

Fundamentals of Single Quadrupole Mass Spectrometry

Waters Corporation

Noud van der Borg

Introduction

- In this presentation we will review the fundamentals of mass spectrometry for current liquid chromatography users

What Is Single Quadrupole Mass Spectrometry and How Is It Used?

Short explanation on other MS techniques and MS-sources

Overview

- What is mass spectrometry?
 - Who uses it and why?
- How do single quadrupoles work?
- What type of data is collected? What impacts the observed mass-to-charge ratio?

Who Uses MS and Why?

■ What industries use MS?

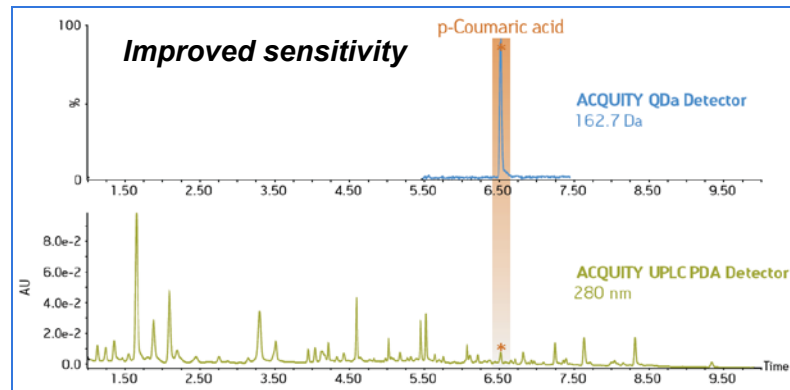
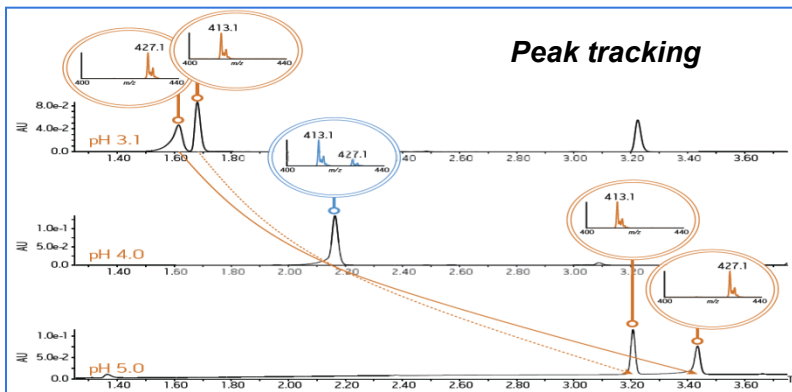
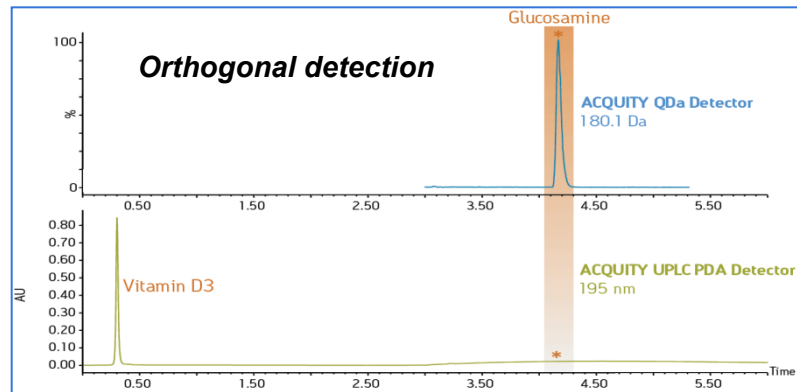
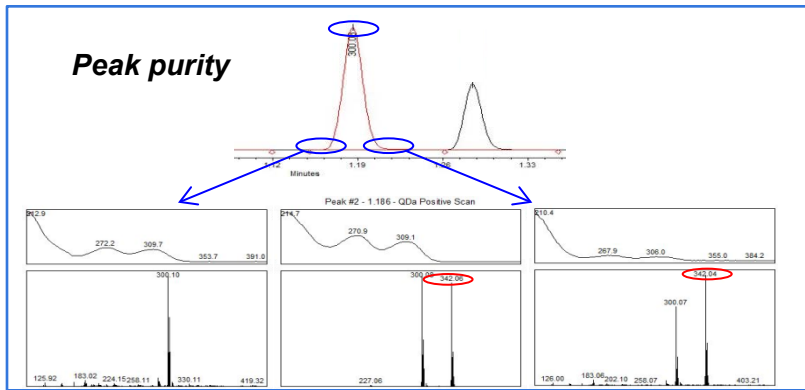
- Pharmaceutical
 - Reaction Monitoring, Purification, Characterization
 - Drug Metabolism & Pharmacokinetics, Degradation and Bioavailability
- Food and Environmental
- Industrial
- Health Science



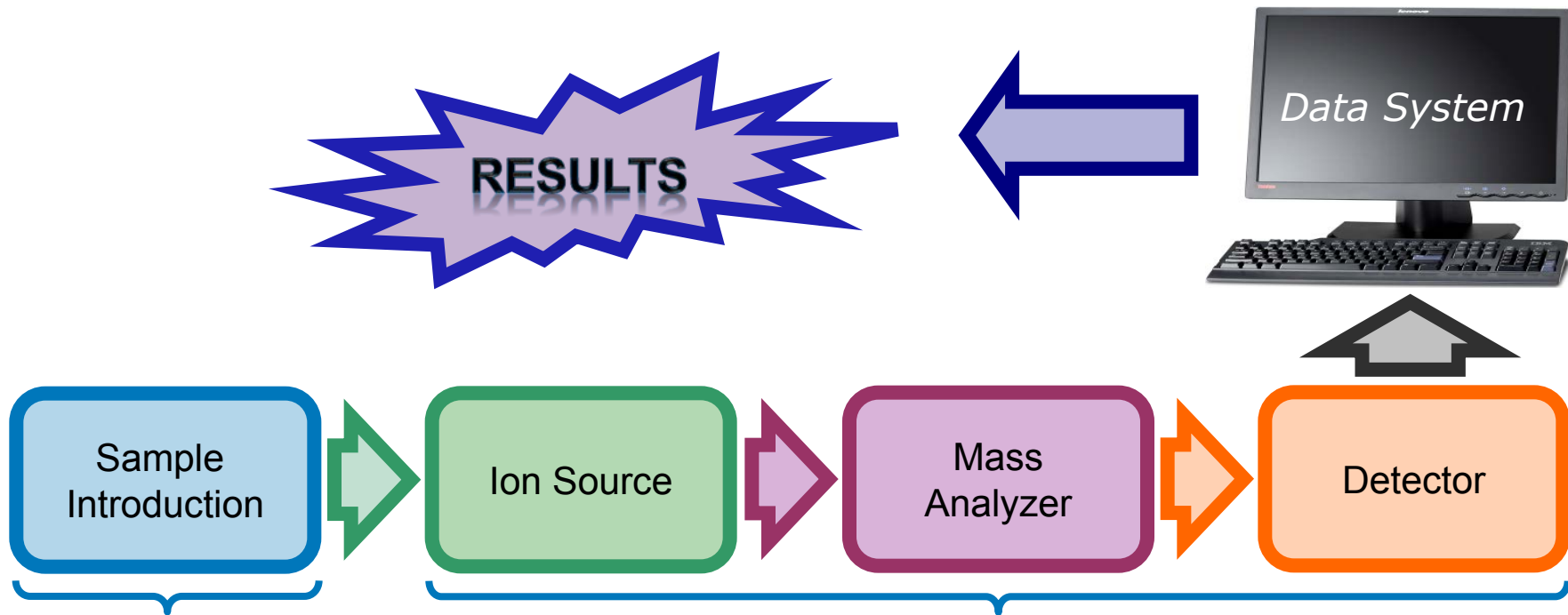
■ Why?

- QC
 - Confirmation of analytes
 - Check raw ingredients
- Synthesis Labs or Reaction Monitoring
 - Confirmation of products
- Method Development
 - Basic peak tracking
 - Peak purity/co-elutions

Advantages of Adding Mass Spectrometry



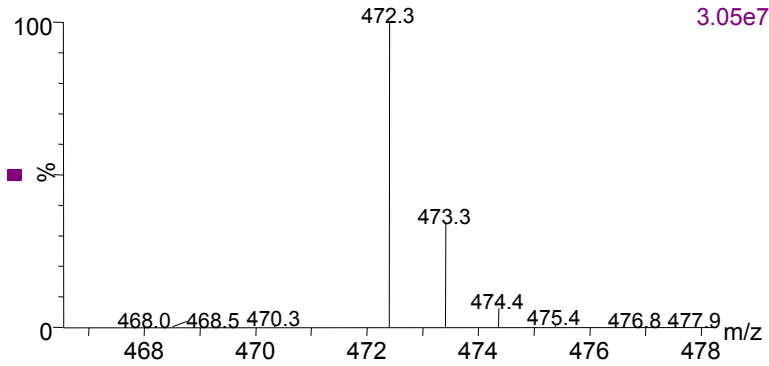
What is Mass Spectrometry?



LC, GC, direct infusion, etc..

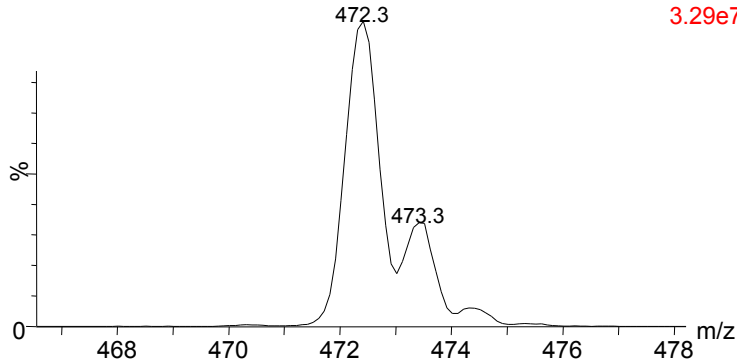
Mass Spectrometer (MS Detector)

History of MS



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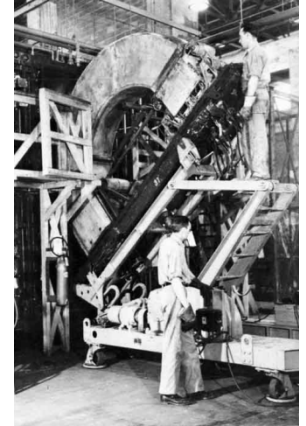
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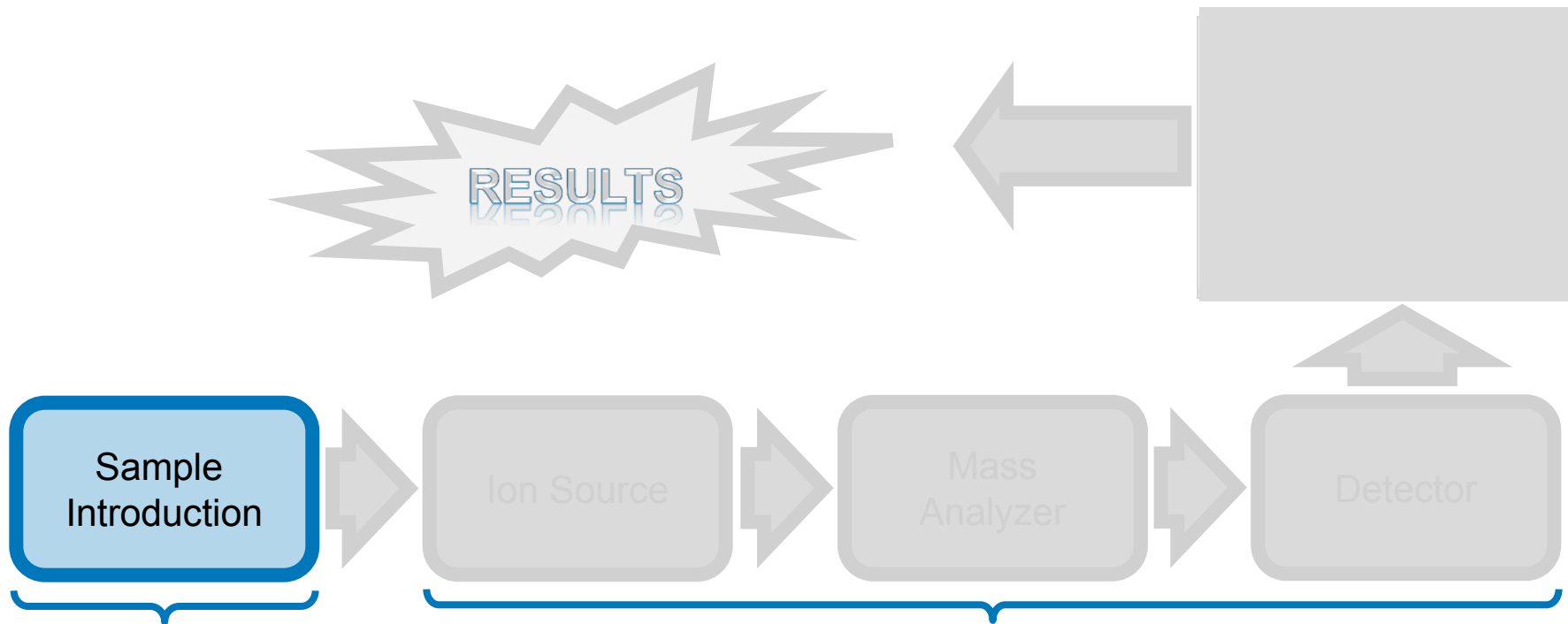
Started in 1886
First only GC-MS systems,
ionization of liquid spray was
difficult

Computers changed MS
TOF was invented before 1940
but start to be useful after 1990
(use of PC's)

PC's could not handle
continuum and centroid was
developed, now easier to read



Sample Introduction

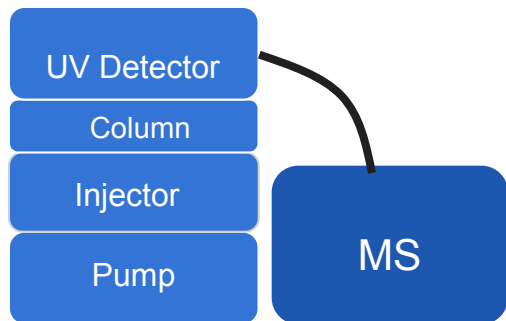


LC, GC, direct infusion, etc..

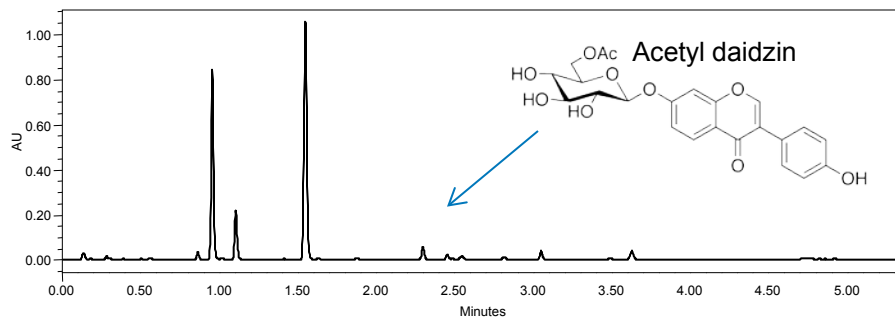
Mass Spectrometer (MS Detector)

Sample Introduction

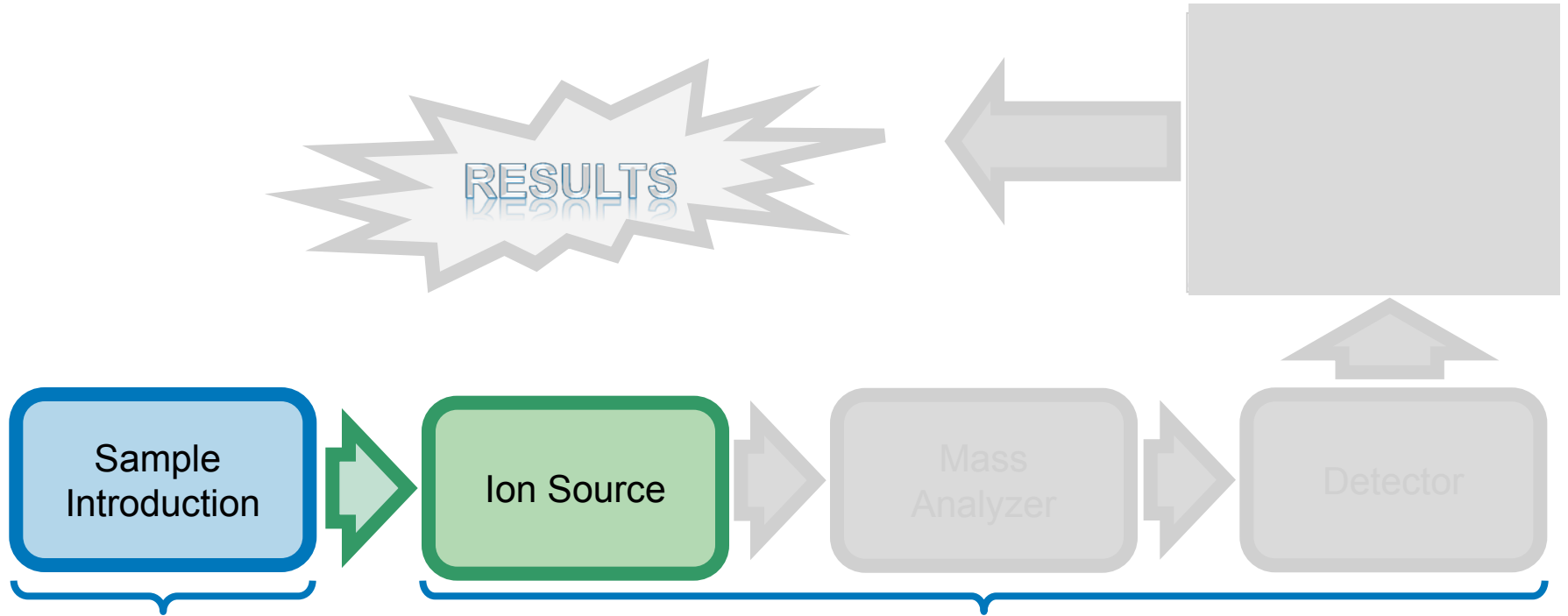
- To analyze a compound by MS detection the sample must be introduced into the system
- For the examples used in this presentation, the MS is coupled to an LC system
- Regardless of the mode of sample introduction, the same data is produced
 - The sample shown contains isoflavones, such as acetyl daidzin, in a dietary supplement



Isoflavones in soy extract



Ion Source

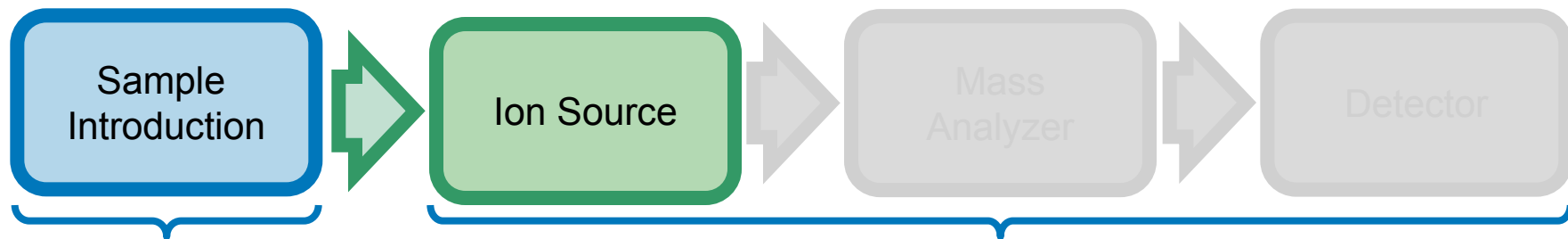


LC, GC, direct infusion, etc..

Mass Spectrometer (MS Detector)

Ion Source

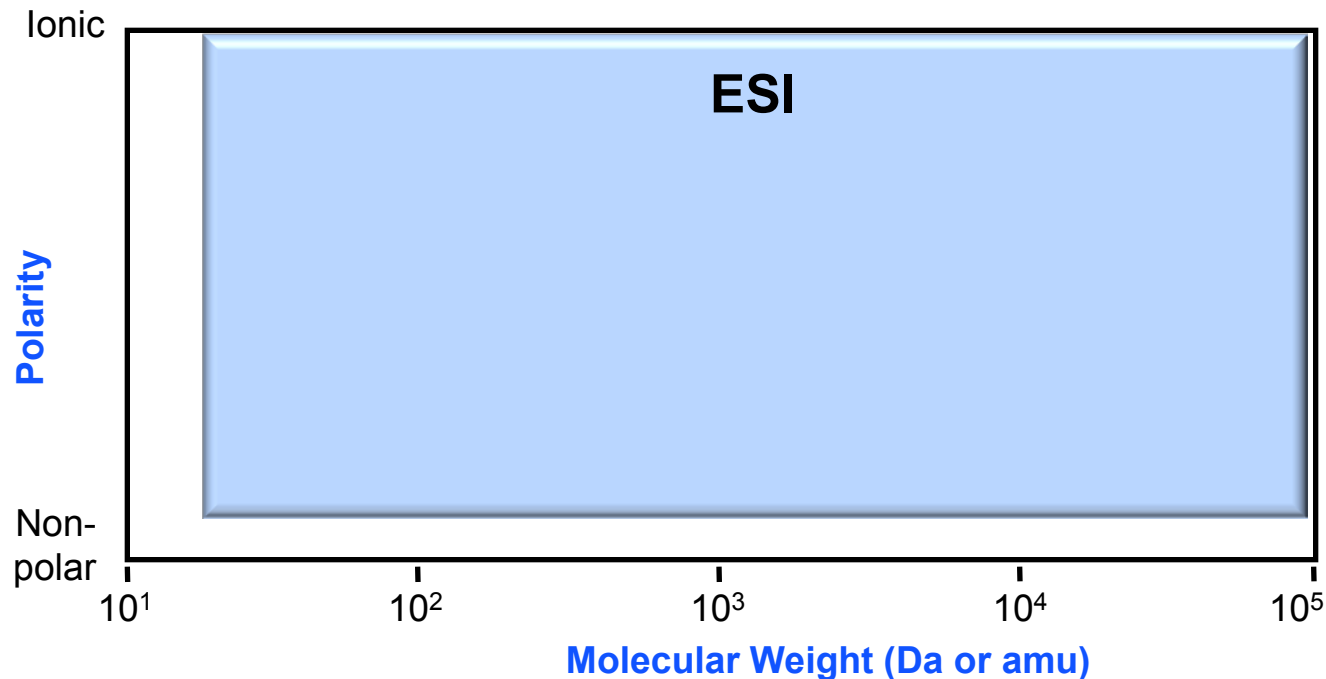
- Some common ionization techniques
 - Electrospray ionization (ESI)
 - Electron ionization (GC, EI)
 - Atmospheric pressure chemical ionization (APCI)
 - Desorption/ionization on silicon (DIOS)
 - Direct analysis in real time (DART)



LC, GC, direct infusion, etc..

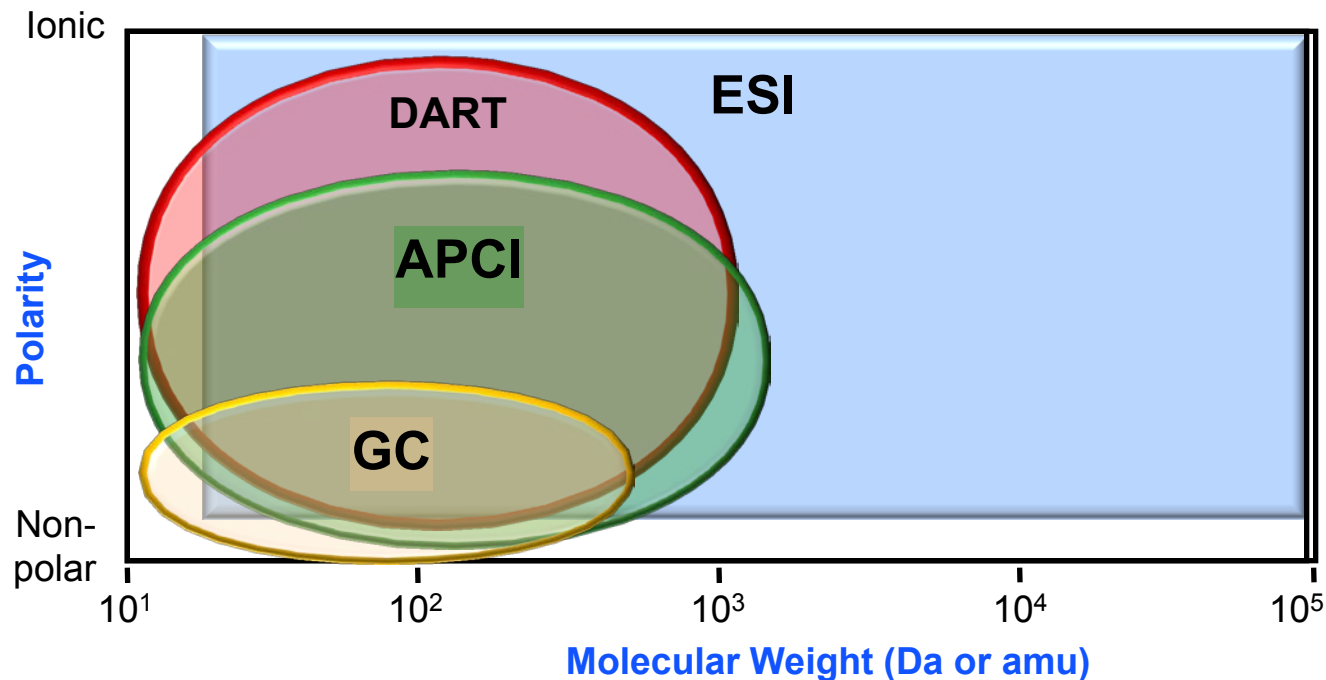
Mass Spectrometer (MS Detector)

Electrospray Ionization: Sample Properties



- Electrospray can ionize compounds up to 200,000 – 500,000 Da
- Better for polar or charged compounds (as compared to non-polar analytes)
- Other techniques may have a more limited range

Electrospray Ionization vs. Other Ionization Techniques: Sample Properties

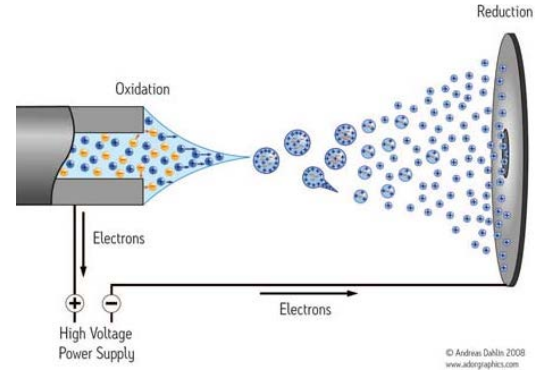
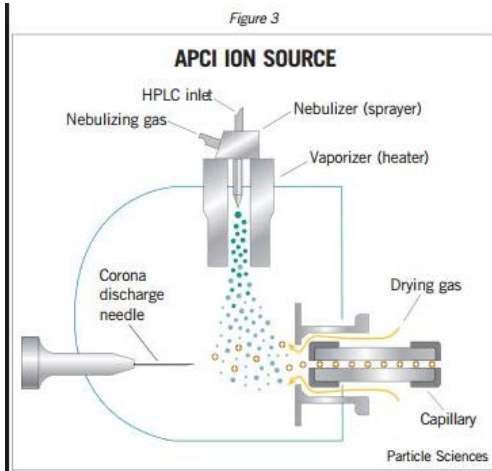


- GC and APCI ionization techniques are optimum for molecules at lower MW range and less polar molecules
- DART ionization technique is applicable for lower MW analytes

Different MS sources

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ESI is mostly used



Multimode source
ESI – Electrospray Ionization
APG – Atmospheric Pressure
Chemical Ionization
ESQ™ – Dual ESI and APG



Dual mode source
APPI – Atmospheric
Pressure Photo
Ionization
APCI – Atmospheric
Pressure Chemical
Ionization

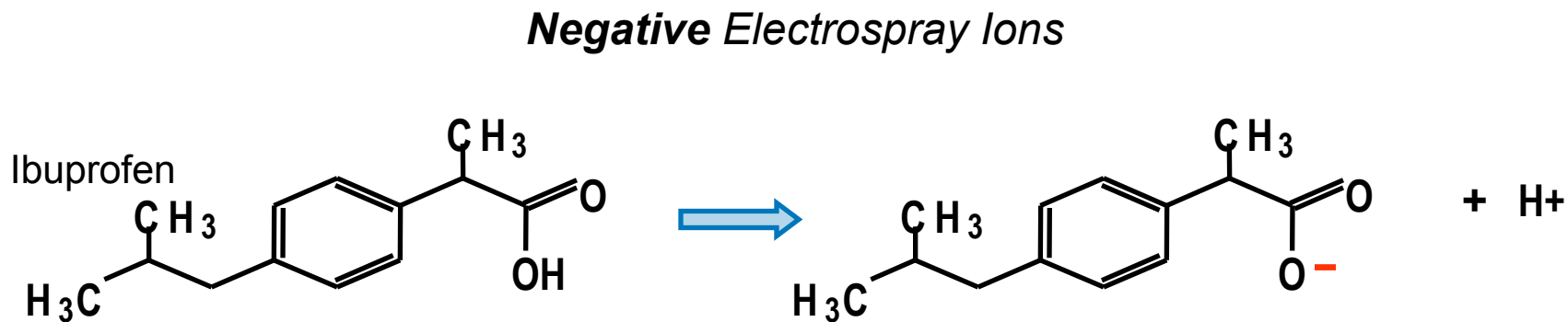
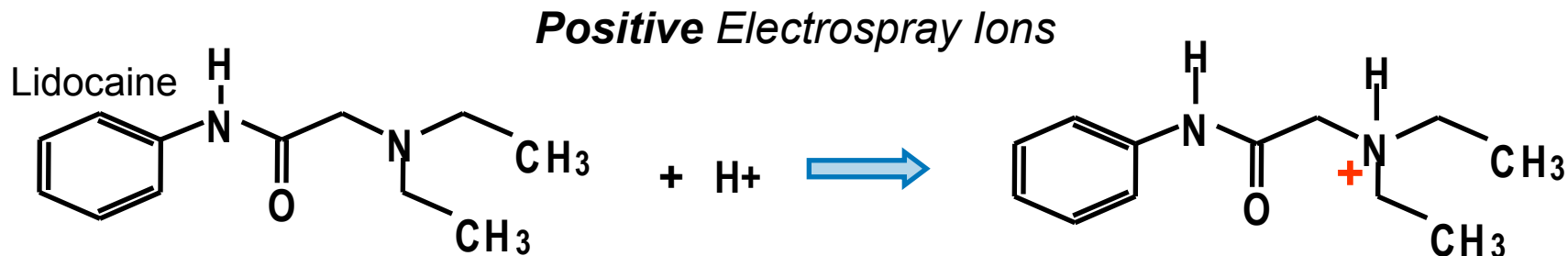


IonKey Source
with nanoTile Technology.
Plug & Play nanoFlow

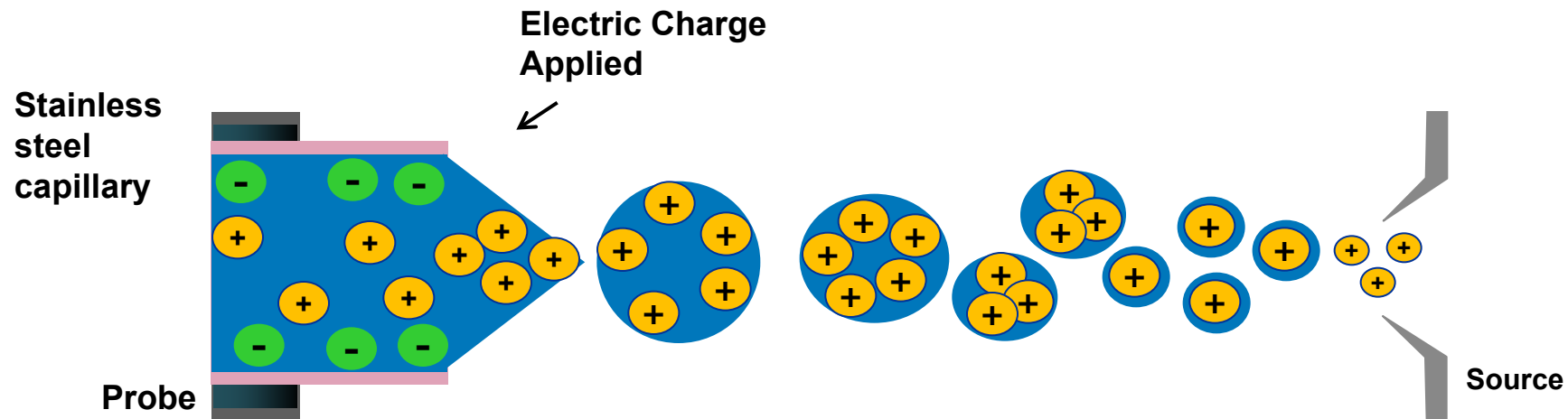


nanoFlow™ ESI

Electrospray Ionization Mechanisms of Ion Formation

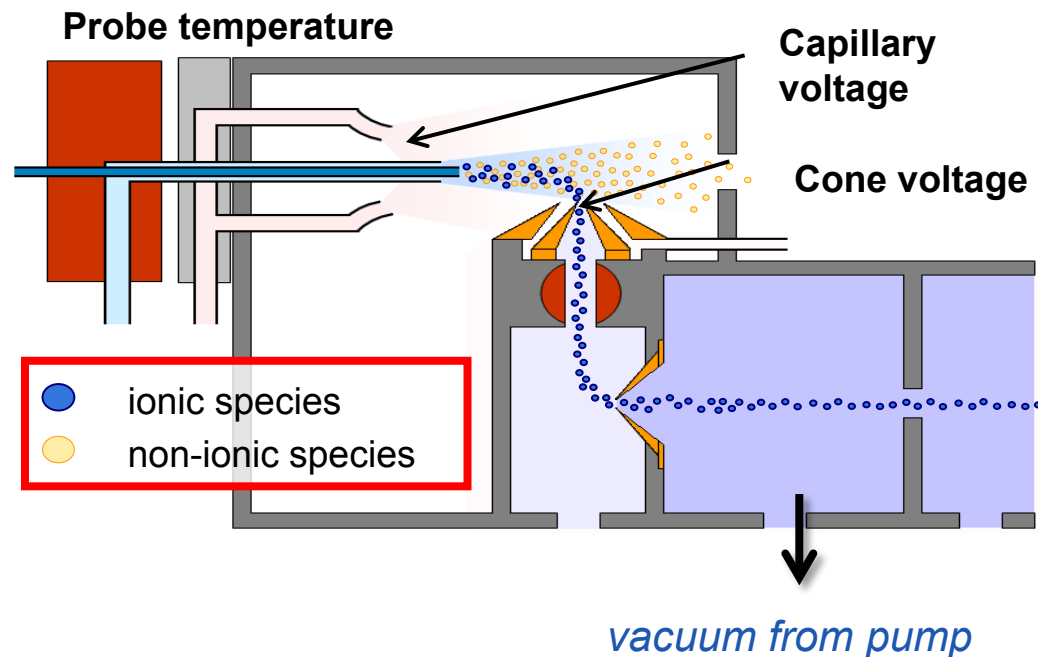


Forming Gas-Phase Ions



- Ionization takes place at atmospheric pressure and has three stages:
 - Formation of charged droplets
 - Solvent evaporation and droplet fission
 - Formation of gas-phase ions
 - Generally produces $(M+H)^+$ or $(M-H)^-$ ions

Introducing Sample into Mass Analyzer

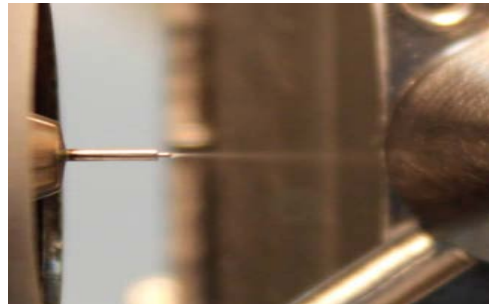
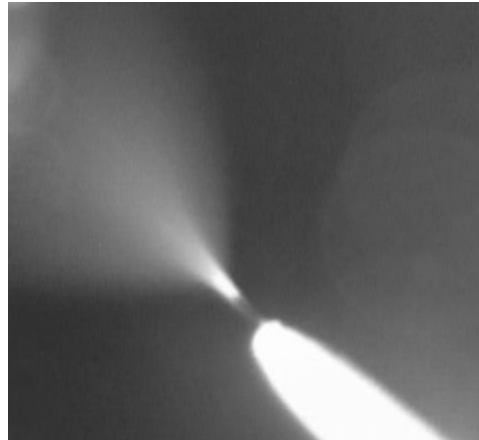


- **Probe temperature:** High temperature applied to the probe
- **Capillary voltage:** Potential used in the capillary of the ESI probe
- **Cone voltage:** Potential applied to the cone located at the entrance of the source
- After ionization, the ions are directed to mass analyzer by vacuum from a pump

Spray, ionization and use of salts

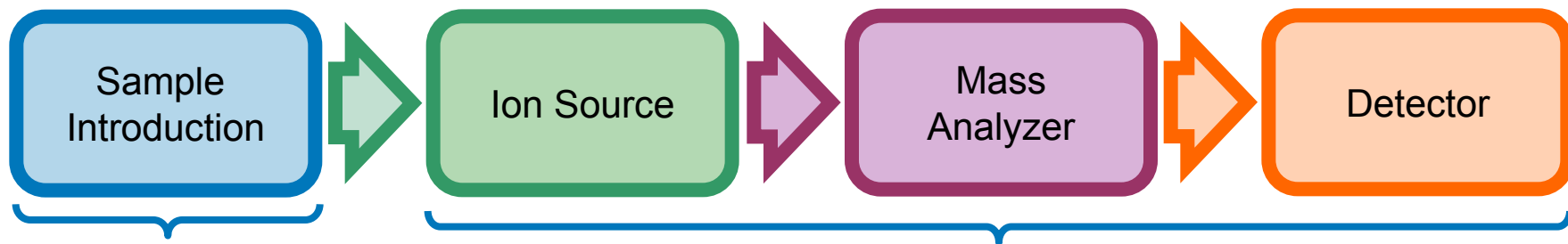
To get a good ionization a spray with small particles is needed.

Creating this, salts will fall out and crystallize



Single Quadrupole Mass Analyzer and Detector

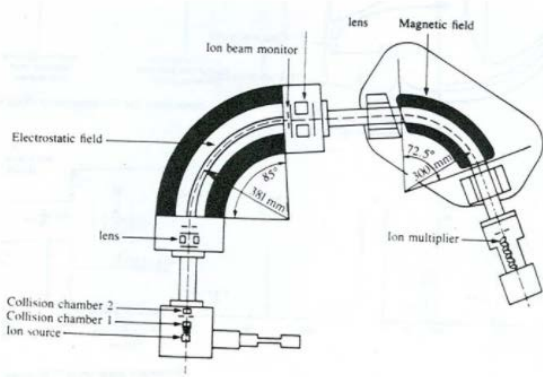
- To detect ions, the ions must be filter or separated
- This occurs in the mass analyzer which includes a pre-filter and the quadrupole
- The measurement of the ions occurs in the detector



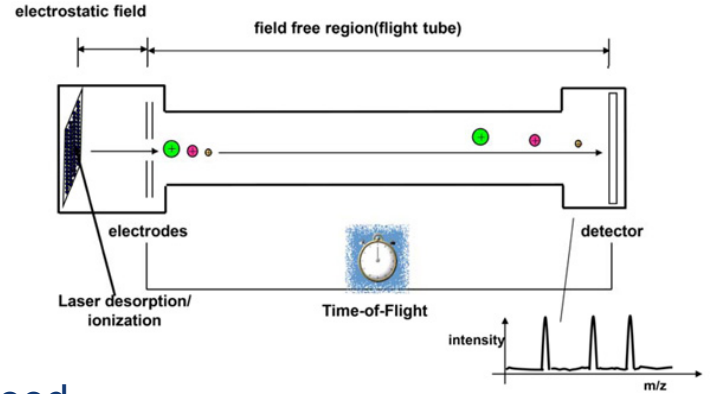
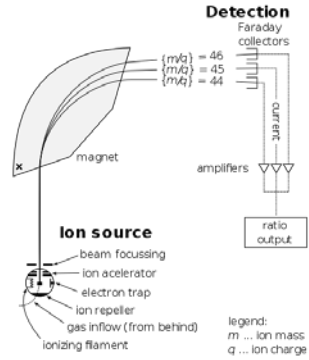
LC, GC, direct infusion, etc..

Mass Spectrometer (MS Detector)

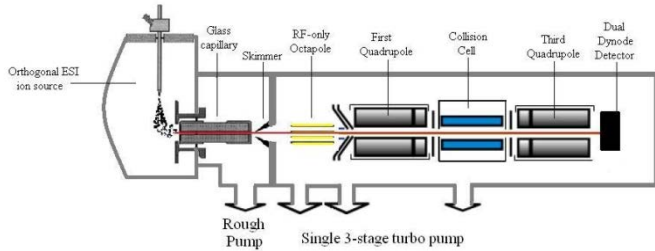
Different mass selection options



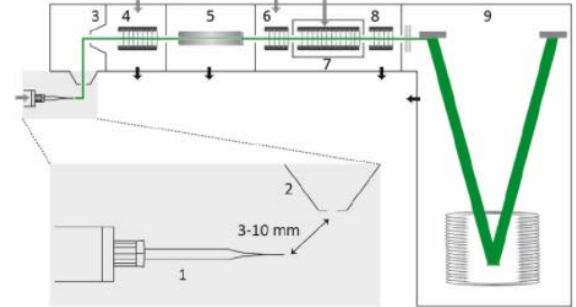
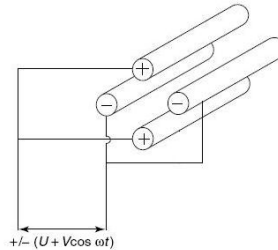
Curve



Speed



Slalom

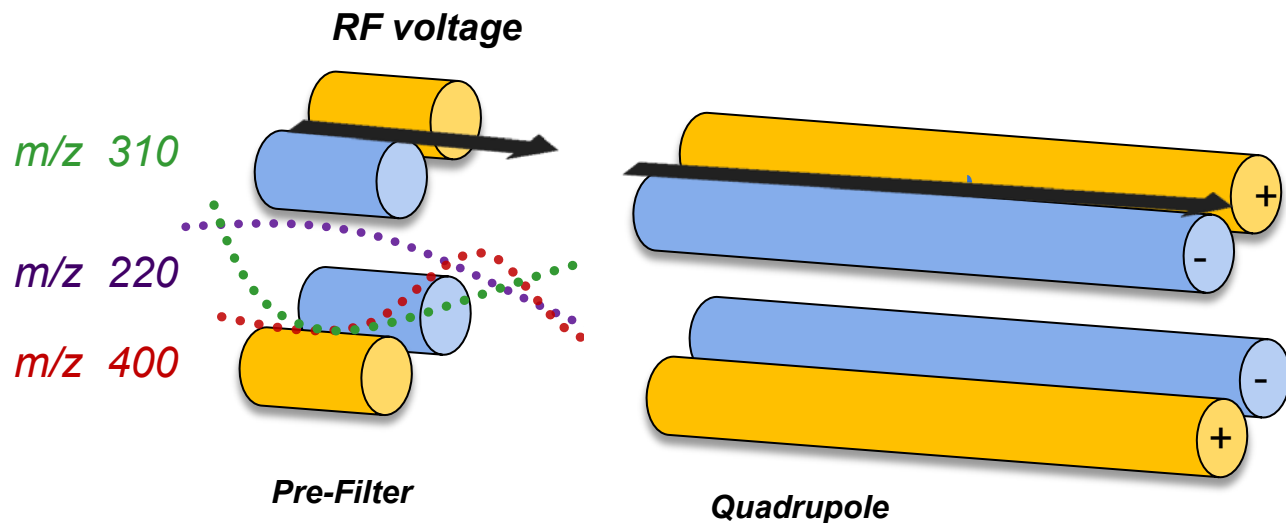


m/z explained 😊

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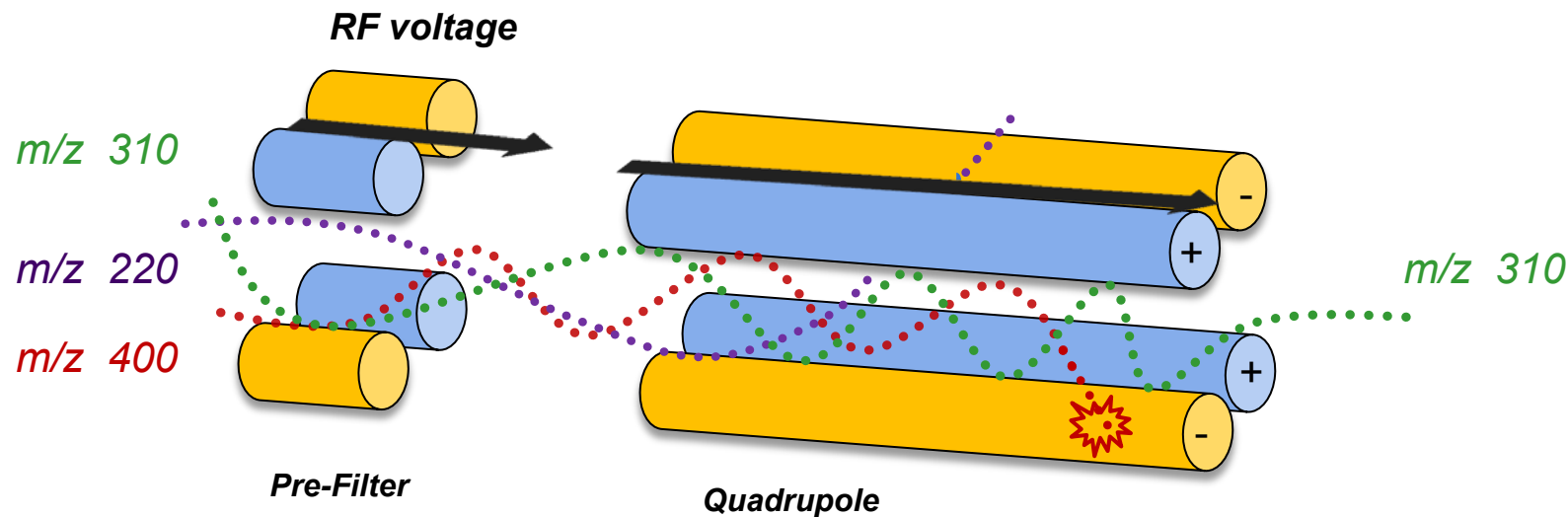


Separation or Filtering of Ions in the Quadrupole



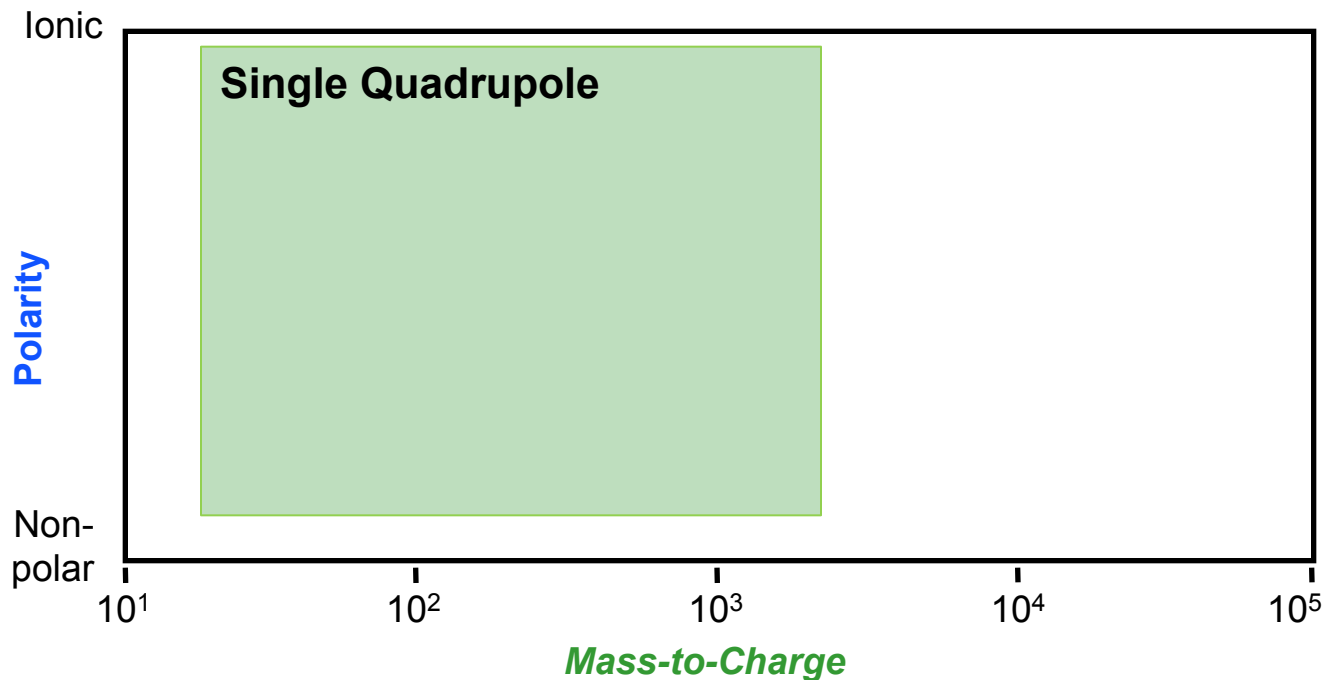
- After the sample is ionized, the sample enters the ion guides and quadrupole
- A combination of RF and DC voltages are applied to the quadrupole to create a fluctuating field
- Opposite rods have same charge applied
- The alternating field guides ions based on *mass-to-charge (m/z) ratio*

Separation or Filtering of Ions in the Quadrupole



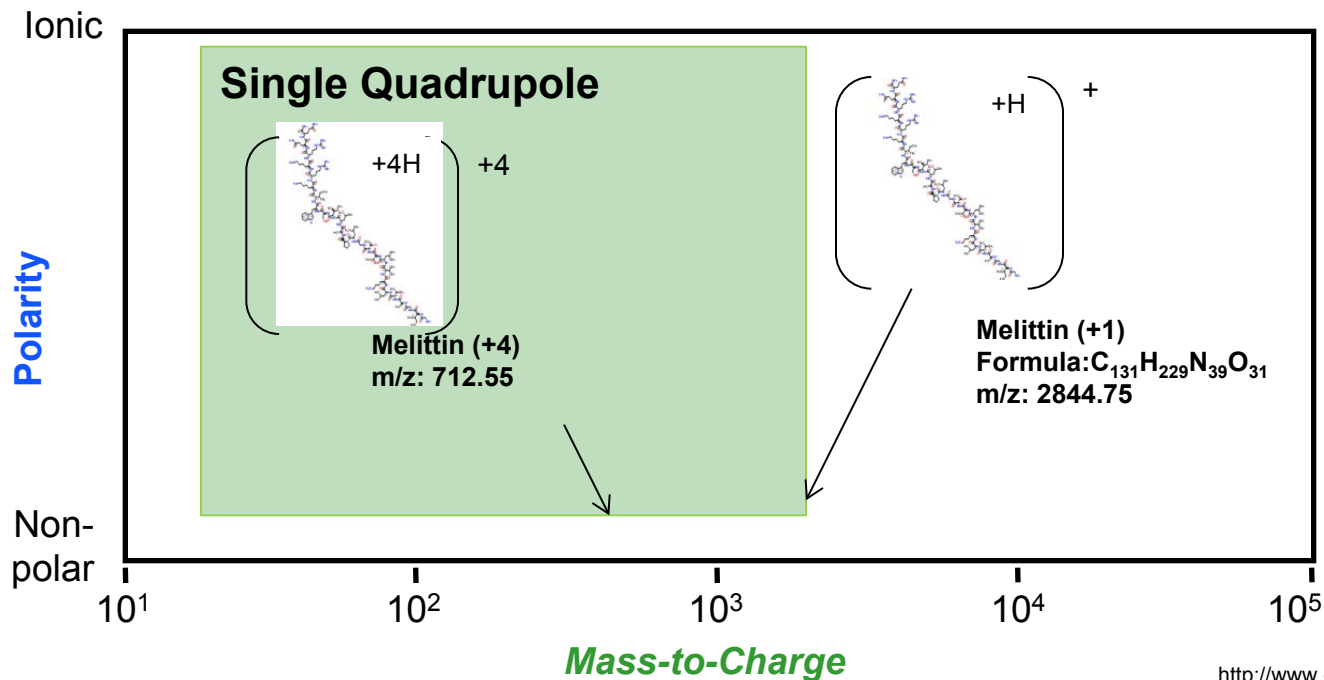
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Single Quadrupole Mass Range: Sample Properties



- Single quadrupoles typically have a maximum range of up to 2,000 - 3,000 m/z
- Higher molecular weights analytes with charged states greater than 2 might be observed on a single quadrupole, provided the charged state is less than 2,000 - 3,000 m/z

Single Quadrupole Mass Range: Sample Properties



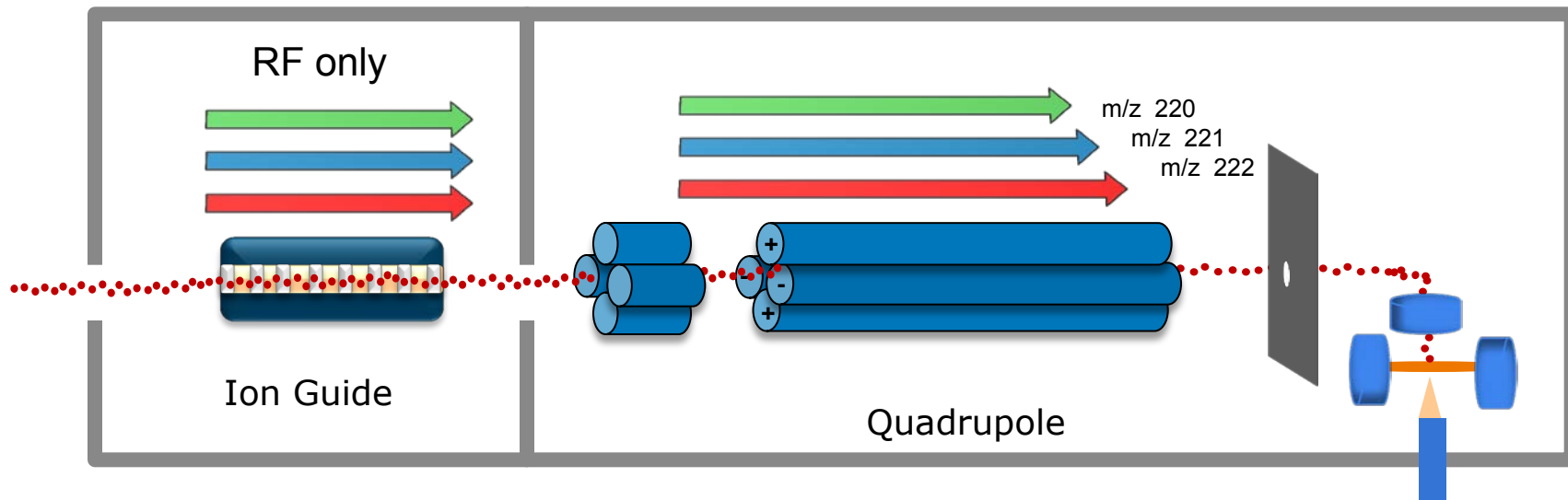
<http://www.chemspider.com/Chemical-Structure.17290230.html>

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Acquiring Single Quadrupole Mass Spectra

- Mass data can be collected in two modes:
 - Full scan mode
 - Static mode
- Full scan is analogous to photo diode array (PDA) spectra
 - Mass range unit is mass-to-charge (m/z) (e.g. 50-2000)
 - The data can be viewed in a **Total Ion Chromatogram (TIC)**
 - Individual mass-to-charge (m/z) spectra can be extracted from the TIC
- Static mode is analogous to a single 2D UV channel
 - Static mode produces a **Single Ion Recording (SIR) or Single Ion Mode (SIM)**
 - A single mass-to-charge (m/z) channel is selected

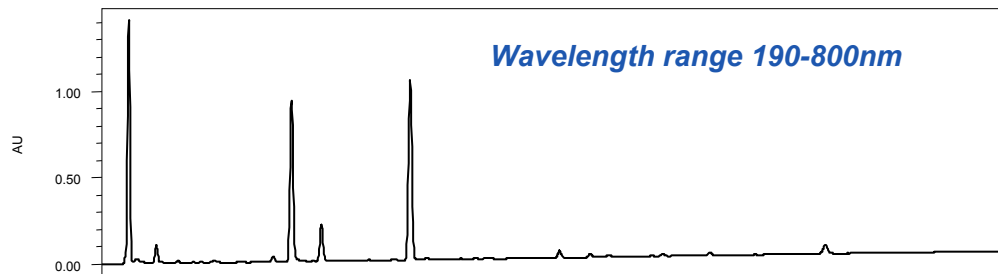
Scanning Mode



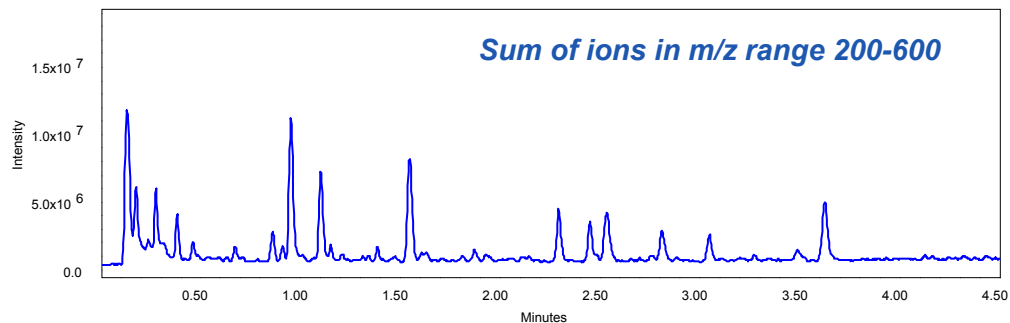
- In full scan, the RF and DC voltages are scanning transmitting ions in sequence
- The quadrupole can scan an ion in milliseconds
- More ions are transmitted in scan mode than static but the sensitivity will be lower
- Ideal for screening or scouting

Total Ion Chromatogram (TIC): Comparison to Data in UV Detection

UV 190-800 nm

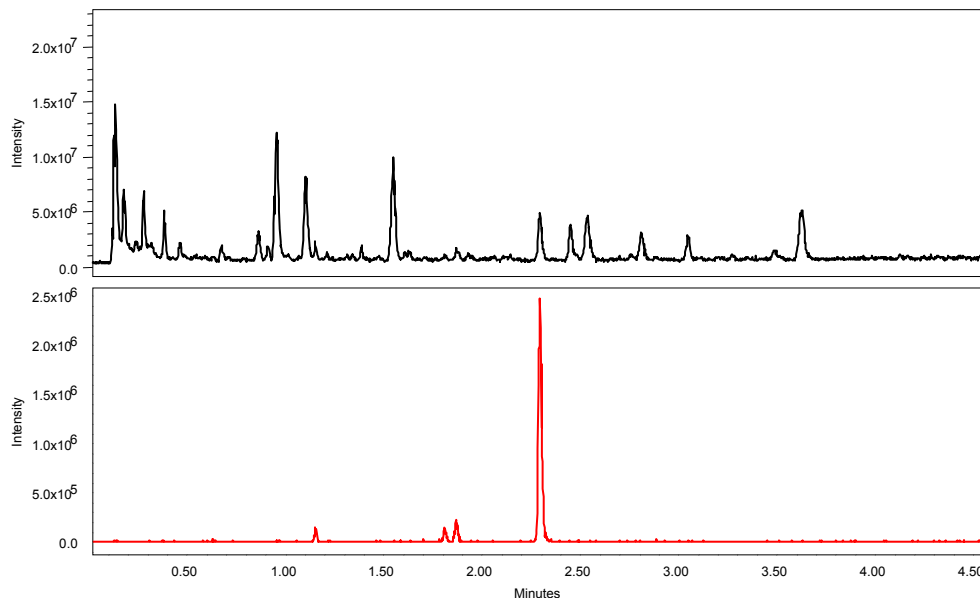


ESI+ TIC



- Relative response of analytes varies in each technique
- Y-scale equals Counts in MS vs. AU in UV
- Sample: Isoflavones in dietary supplement. Data was collected in ESI+

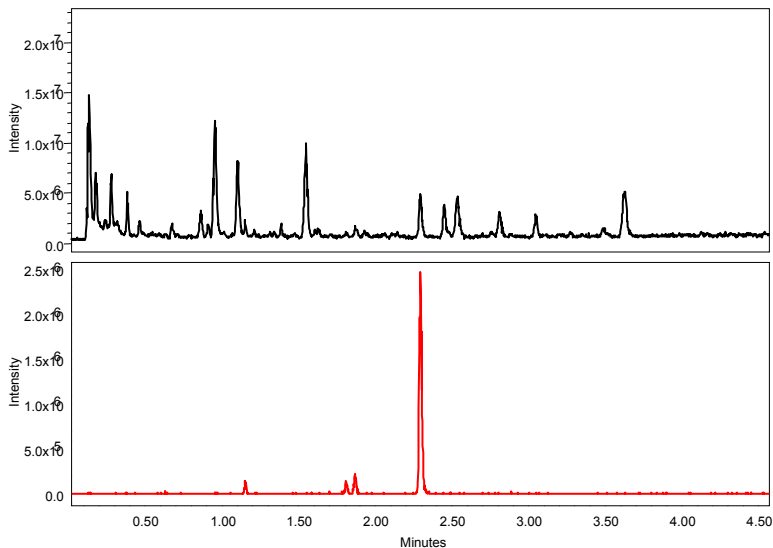
Extracted Ion Chromatogram (XIC)



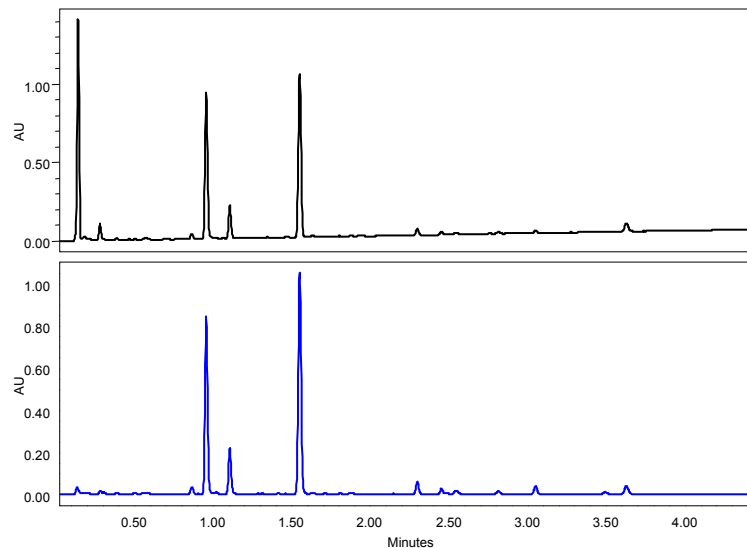
- Extracting a specific ion from the chromatogram indicates whether the m/z is present
- In this case, the m/z corresponding to 459.0 is present in multiple peaks
- Sample: Isoflavones in dietary supplement

Comparison of Extracting Single Channel in MS and PhotoDiode Array Detector (PDA)

MS

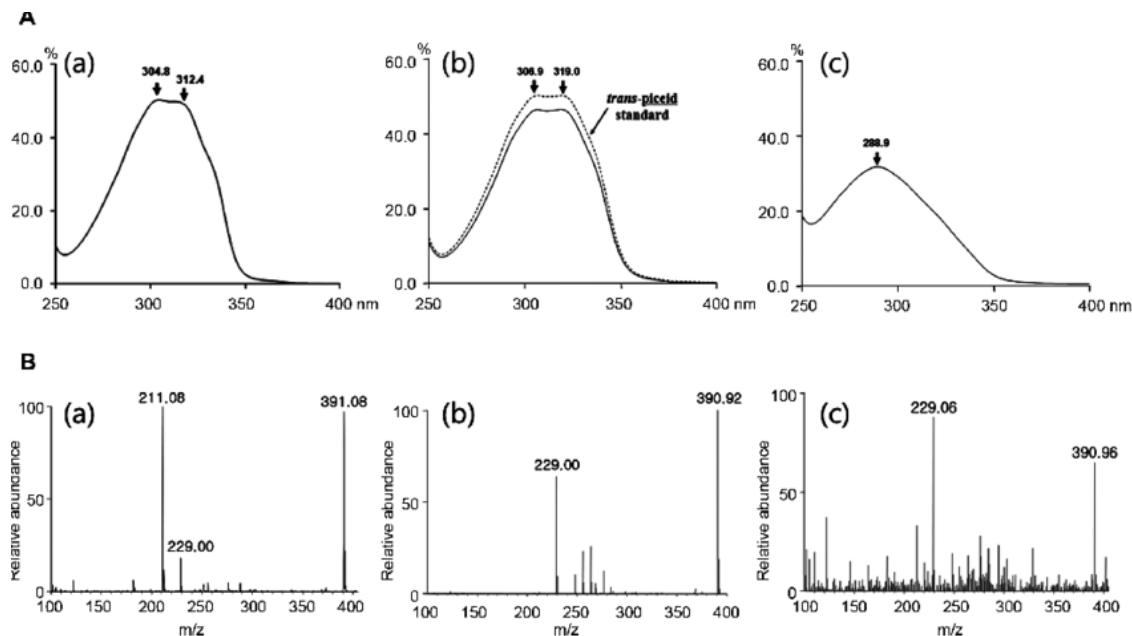


PDA



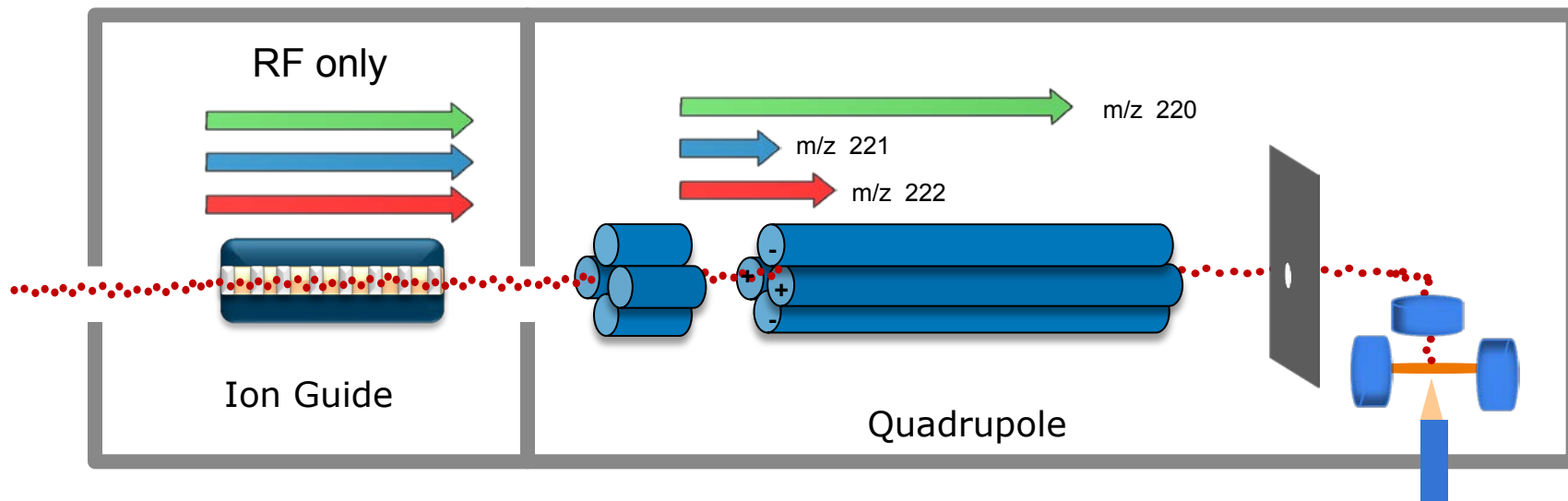
- Extracting a single mass-to-charge channel from a MS is analogous to extracting a single wavelength from a photo diode array detector
- Each may show different responses than the full TIC or full scan in PDA
- Full scan in PDA or Max Plot is maximum spectral absorbance measured at each time point

Difference UV and MS spectra



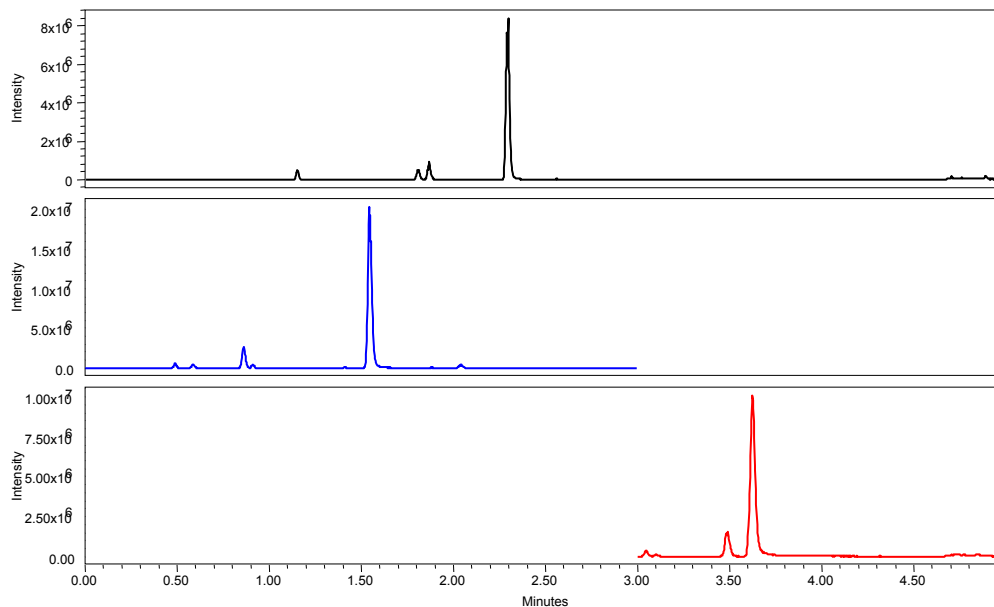
If these 3 peaks would not be chromatographically separated you could not do any quantification on the Spectral difference, With MS you could measure a) easily and b) and c) most likely. The more MS resolution the better change you can detect or measure the compounds.

Static Mode: Single Ion Recording or Single Ion Mode



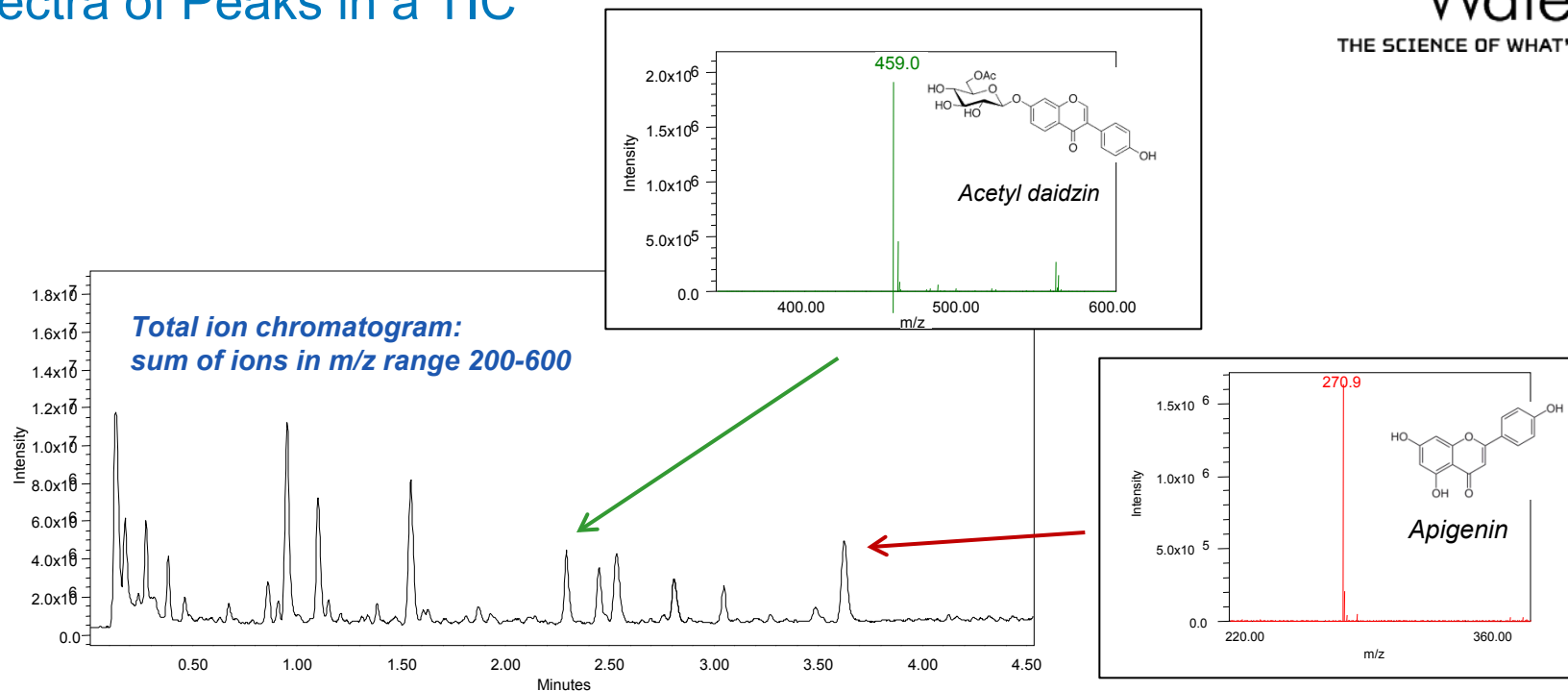
- In this example, the quadrupole settings are fixed so only a single mass-to-charge (m/z) ratio is transmitted
- Highest sensitivity because entire dwell time is spent on single m/z
- *Most common for quantitative analysis*

Single Ion Recording (SIR) Channels



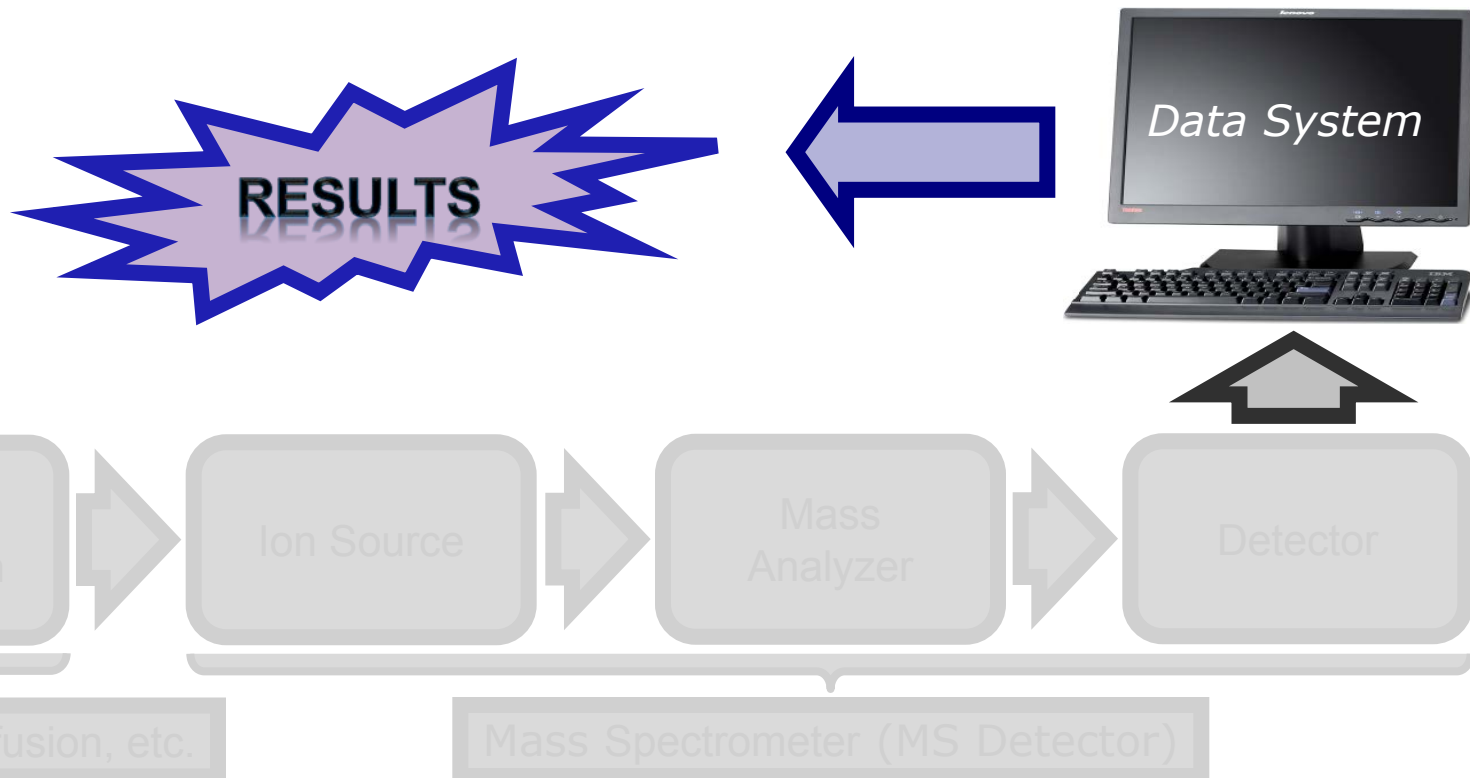
- In static mode a single ion recording channel is collected
- Each single ion recording channel represents a specific mass-to-charge ratio
- Multiple single channels can be collected in a single run
- Single channels can be collected over the entire analysis or for a specific amount of time

Spectra of Peaks in a TIC



- Each peak in the TIC has a spectrum associated with it
- The spectrum includes the observed mass-to-charge ratios present for that peak

What Impacts Results?



Ionized Molecules and Adducts

Positively charged adducts

Adduct Ion Formed	m/z of ion
M+H	M + 1.00
M+NH ₄	M + 18.03
M+Na	M + 22.99
M+CH ₃ OH+H	M + 33.03
M+K	M + 38.96
M+ACN+H	M + 42.03
M+2Na-H	M + 44.97
M+IsoProp+H	M + 61.06
M+ACN+Na	M + 64.02

Negatively charged adducts

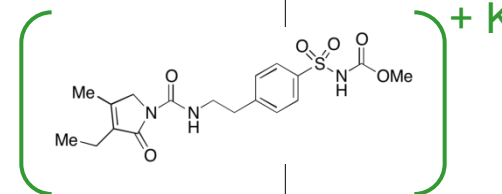
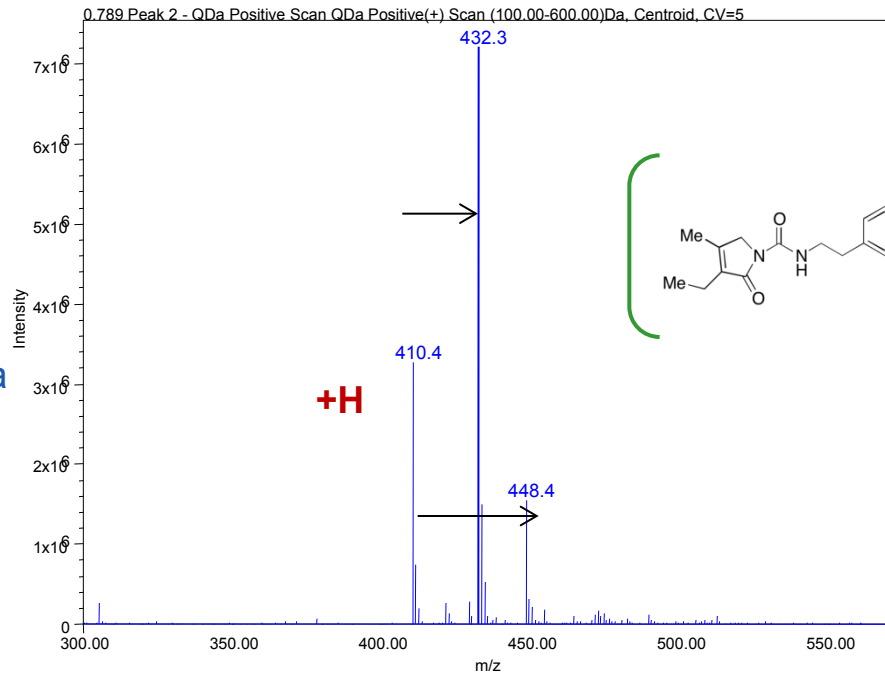
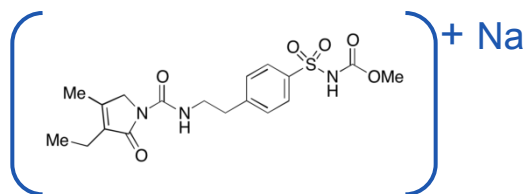
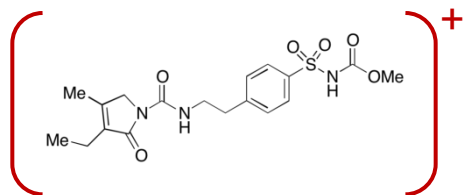
Adduct Ion Formed	m/z of ion
M-H	M - 1.00
M+Na-2H	M + 20.97
M+Cl	M + 34.97
M+K-2H	M + 36.95
M+FA-H	M + 45.00
M+Br	M + 78.92
M+TFA-H	M + 112.99

- REF <http://fiehnlab.ucdavis.edu/staff/kind/Metabolomics/MS-Adduct-Calculator>

Adducts Observed in MS

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- Some compounds are more likely to form adducts other than H⁺
- Common adducts observed in positive mode include Na and K adducts

Glimepiride related compound C
Formula: C₁₈H₂₃N₃O₆S
MW:409.46

Multiple Charging in Electrospray Ionization

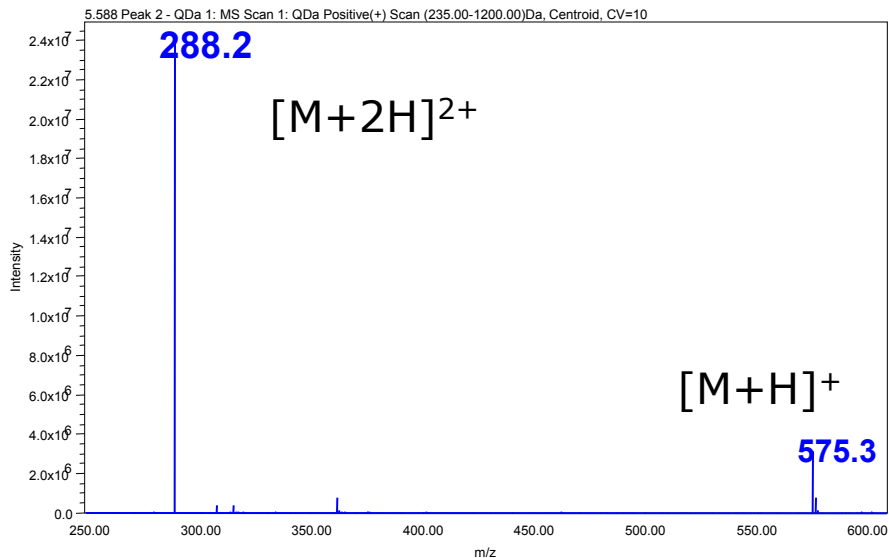
- Mass spectrometers separate ions on the basis of mass-to-charge ratio (m/z)
 - Singly charged ion: $(M + H)^+$ $m/z = (M + H)/1$
 - Doubly charged ion: $(M + 2H)^{2+}$ $m/z = (M + H)/2$
 - n charged ion: $(M + nH)^{n+}$ $m/z = (M + nH)/n$

- Isotope peaks of an ion with n charges are separated by $1/n$ m/z
 - e.g. isotope peaks of a doubly charged ion would be separated by 0.5 m/z

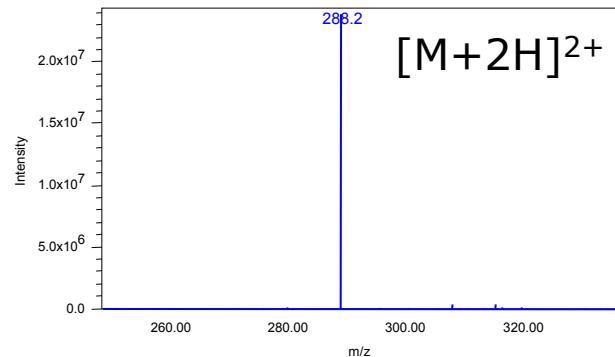
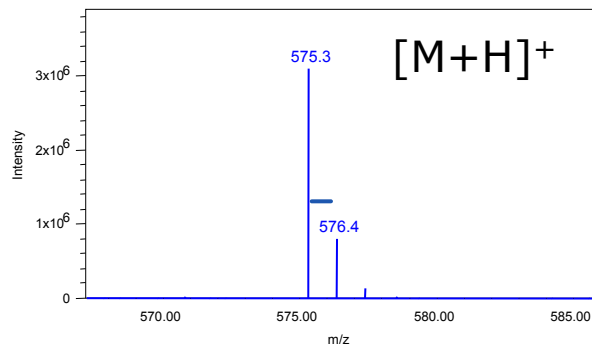
- Multiple charging of an analyte molecule may happen when more than one location on a molecule can accept a charge

- Multiple charge states are more likely observed with analytes of higher mass (peptides, proteins, etc.)

Distinguishing Between Multiply Charged Peaks

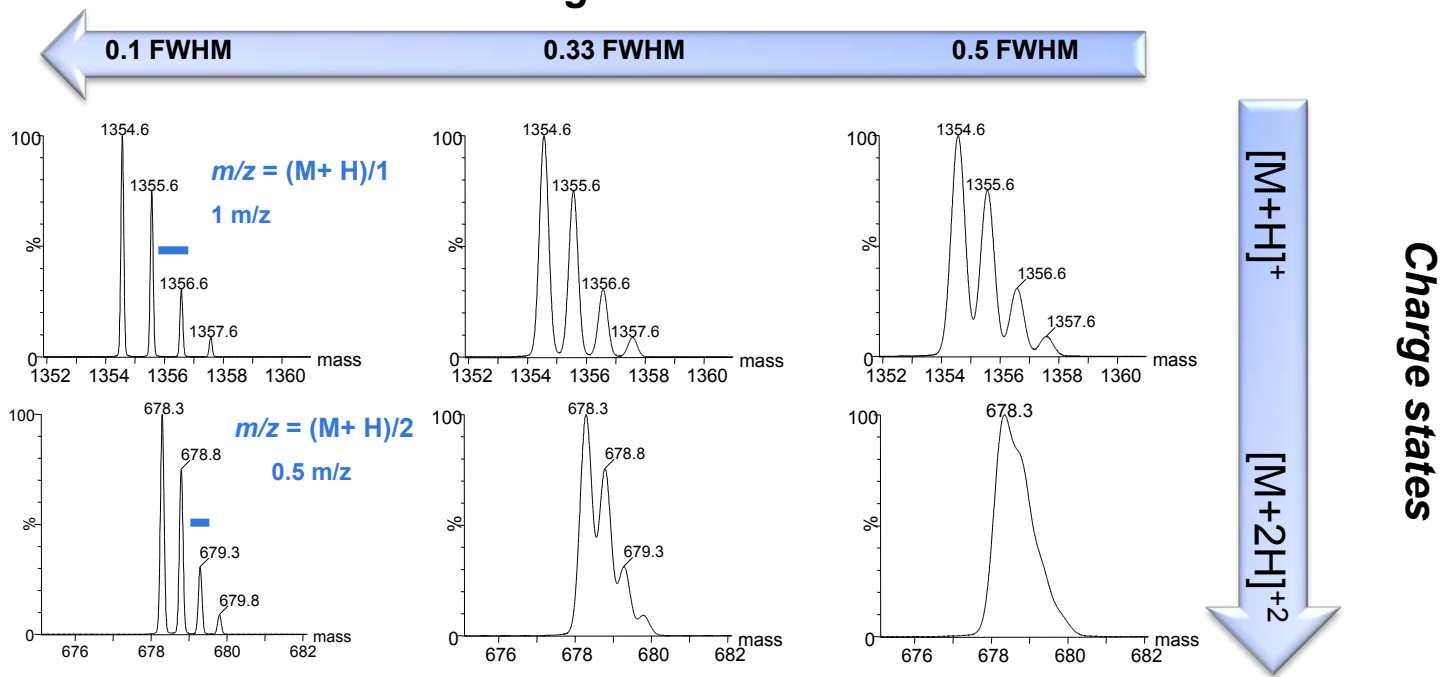


- NIST MAb peptide
 - Peptide⁺ – 574.3
 - Single charged: $(574.3+1)/1=575.3$
 - Doubly charged: $(574.3+2)/2 = 288.2$



Impact of Instrument Resolution on Charge State

increasing resolution



- Simulation of Vitamin B¹² isotope models for singly and doubly charged isotopes
- Singly charged species separated by 1 m/z, doubly charged species isotopes separated by 0.5 m/z

Questions??