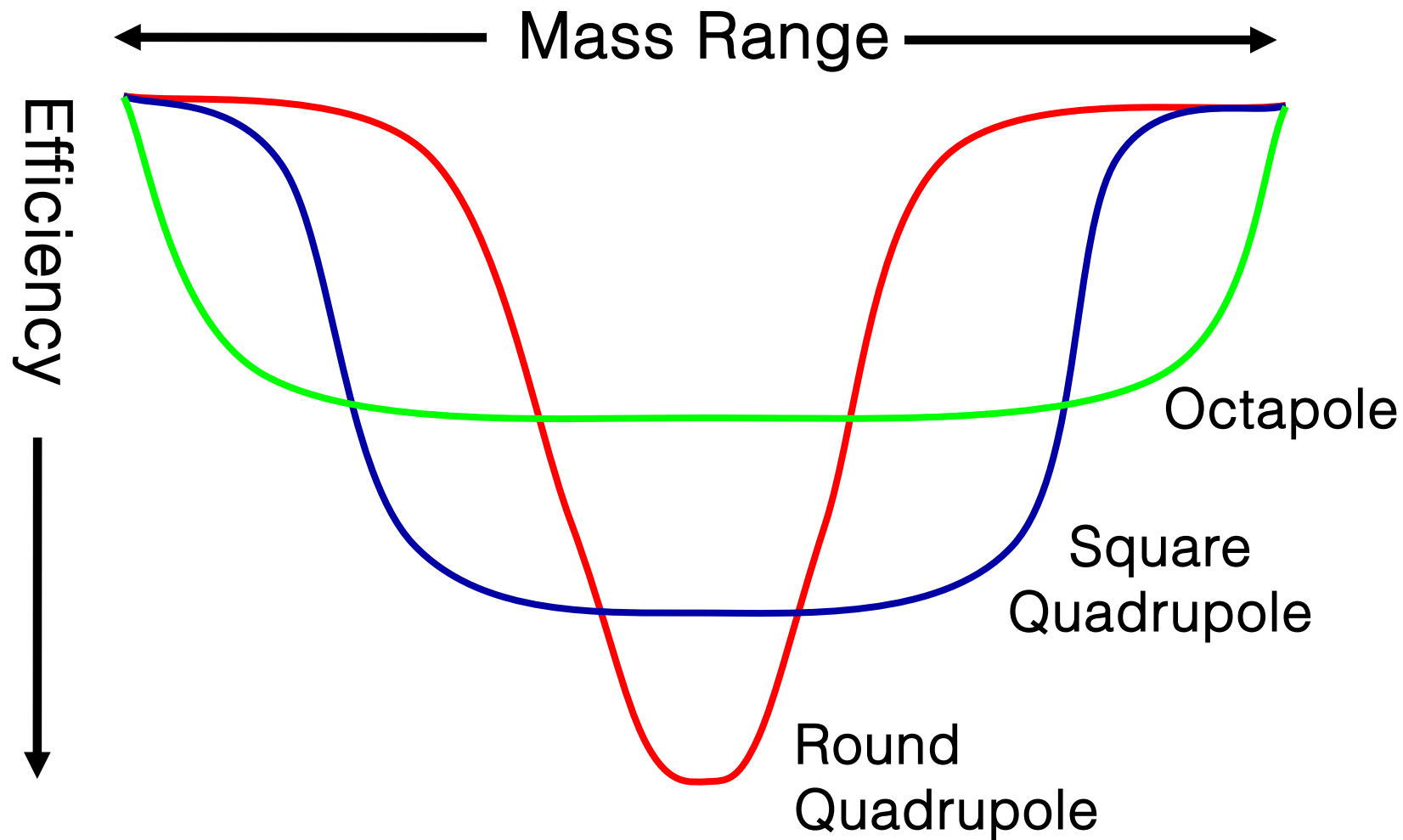


# Linear Ion Trap Quadrupole (2D) Mass Spectrometer

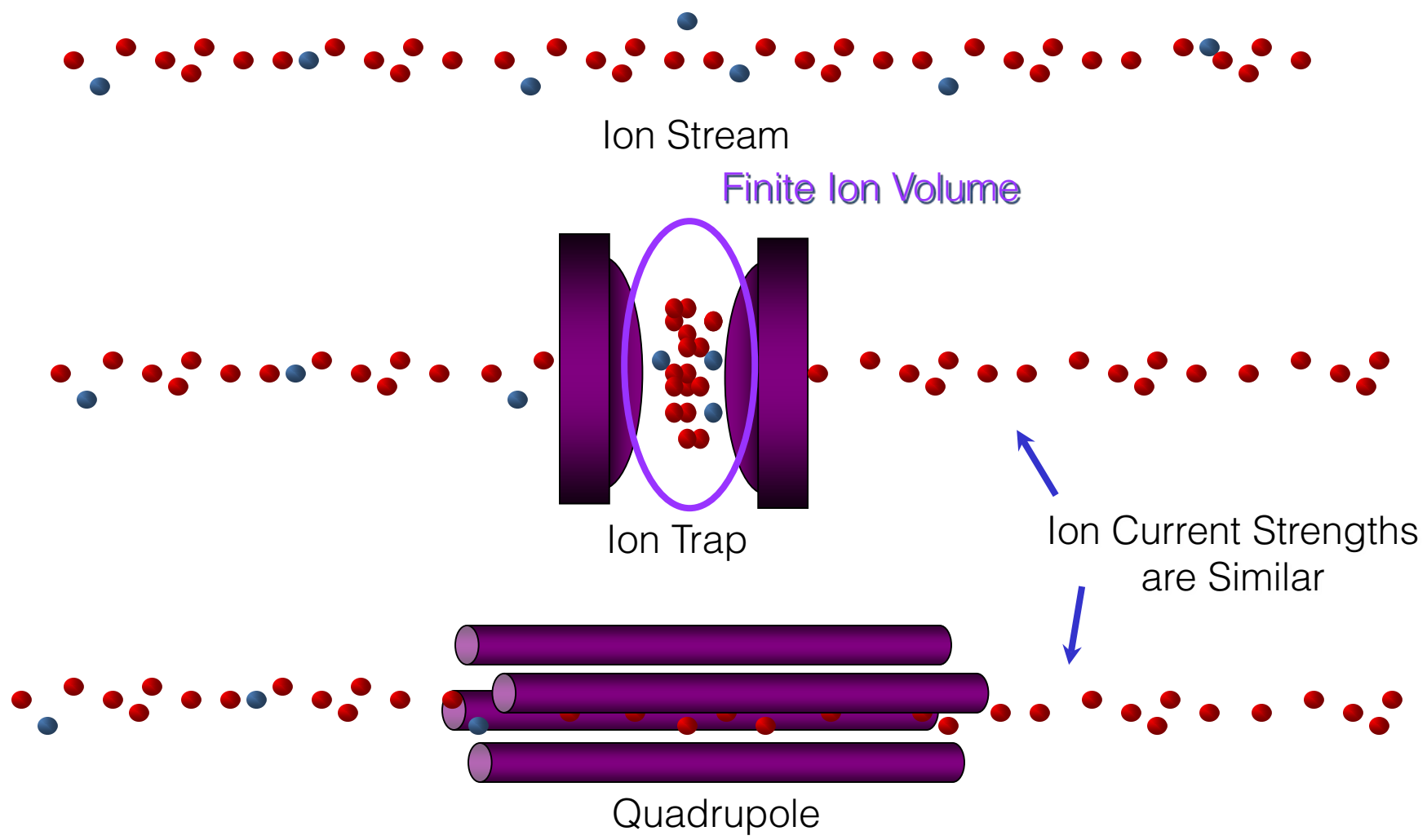




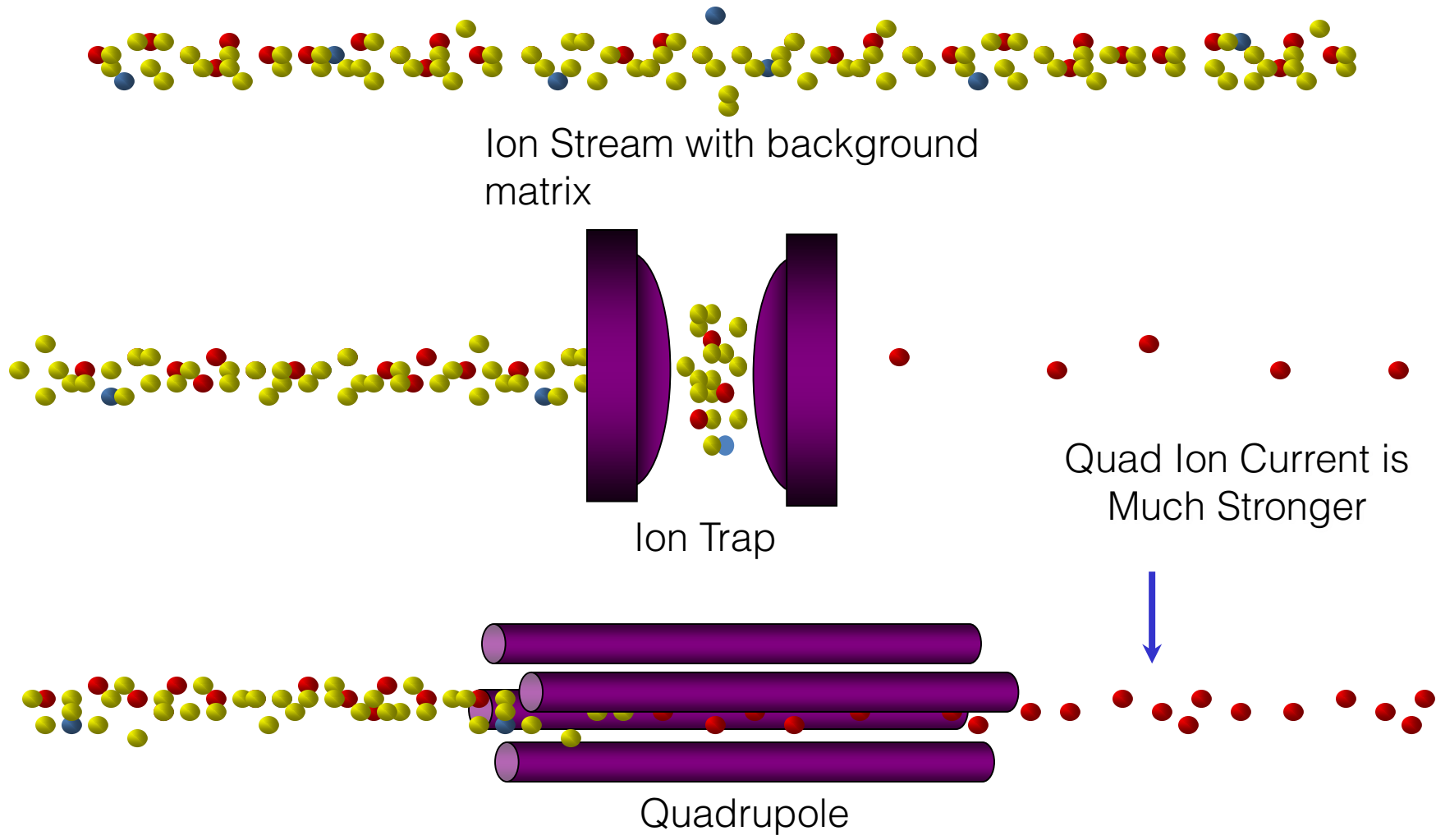
# Why Use Square Rods ? – Pre-filter



# Trap vs. Quad



# Trap vs. Quad



# Advantage of Ion Trap Vs QqQ

Ion Trap	QQQ
Suffered from Matrices	Suitable for high matrices
Target & Non-target	Best for Target, compromise from Non-target
MS <sup>n</sup>	MS/MS
Higher LOQ, LLD	Lowest LOQ, LLD

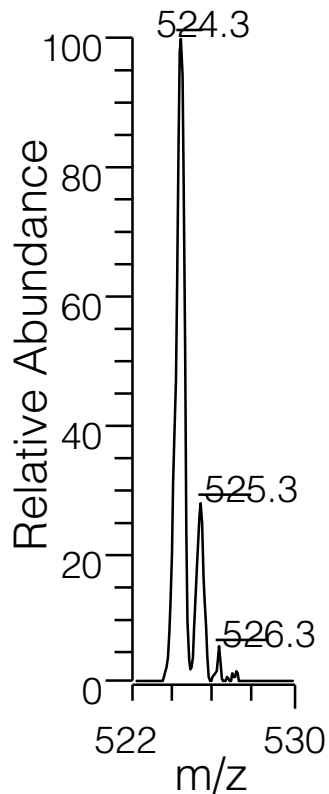
QTRAP, Hybrid MS is the compromised of Ion trap for MS<sup>n</sup> & Single Quadrupole for ion transmitting or Trapping without Quadrupole filtered

# Ion Trap Scan Functions

- Trapping - all scan modes
- Isolation - SIM and MS<sup>n</sup>
- Excitation - MS<sup>n</sup>
- Ejection - all scan modes

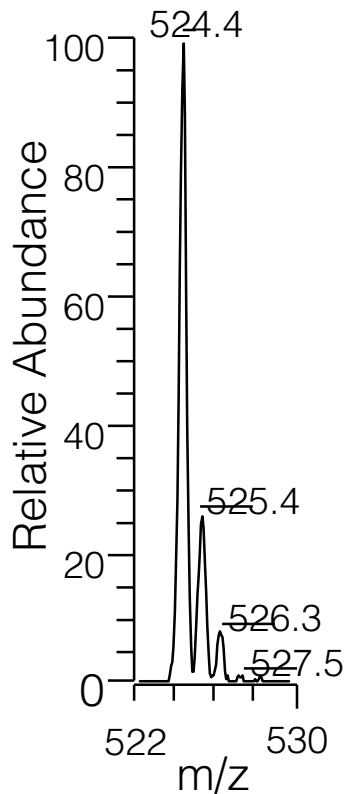
# Space Charge Effects...

~ 300 Ions

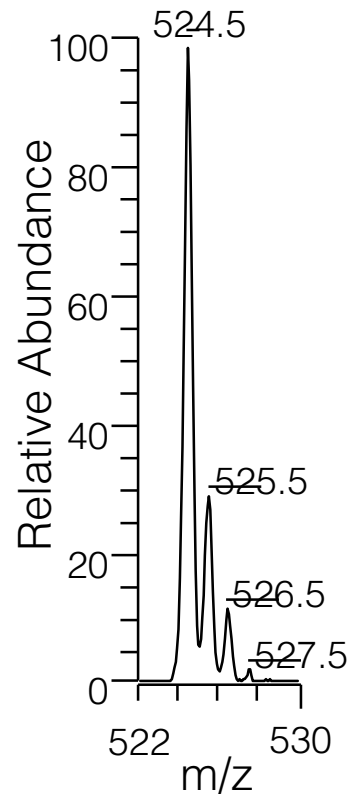


Good Resolution

~ 1500 Ions

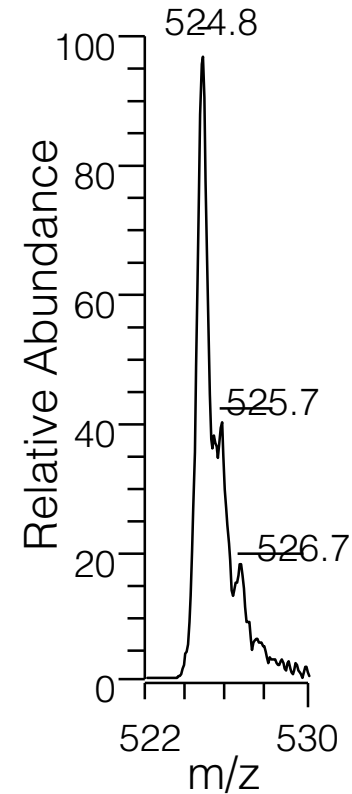


~ 3000 Ions



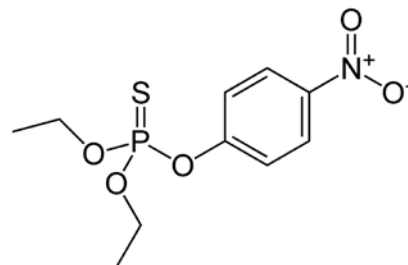
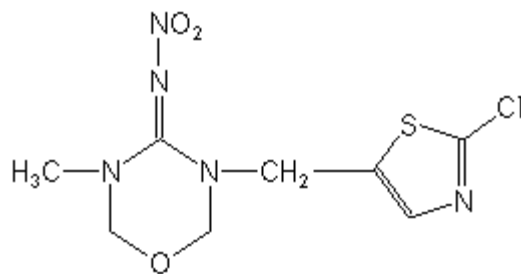
Poor Resolution

~ 6000 Ions



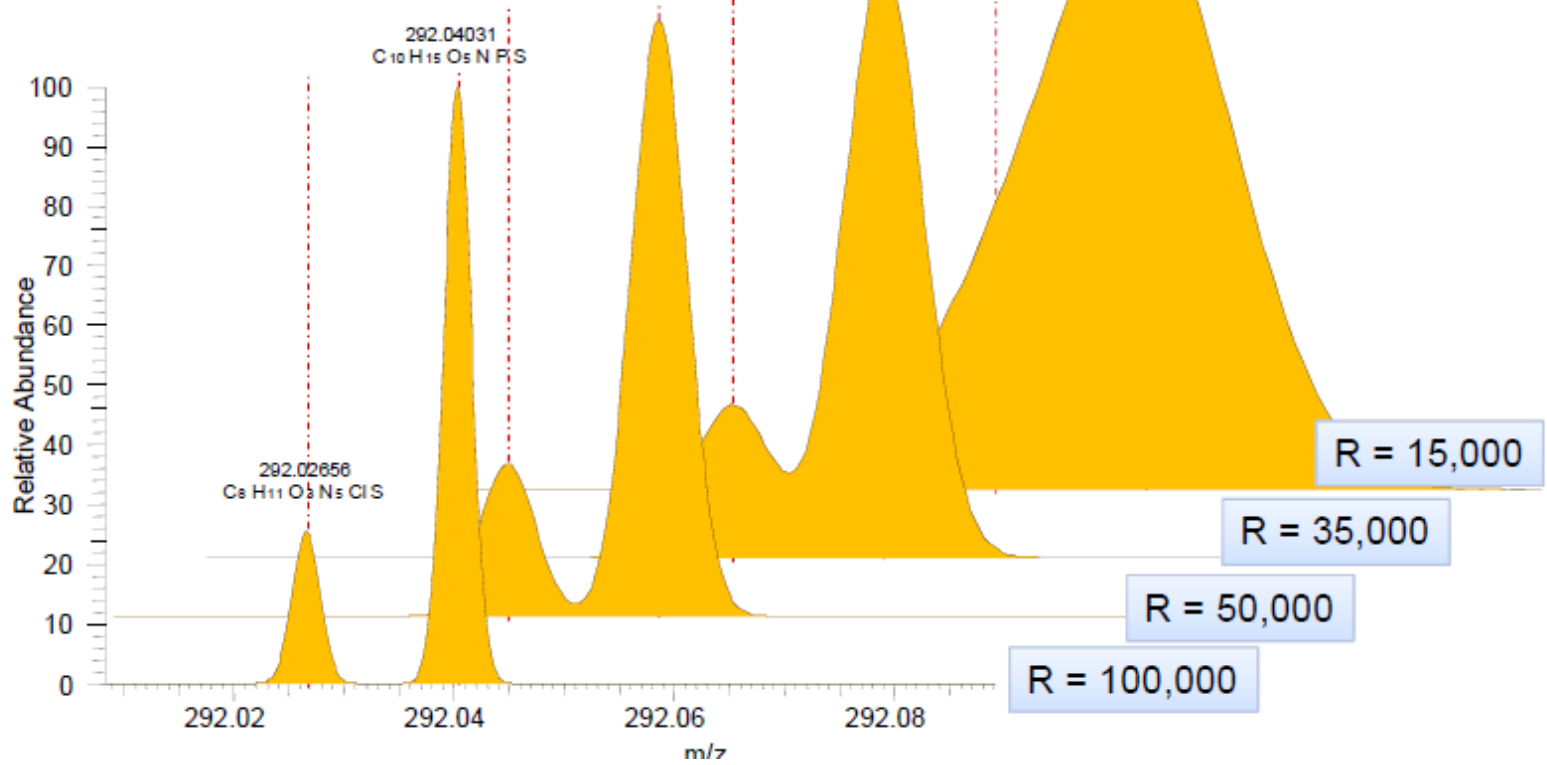
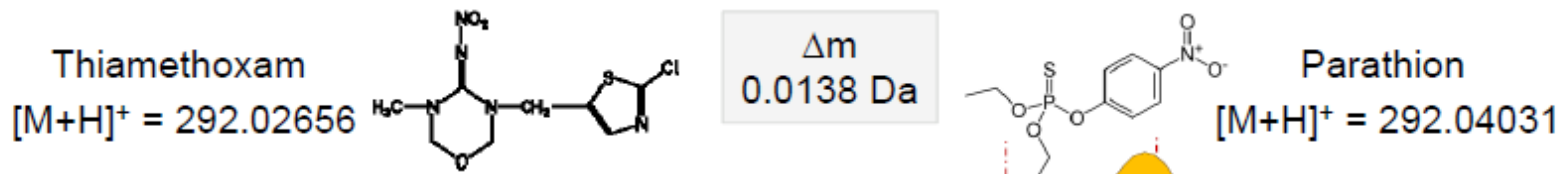
# Isobaric Pesticides

Thiamethoxam:  $[M+H]^+ = C_8H_{11}ClN_5O_3S$   
(292.02656)



Parathion:  $[M+H]^+ = C_{10}H_{15}NO_5PS$  (292.04031)

# Isobaric Pesticides 3:1 Mix

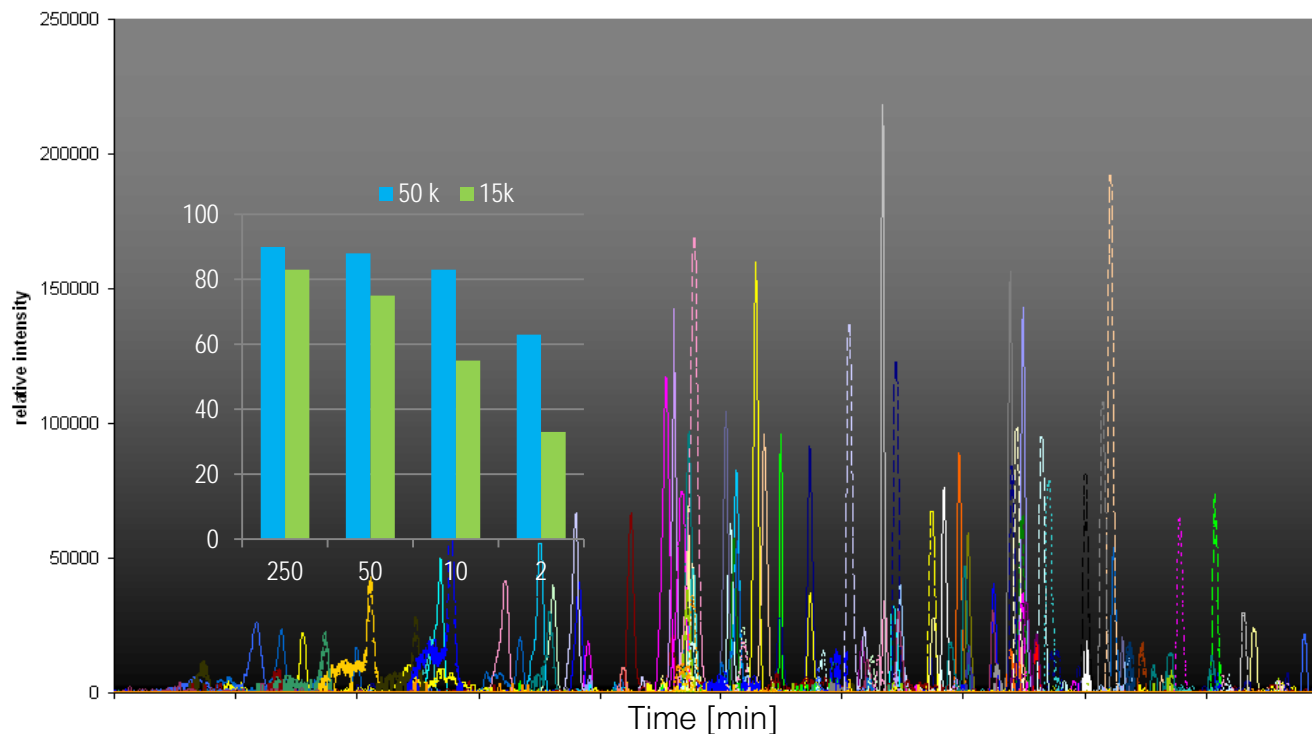




# Pesticide Analysis at different Resolution Settings

Overlaid extracted ion chromatograms from a mixture of 116 pesticides and mycotoxins at a 100ppb level. Extraction was done with 3 ppm mass window. The inset chart shows the number of detected compounds at different concentrations (in matrix) at two different resolution settings

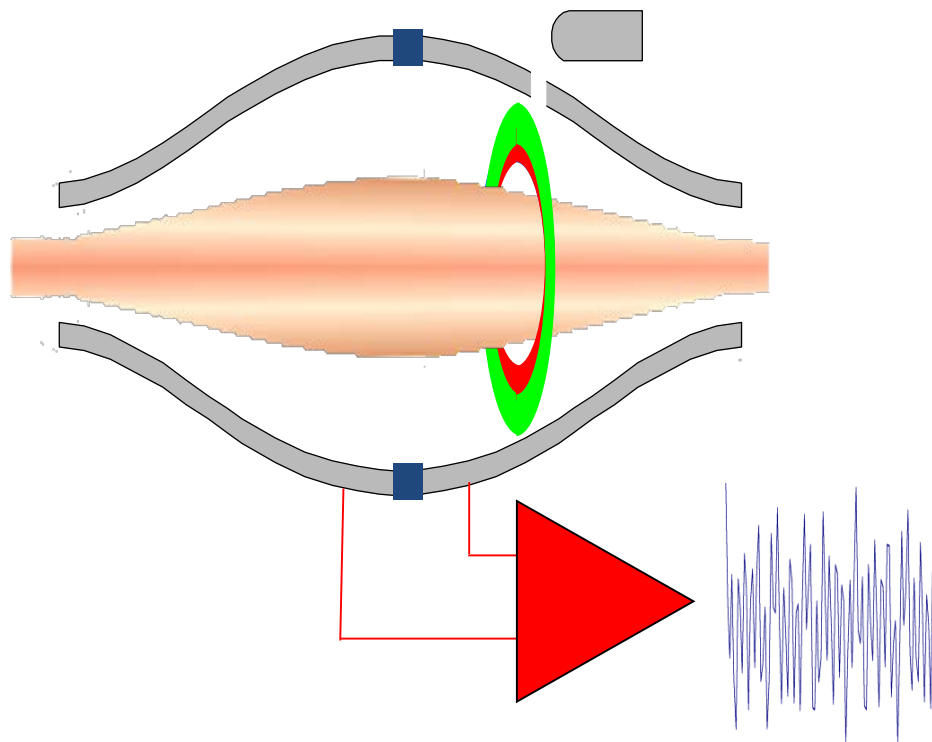
- More Resolution, more Identification !



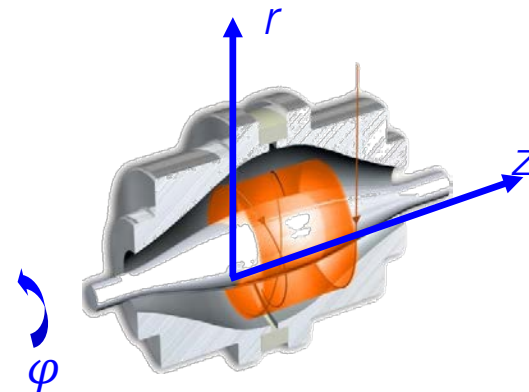
# Typical Mass Accuracy

Type of MS	Mass accuracy	Utility for
Quadrupole	0.1 $\mu$	Identify
Traps	0.1 $\mu$	Identify
TOF	0.0001 $\mu$	Empirical formula/ composition
Sector	0.0001 $\mu$	Empirical formula/ composition
FT-MS	0.0001 $\mu$	Empirical formula/ composition

# Orbitrap MS – Principle of Operation



$$\omega_z = \sqrt{\frac{k}{m/q}}$$



Hyper-logarithmic potential distribution:  
"ideal Kingdon trap"

$$U(r, z) = \frac{k}{2} \left\{ z^2 \left[ \frac{r^2}{2} + R_m^2 \right] \ln(r/R_m) \right\}$$

■ Characteristic frequencies:

- Frequency of rotation  $\omega_\phi$
- Frequency of radial oscillations  $\omega_r$
- Frequency of axial oscillations  $\omega_z$

$$\omega_\phi = \frac{\omega_z}{\sqrt{2}} \sqrt{\frac{R_m}{R}} \approx \omega_z \quad \omega_r = \omega_z \sqrt{\frac{R_m}{R}} \approx \omega_z \sqrt{2}$$

# Technologies Shift - Key Performance of ICRMS

- Very Precise Mass (<1 ppm)
- High Resolution (up to 1,000,000 FWHM)
- Excellent Matrices Elimination
- Isotopic Analysis
- MS<sup>n</sup> capabilities – *Alexandre Makarov invented C-Trap for linkage between collision cell and ICR*
- *Step Over TOF/Q-TOF/TOFTOF limitation; Major drawback – TDC (time to digital converter)*
- *Although very fast, the TDC is a low cost, ion counting detector – its dynamic range is limited due to its inability to properly count the events, particularly when more than one ion simultaneously hits the detector and also b/c of the deadtime incurred after each count. Additionally, with higher sample concentration, two or more distinct isobaric peaks will not be detected when hitting the detector at the same time, resulting in improper peak height and inaccurate m/z reported*

# Mass Spectrometer Database

- [www.mzcloud.org](http://www.mzcloud.org)
- [www.chemspider.com](http://www.chemspider.com)
- [www.massbank.jp](http://www.massbank.jp)
  
- Different dataset in each type of instrument