

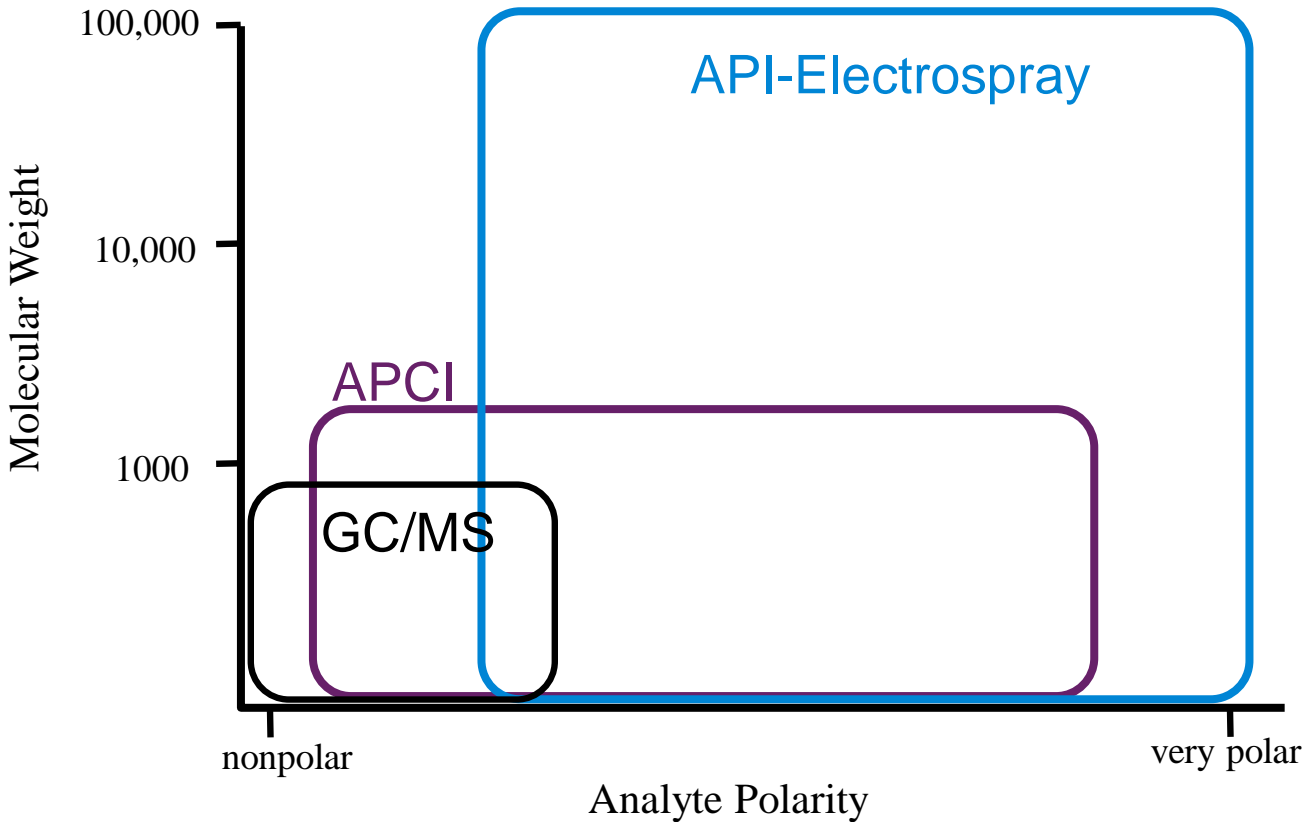


# Innovative Screening Workflows in Liquid Phase Mass Spectrometry

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Dr Shaun Bilsborough

# Relative Applicability of LC/MS Techniques






# Guiding Principles in MS Analyzer Selection

The correct choice of analyzer depends on whether one is “LOOKING for **UNKNOWN**S” or whether one is “CONFIRMING & MEASURING **KNOWN**S”.

The correct choice of analyzer depends on the **COMPLEXITY** of the **MATRIX**.



# Screening Workflow Solutions

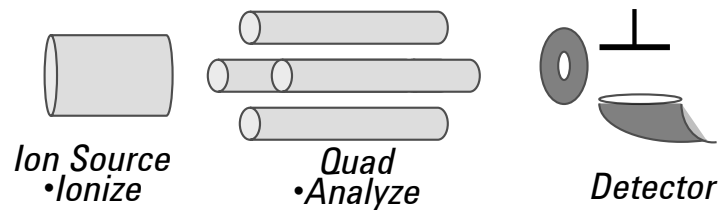
Screening	Target	Suspect	Discovery
SQ or QQQ	 <p>Using SIM, MRM and tMRM databases, method can be quickly developed for implementation. Standards required for QQQ method development and quantitation</p>		
TOF	<p>Using PCDLs and an All-Ions workflow Quantitation and Qualification is possible using four points for confirmation (Rt, AccMass, Fragments, IP).</p>	<p>Suspect Screening requires the use of Find by Formula referring to a suspect PCDL where MS/MS spectra are available.</p>	
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# Single Quad (SQ)

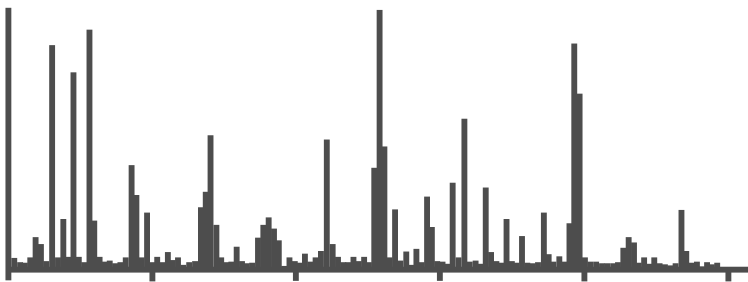
Selected Ion Monitoring selects a single mass

Allows sensitive, selective quantitation in relatively clean samples

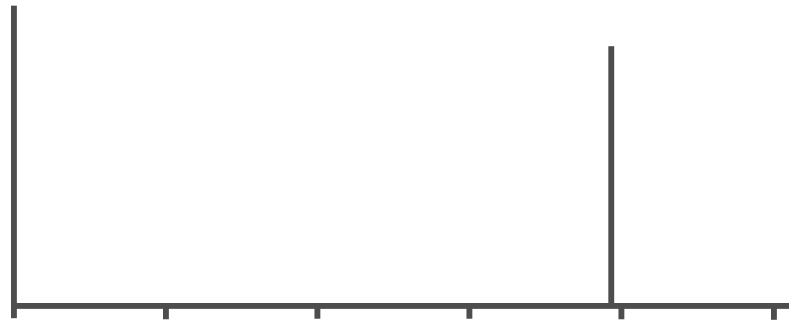
*a single quad instrument...*



## Full Scan

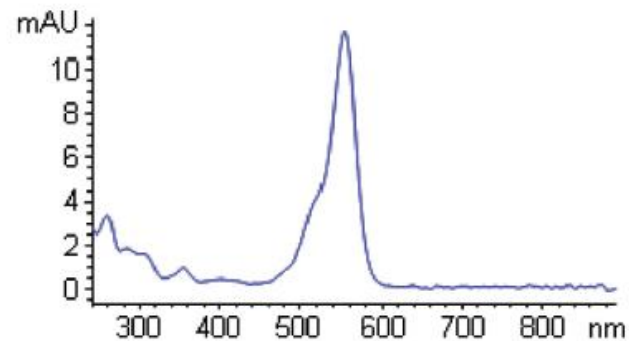
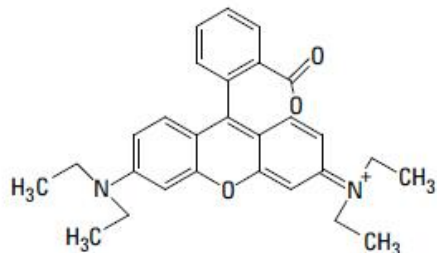


## Selected Ion Monitoring (SIM)



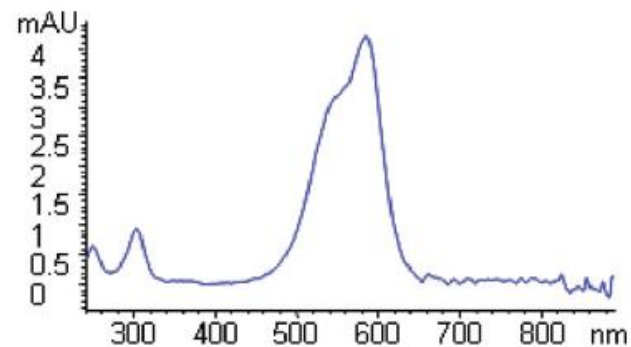
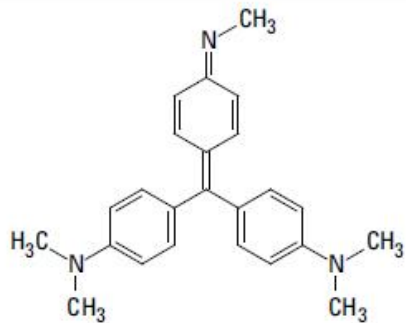
# LC + DAD

Rhodamine B (RB)



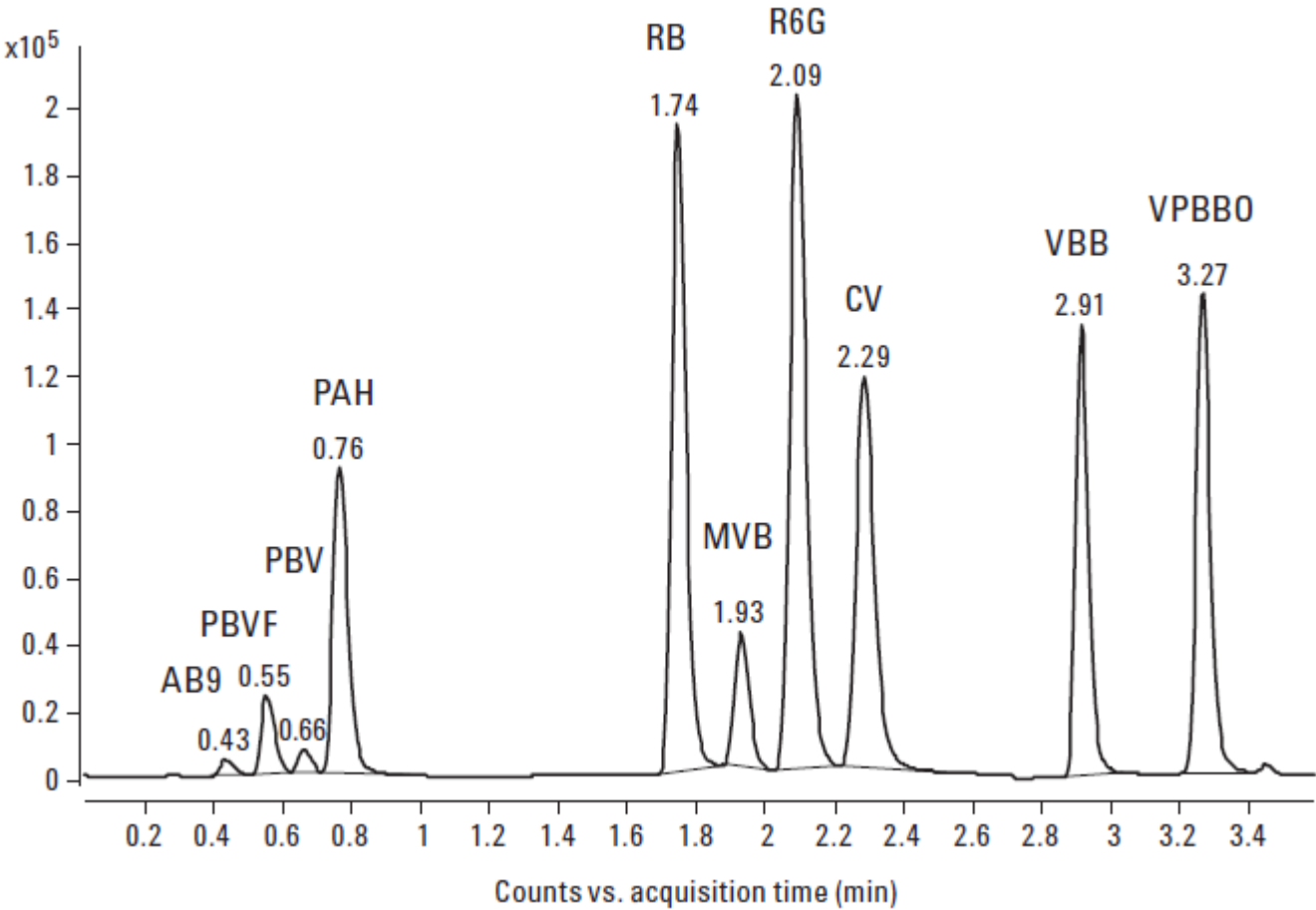
λ max: 554

Methyl violet B base (MVB)






λ max: 580

# LC/SQ MS + DAD



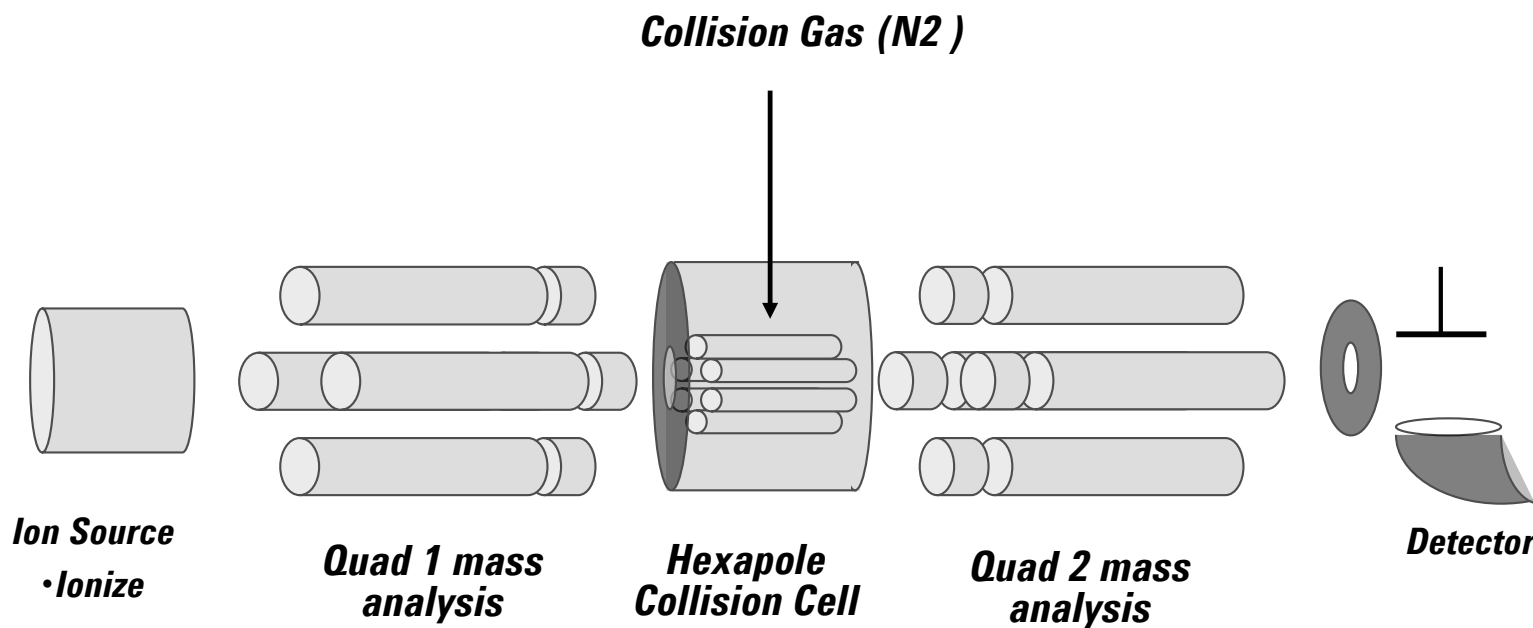
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# Triple Quad (QQQ)

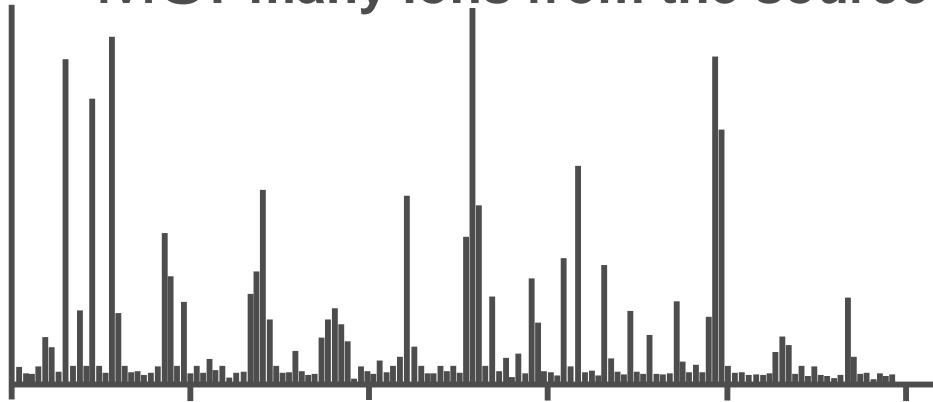
Quad – Hex - Quad



# What is MRM MS/MS using QQQ?

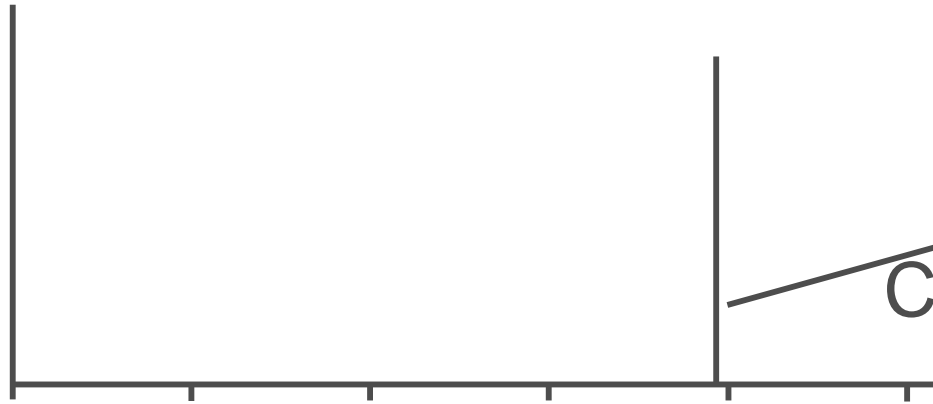
Multiple Reaction Monitoring

MS: many ions from the source



Q1 SIM

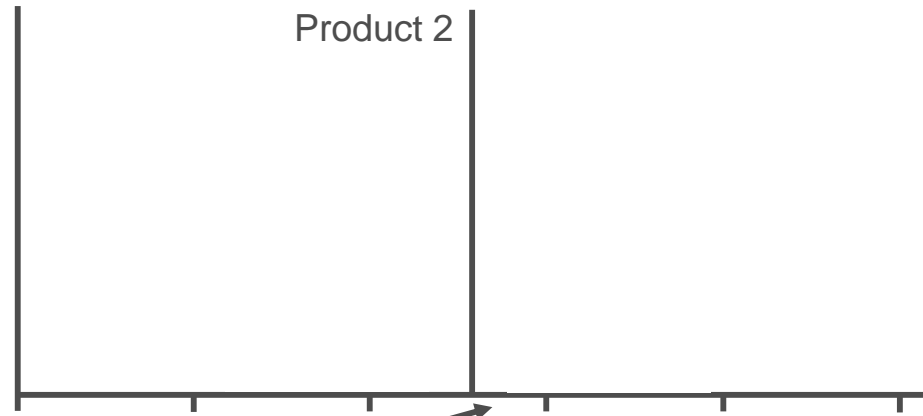
isolate precursor  
before CID



CID + Q2 SIM

Q3: SIM

Product 2



# Why MS/MS?

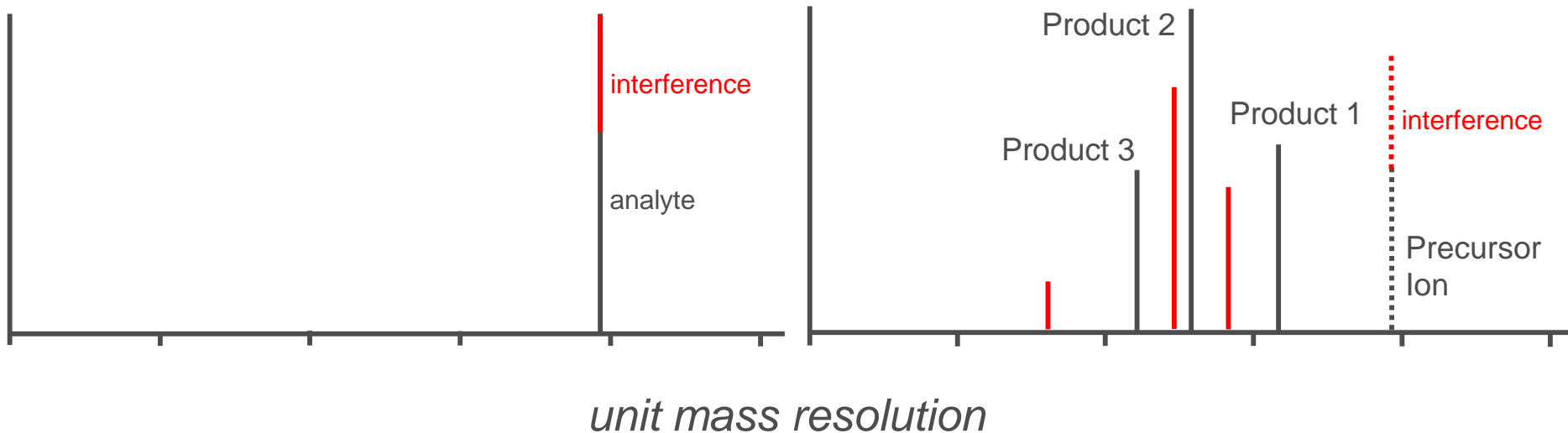
## Greater Selectivity Than SIM

### SIM (Single Quad)

- selectivity proportional to spectral resolution
- no selectivity against ions with same  $m/z$

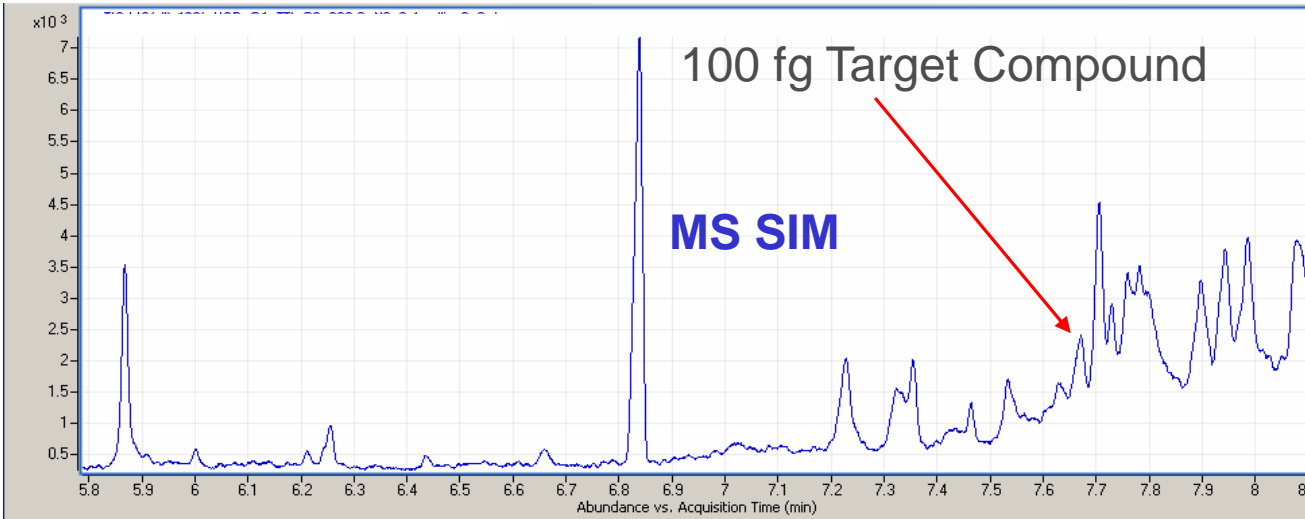
### MS/MS (QQQ)

- precursor selectivity same as SIM
- high probability that at least one product ion will be a unique fragment of the precursor BUT not the interference

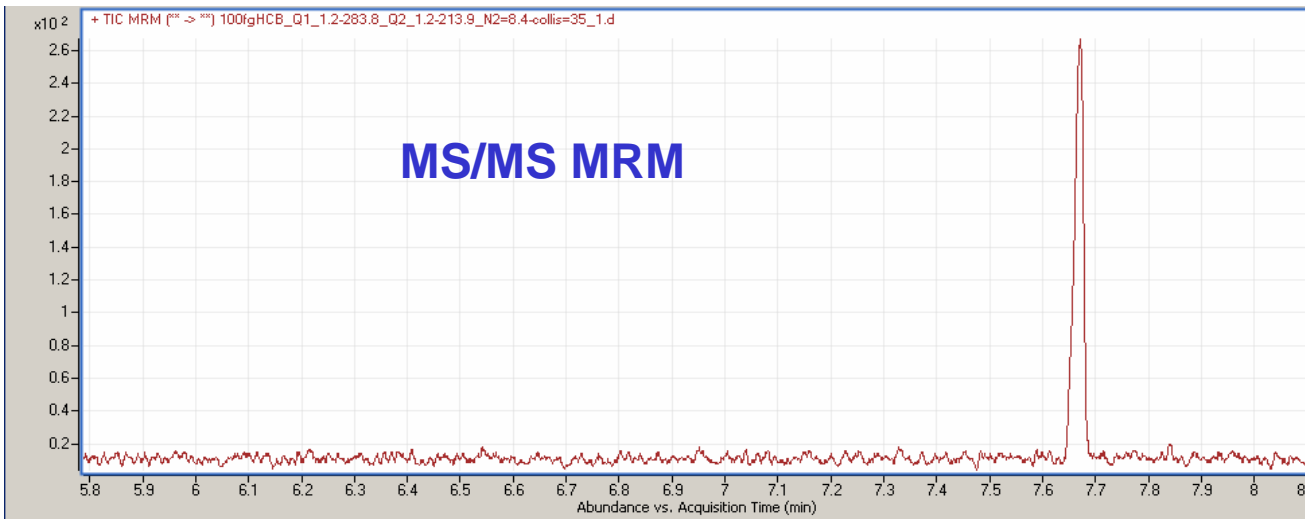


# Why MS/MS?

A Picture Is Worth a Thousand Words






**Single Quad SIM of  
single m/z**



**MS/MS QQQ MRM  
of single transition**

*Single peak on flat  
baseline*

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# TOF and QTOF

## Performance Summary and Overview

Q-TOF

[RETURN TO OVERVIEW](#)

Q-TOF

TOF

[Sensitivity](#)

[Mass Accuracy](#)

[Dynamic Range](#)

[Speed](#)

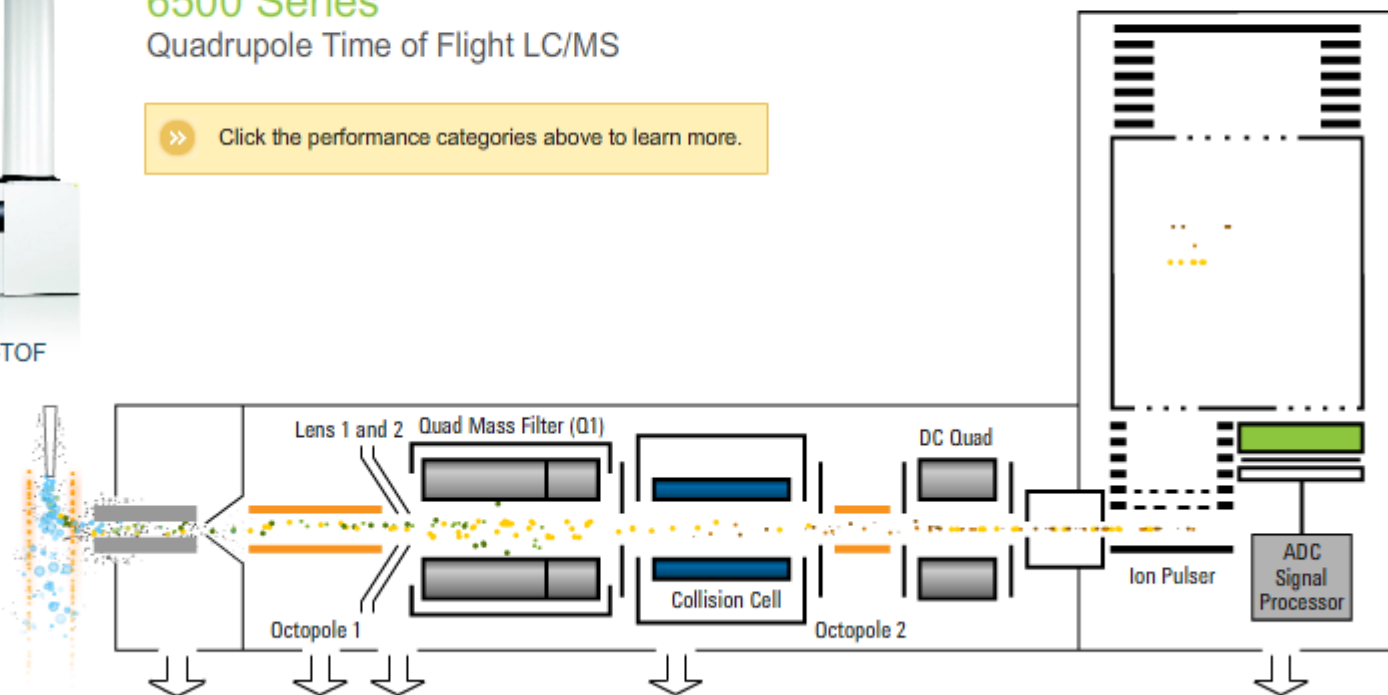
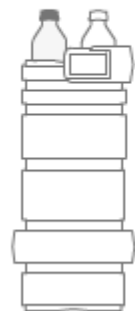
[Resolution](#)



### 6500 Series

Quadrupole Time of Flight LC/MS

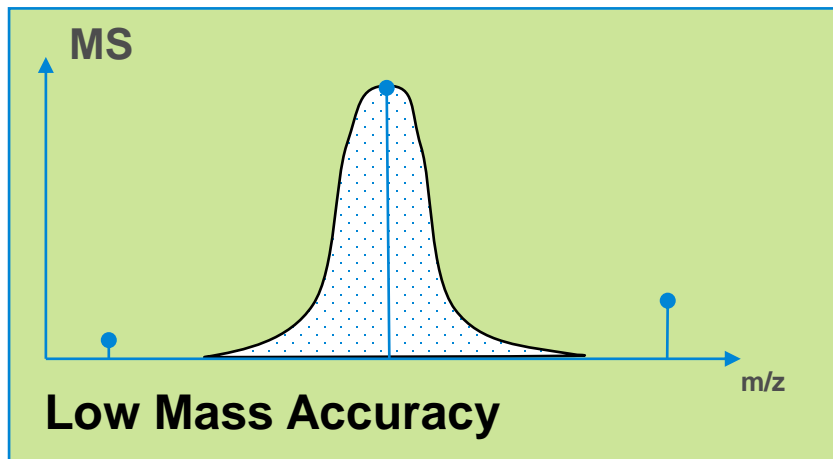
» Click the performance categories above to learn more.



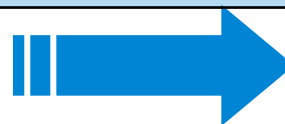
Agilent Technologies



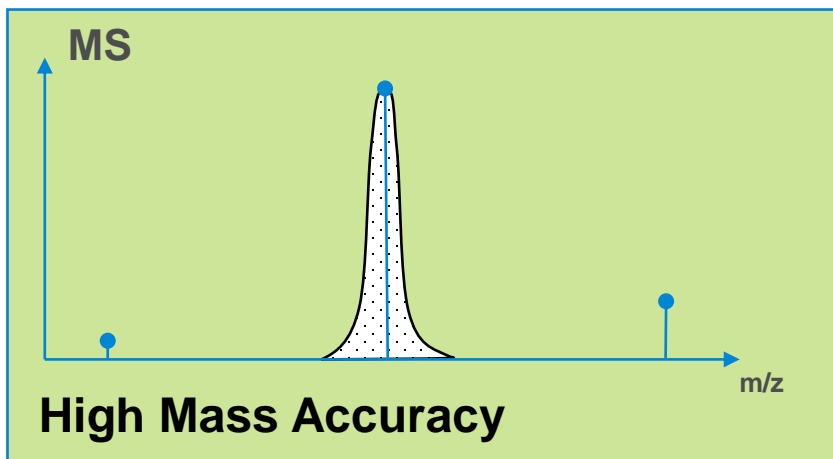
# Why Accurate Mass?



Unit accuracy instrument  
e.g. ion trap, QQQ, Q



C12 H14 N9 Cl  
C13 H23 N3 P S Cl  
**C14 H24 N2 S2 Cl**  
C14 H22 N O3 S Cl  
C14 H16 N6 O Cl  
C15 H25 O P S Cl  
C16 H19 N3 P Cl  
C17 H20 N2 S Cl  
C18 H21 O P Cl



High mass accuracy  
instrument  
e.g. QTOF, TOF



C13 H23 N3 P S Cl  
**C14 H24 N2 S2 Cl**  
C14 H22 N O3 S Cl  
C14 H16 N6 O Cl  
C15 H25 O P S Cl  
**C16 H19 N3 P Cl**  
C17 H20 N2 S Cl  
C18 H21 O P Cl

# Molecular Formula Generation (MFG) Algorithm Uses All Available Information

Scoring based on

**Method Editor: Generate Formulas**

Method Items

Allowed Species Limits Charge State Scoring

Contribution to overall score

Mass score: 100.00

Isotope abundance score: 60.00

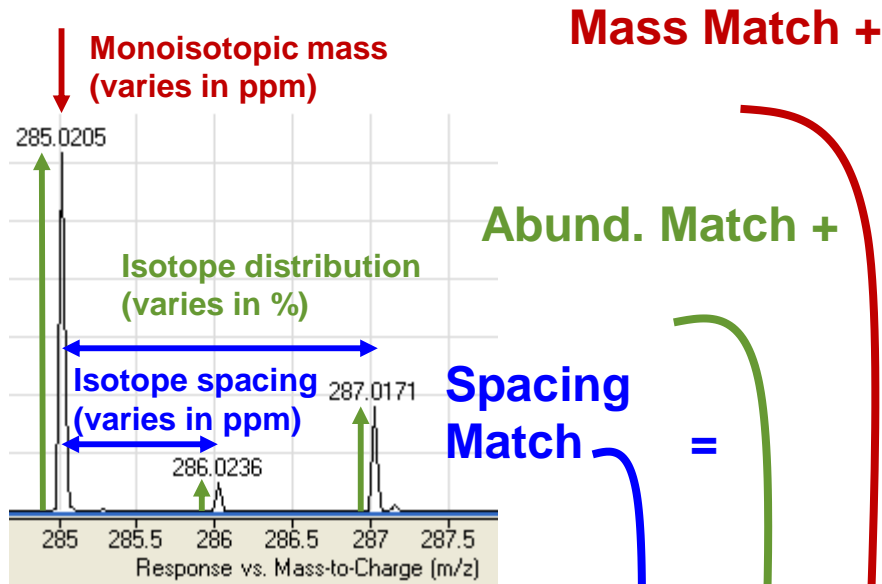
Isotope spacing score: 50.00

Expected data variation

MS mass: 2.0 mDa + 5.6 ppm

MS isotope abundance: 7.5 %

MS/MS mass: 5.0 mDa + 7.5 ppm



Overall Score

**MS Formula Results: Cpd 2: C<sub>10</sub>H<sub>9</sub>ClN<sub>4</sub>O<sub>2</sub>S**

m/z	Ion	Formula	Abundance
285.021	(M+H) <sup>+</sup>	C <sub>10</sub> H <sub>10</sub> ClN <sub>4</sub> O <sub>2</sub> S	24506.1

Best	Formula (M)	Calc m/z	Score	Cross Score	Mass	Calc Mass	Diff (ppm)	Abs Diff (p)	Spacing Matc	Abund Matc	Mass Match	m/z	DBE
<input checked="" type="checkbox"/>	C <sub>10</sub> H <sub>9</sub> ClN <sub>4</sub> O <sub>2</sub> S	285.0208	99.55		284.0137	284.0135	-0.71	0.71	99.19	99.26	99.69	285.021	8
<input type="checkbox"/>	C <sub>7</sub> H <sub>12</sub> N <sub>2</sub> O <sub>6</sub> S <sub>2</sub>	285.021	77.28		284.0137	284.0137	0.01	0.01	99.54	1.93	100	285.021	3
<input type="checkbox"/>	C <sub>7</sub> H <sub>13</sub> ClN <sub>4</sub> O <sub>2</sub> S <sub>2</sub>	285.0241	75.57		284.0137	284.0168	11.12	11.12	99.87	83.87	46.22	285.021	3






# PCDLs by Compounds & Spectra

Accurate Mass (AM) LC/MS PCDL	Market	Compounds	Compounds with MS/MS Spectra	Total number of Spectra
Extractables and Leachables	Pharma, BioPharma, Food Safety	>1,000	>350	>1300
Broecker, Herre & Pragst Forensic Toxicology	Forensic Toxicology	>9,200	>3,900	>13,500
Pesticides	Food Safety / Environmental	1,750	>825	>2,700
Veterinary Drugs	Food Safety	>2,150	>1,525	>5,200
Mycotoxins	Food Safety	>450	>300	>1,350
Water Contaminants	Environmental	>1,400	>1,050	~3,900
METLIN	Metabolomics / Lipidomics	>79,600*	>9,450	>32,000

\* Plus 168k theoretical

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