



**Development of
High Performance
LC-QIT-TOF**

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- The Need for Highly Precise Mass Spectrometry
- Objective of LC-QIT-TOF
- Technology Developed for LC-QIT-TOF
- Fundamental Performance Data

The Need for Highly Precise Mass Spectrometry

- Precision mass spectrometry allows inferring the elemental composition of molecules
- The higher the precision, the more the **number of constituent candidates can be reduced**

Example: (reserpine) The number of elemental combination candidates listed for a mass of $(M+H)^+=609.2822$

Constituent Elements:

Carbon	20 - 40	Hydrogen	30 - 60
Nitrogen	3 - 10	Oxygen	5 - 15
Sulfur	0 - 5		

Precision	10 ppm	5 ppm	2 ppm
Number of Combinations	29	16	4

Objective of LC-QIT-TOF

- Ion Trap MS

Allows MS^n

High precursor resolution

- TOF MS

Highly precise mass values

High resolution

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graph TD; A["• Ion Trap MS<br/>Allows MS<sup>n</sup><br/>High precursor resolution"] --- B["• TOF MS<br/>Highly precise mass values<br/>High resolution"]; B --- C["QIT-TOF<br/>An instrument that allows high<br/>precision MS<sup>n</sup> spectrometry"]; style C fill:#ffff00,stroke:#ffff00
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QIT-TOF

An instrument that allows high
precision MS^n spectrometry

Technologies Developed for LC-QIT-TOF



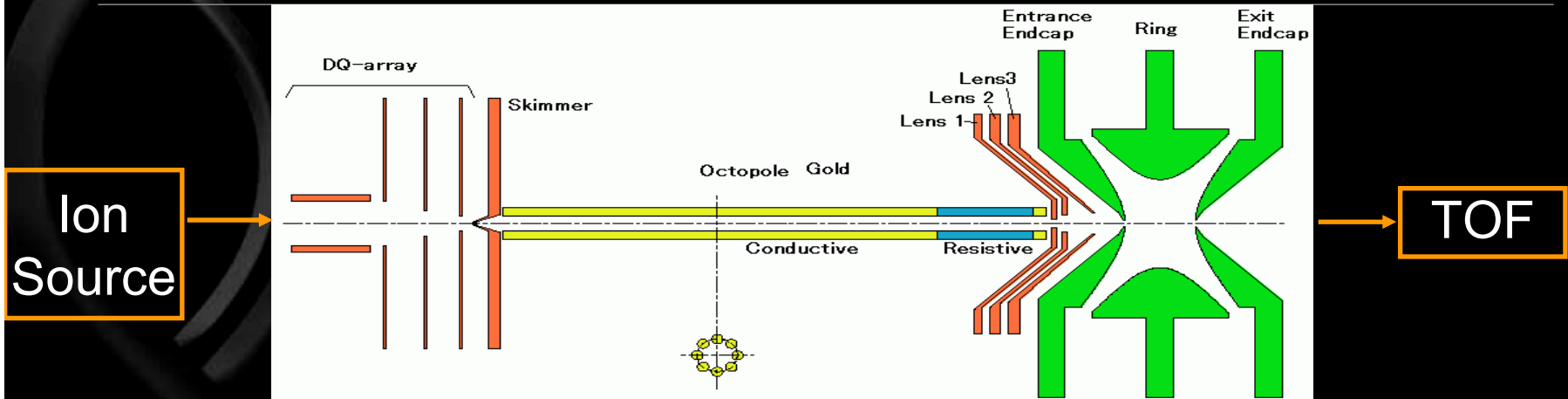
1685 x 685 x 570 mm, 280 kg

Differences Between MALDI-QIT-TOF and ESI-QIT-TOF

Key newly developed technologies:

1. **Compressed ion introduction** (introduction to QIT)
MALDI: Pulsed ionization ESI: Continuous ionization
2. **Improvements to Dual Stage Reflectron**
(High sensitivity and high resolution for QIT-TOF)
3. **Ionic cooling using argon gas**
(High resolution)
4. **Temperature controlled interior of instrument**
(Stability of mass precision)
5. **Ballistic Ion Ejection**
(Accuracy and stability of mass measurements)

Compressed Ion Introduction

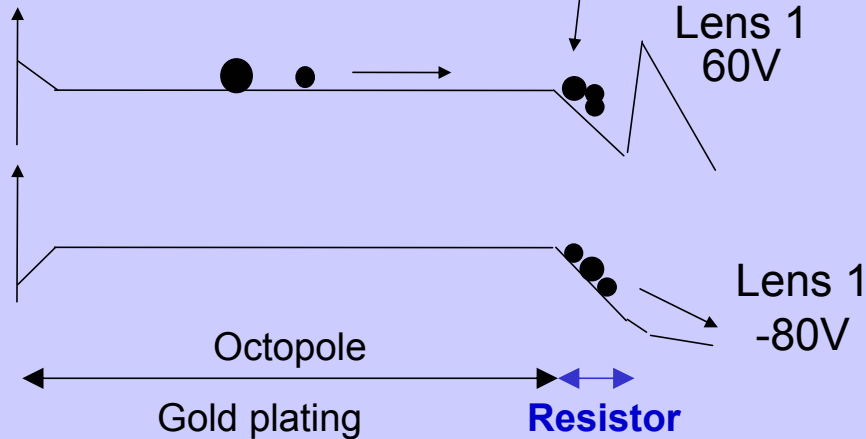


DC potential along central axis

Accumulation of ions to OP

Introduction to QIT

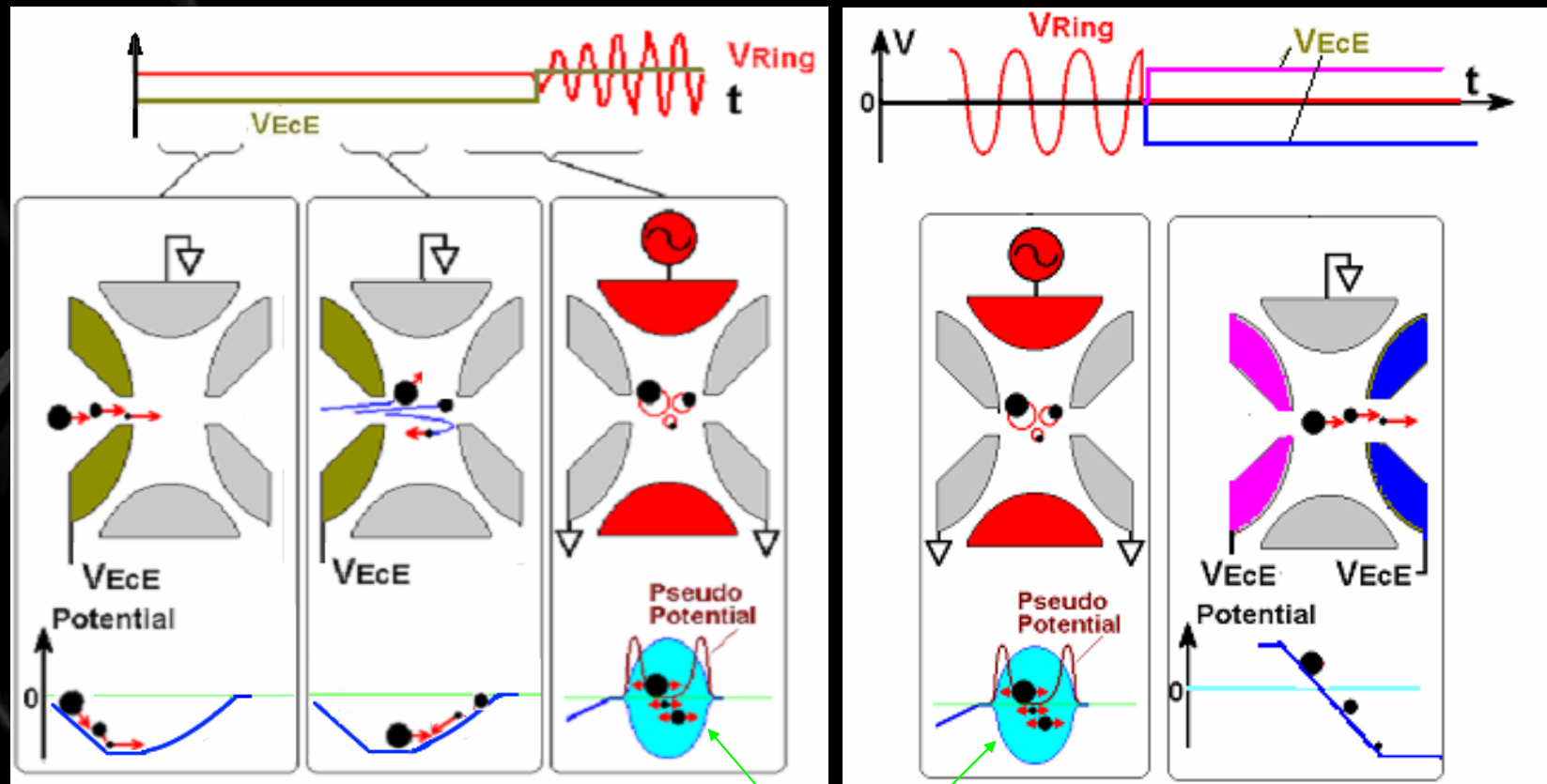
Ions collected at exit



Introducing Ions into the Trap and Discharging Them

Delivery

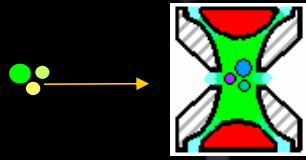
Discharge



Ion trap and localization

Basic Process of QIT-TOF

Ion Pulse



Trap

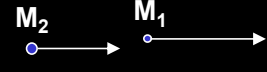
Ions are trapped and cooled



MS



Mass spectrometry



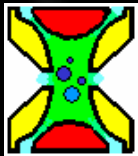
Detector

MS/MS



Isolation

Target ions are selected
Other ions are discarded



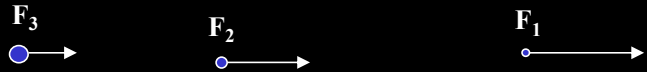
Fragmentation

(CID gas is injected and ions are collided with gas molecules)



Mass spectrometry of fragment ions = MS/MS

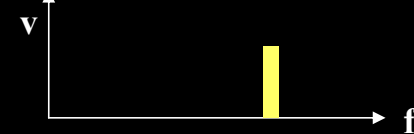
Detector



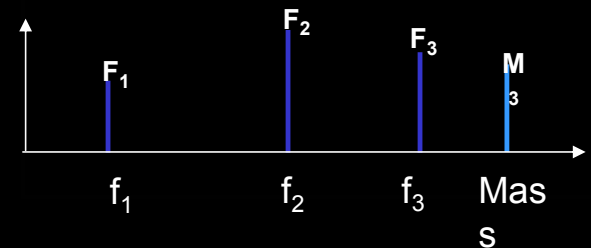
Endcap frequency spectrum 1



Endcap frequency spectrum 2



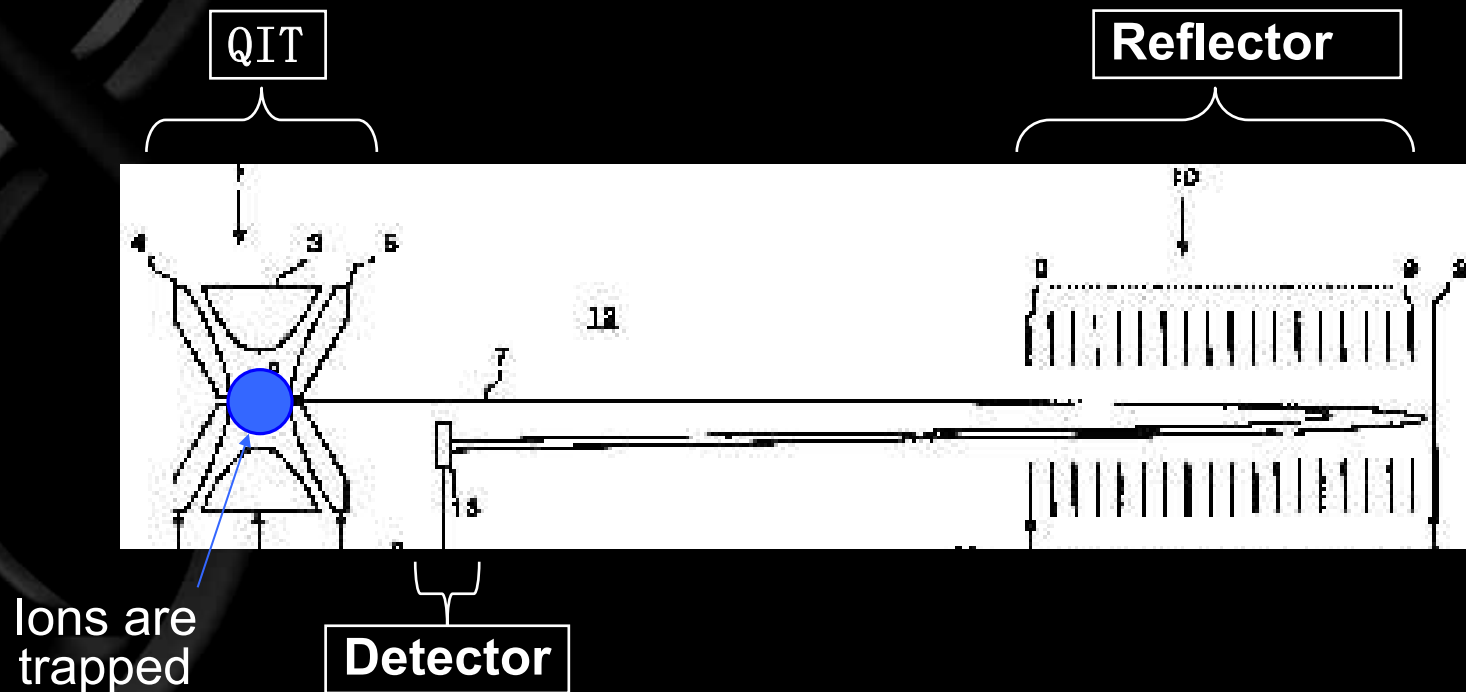
MS/MS Spectrum



Dual Stage Reflectron Improvements

The shape of the electric field was changed for second stage of the dual stage reflectron.

-> Improves the time convergence with respect to ion distribution during acceleration



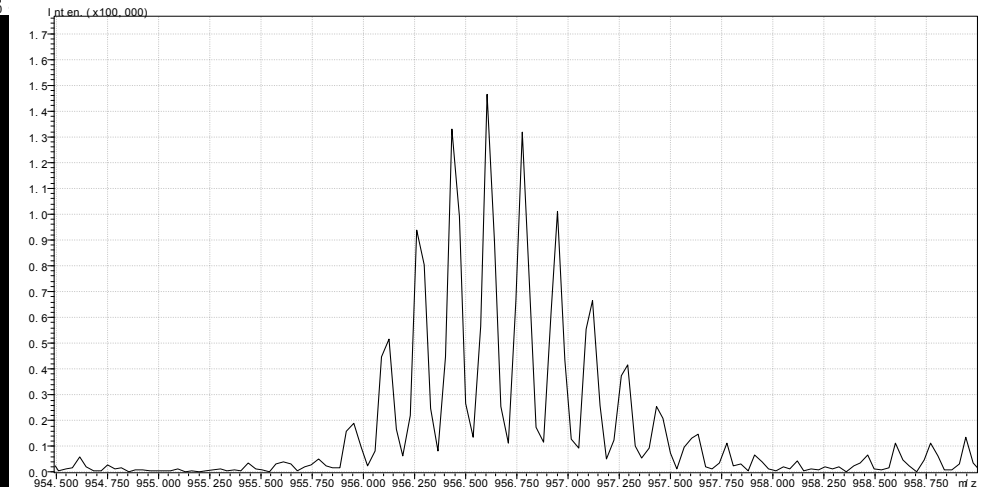
Cooling with Argon Gas

Bovine Insulin m/z 956 $[M+6H]^{6+}$



He: 4.5×10^{-2} Pa

Ar: 7.5×10^{-3} Pa



Controlling the Instrument's Internal Temperature to Stabilize Mass Precision

Temperature Control

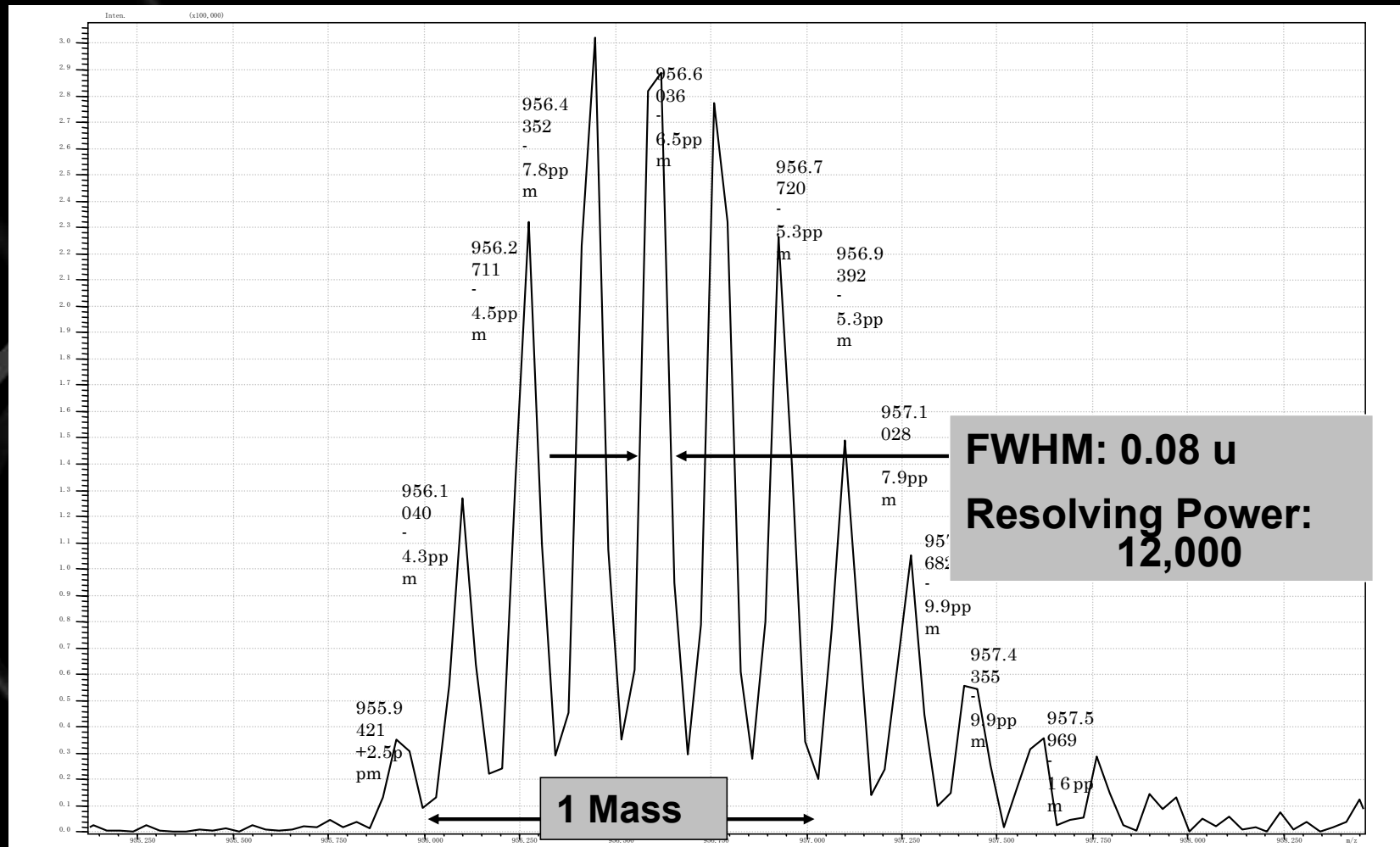
- Flight Tube (Representative temperature point)
- High Voltage Switch (Ion Trap EC, Ring)
- High Voltage Power Supply
(Ion Trap, Flight Tube, Reflectron)
- RF Generator (Ion Trap)

Controls representative temperature point to
 $40 \pm 0.3^{\circ}\text{C}$

A pair of glasses is shown on the left side of a black background. A thin white horizontal line is positioned near the top of the image. The text "Fundamental Performance Data" is centered in white.

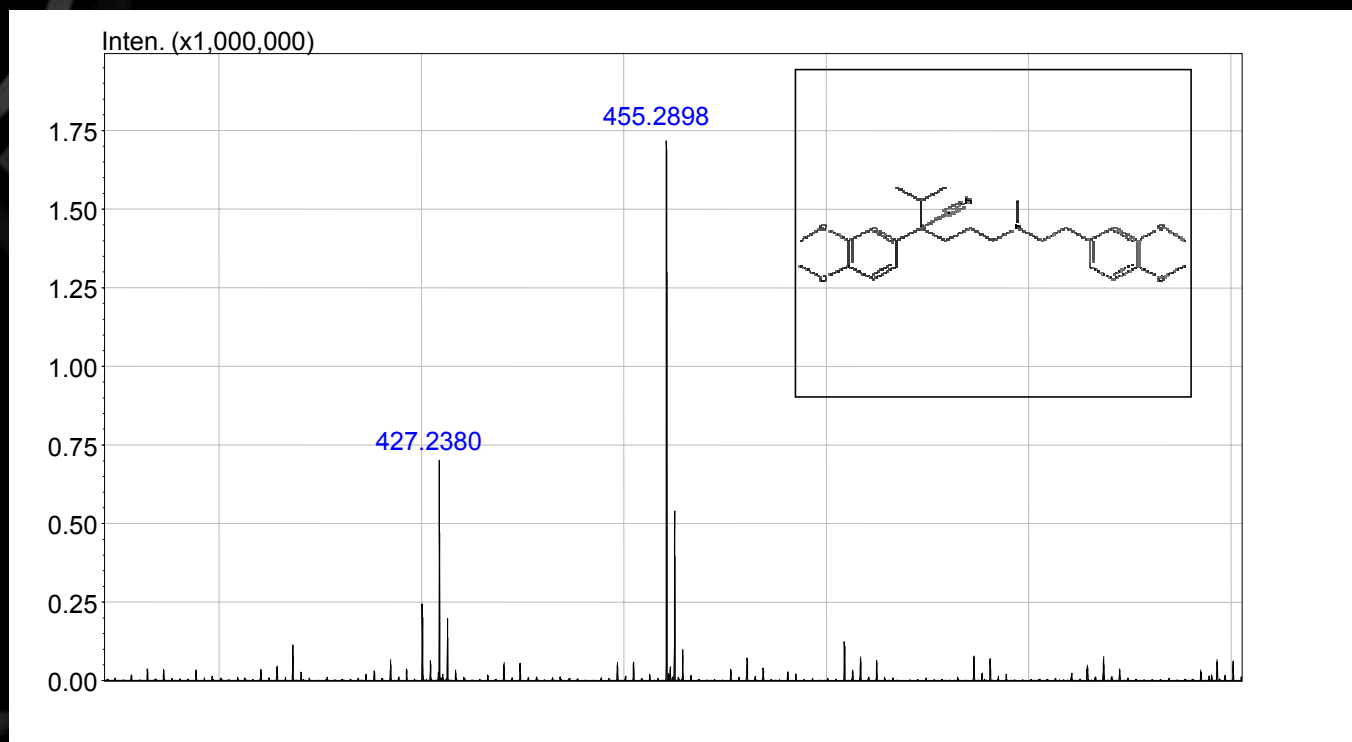
Fundamental Performance Data

Mass Spectrometry of Bovine Insulin 6+(5733m/z)



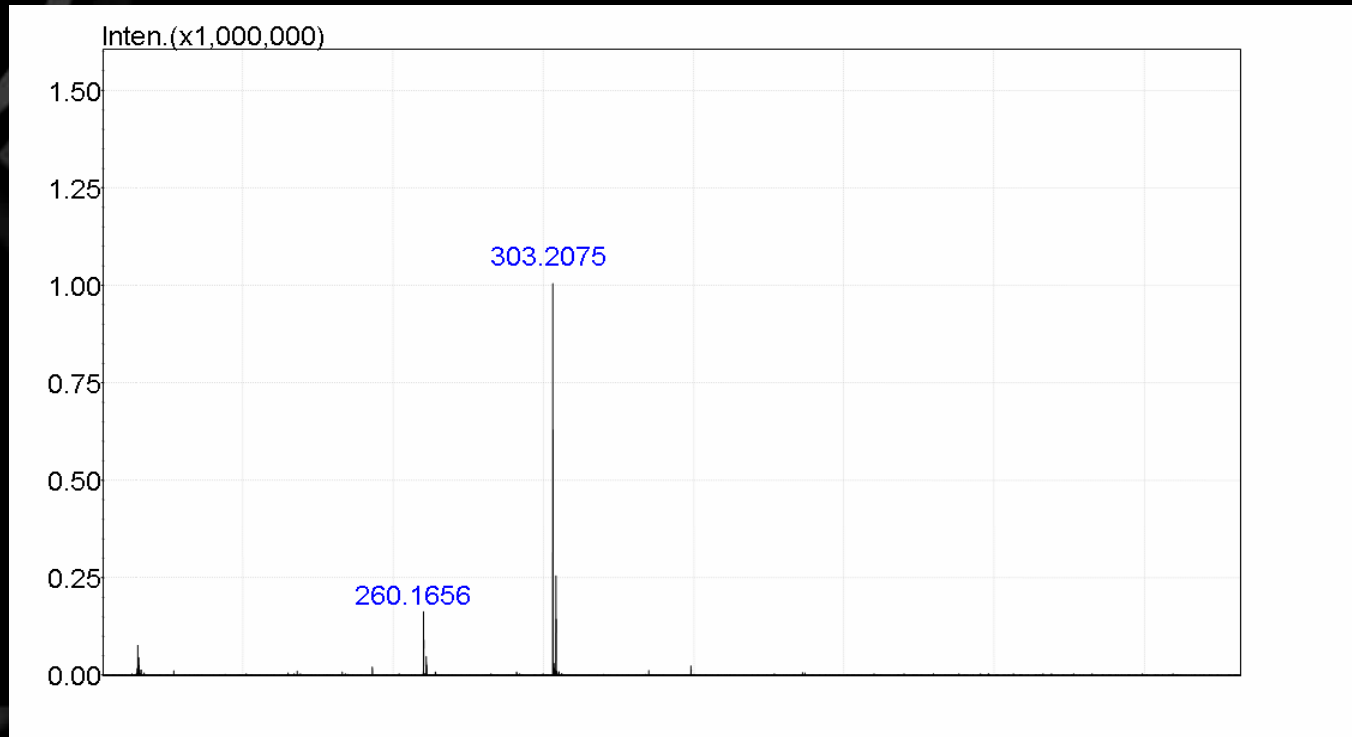
Mass Spectrometry of Verapamil

Sample	Expected	Found	Error
Verapamil $C_{27}H_{39}N_2O_4$	455.2910	455.2898	- 2.64 ppm



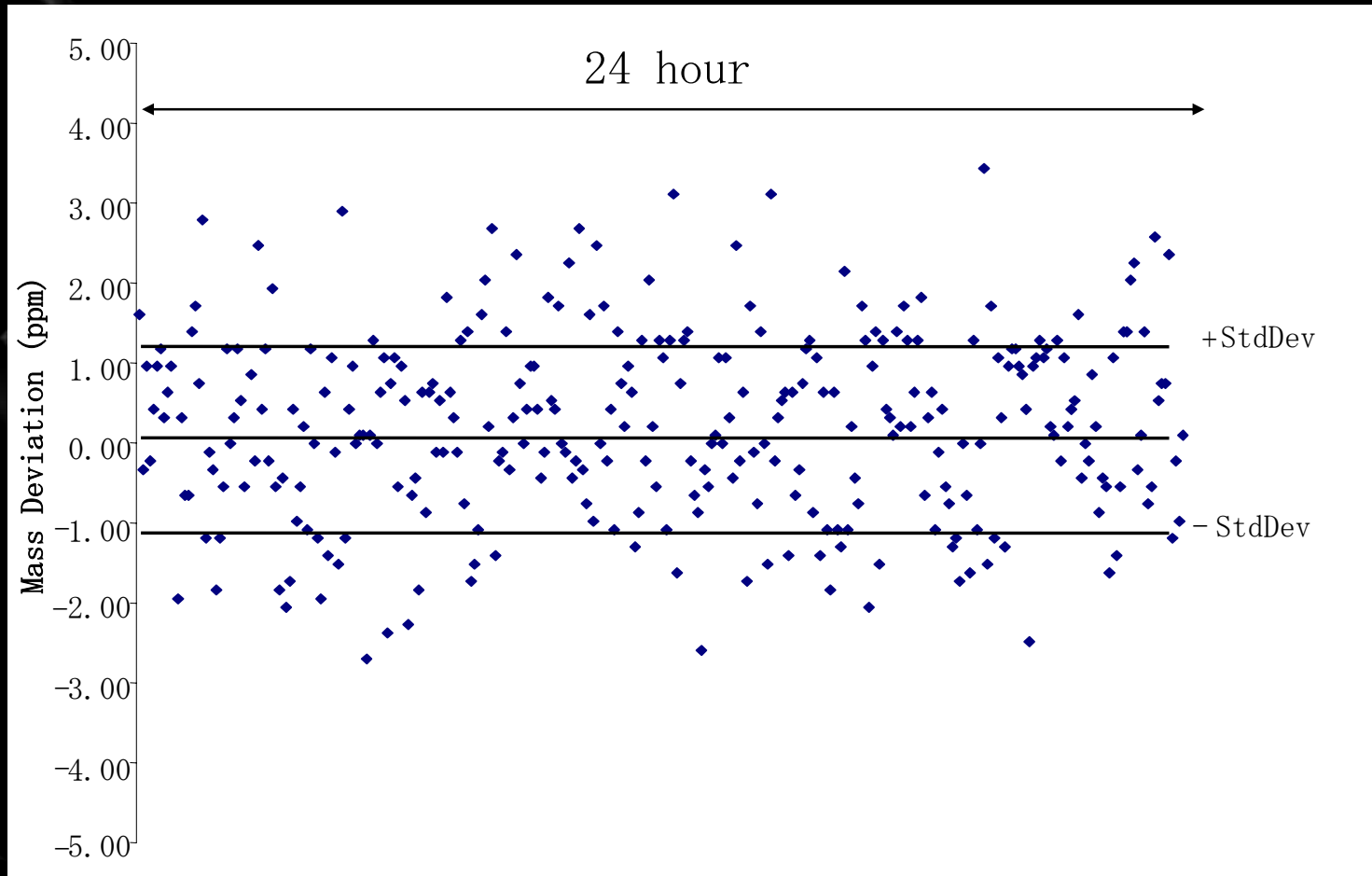
MS/MS Measurement of Verapamil

MS/MS ID	Expected	Found	Error
C18H27N2O2	303.2073	303.2075	+ 0.66 ppm



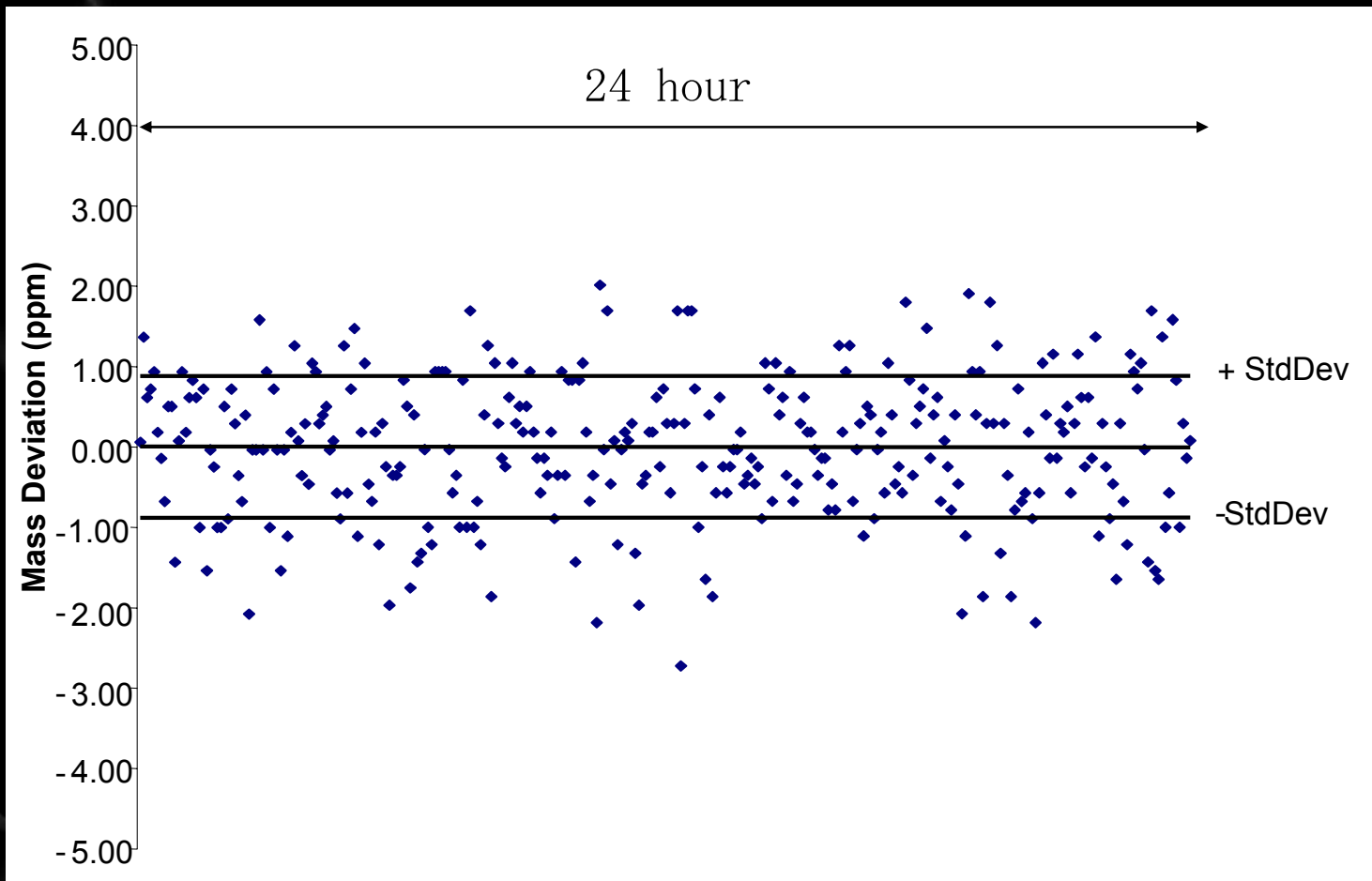
24-Hour Stability of Mass Precision

External Standard Method (m/z928)



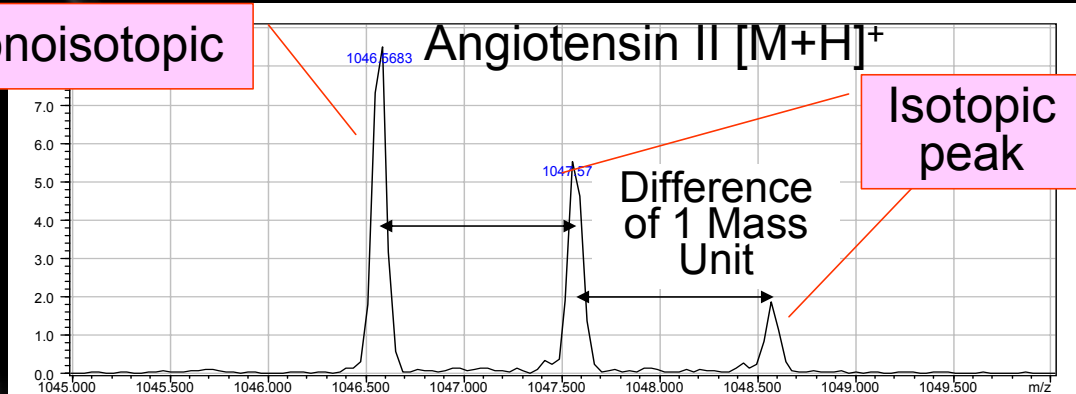
24-Hour Stability of Mass Precision

Internal Standard Method (m/z928)



Precursor Selection Using Ion Trap

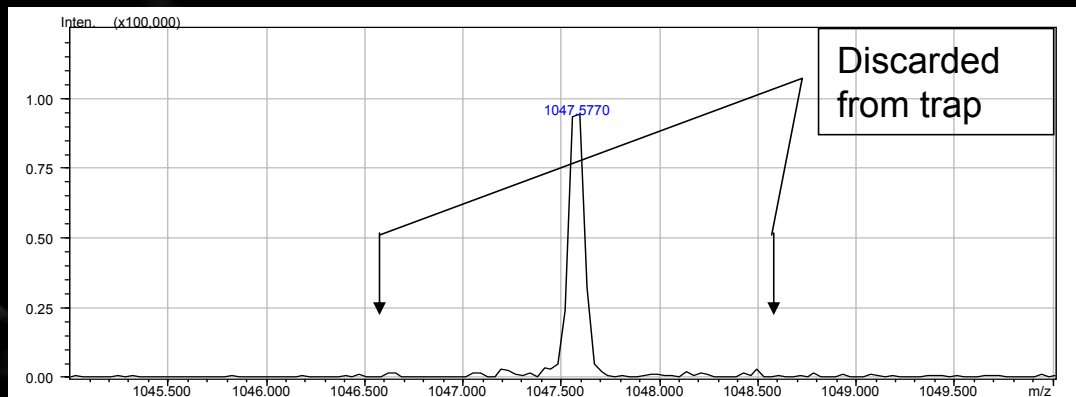
Monoisotopic



Before selecting precursor ion



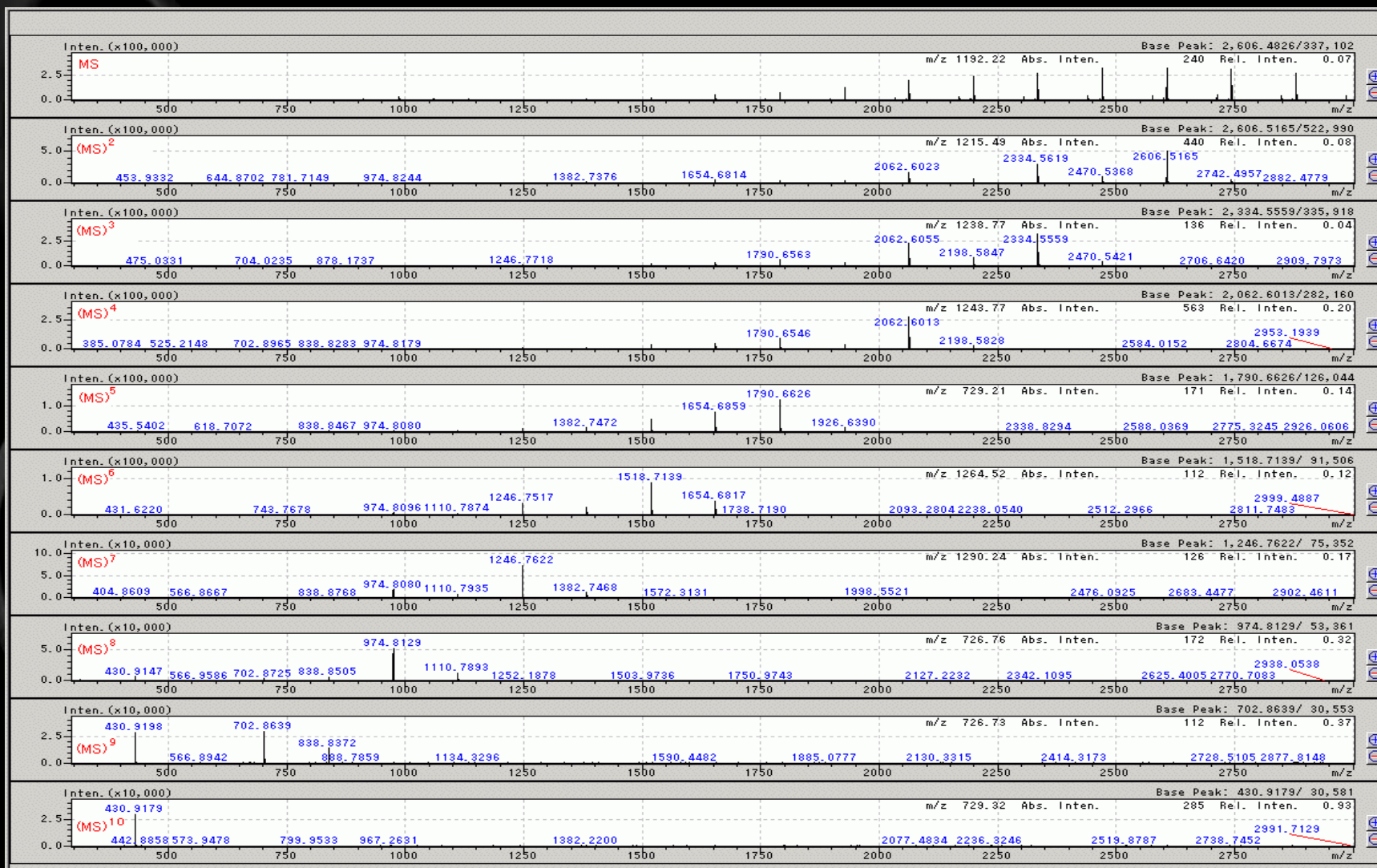
Trap selection of m/z 1047.57



After selecting precursor ion

MS¹⁰

using NaTFA sample



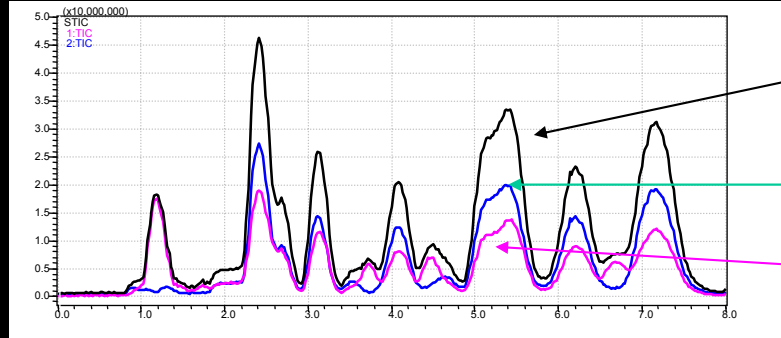
High Speed Pos/Neg Ion Switching (Max 2.5Hz)

Sample: glycosylceramide (C16,18,20,22,23,24:0-d18:1), (C24:1-d18:1) From Riken

Column Intakt Cadenza
CD-C18

Mobile Phase: 5mM
acetate
methanol
solution

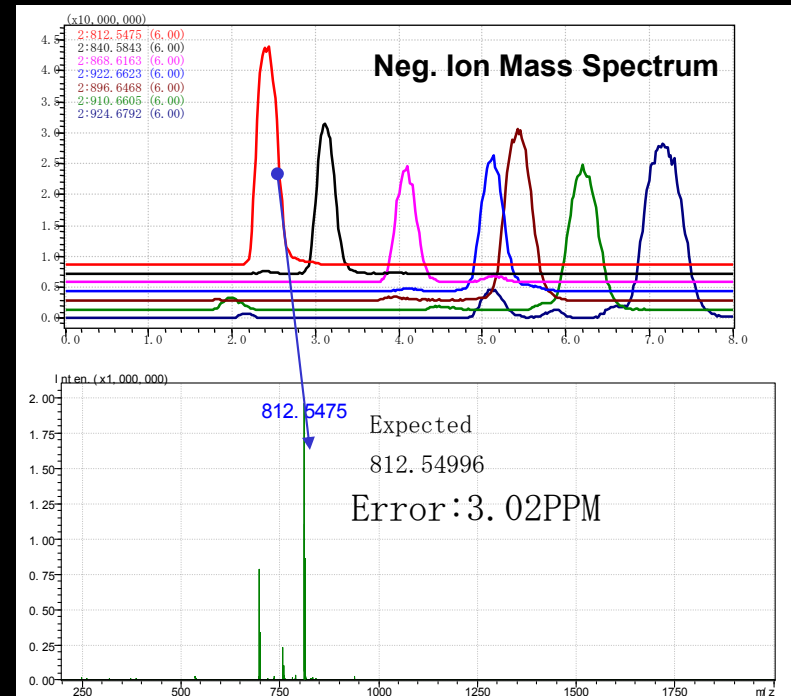
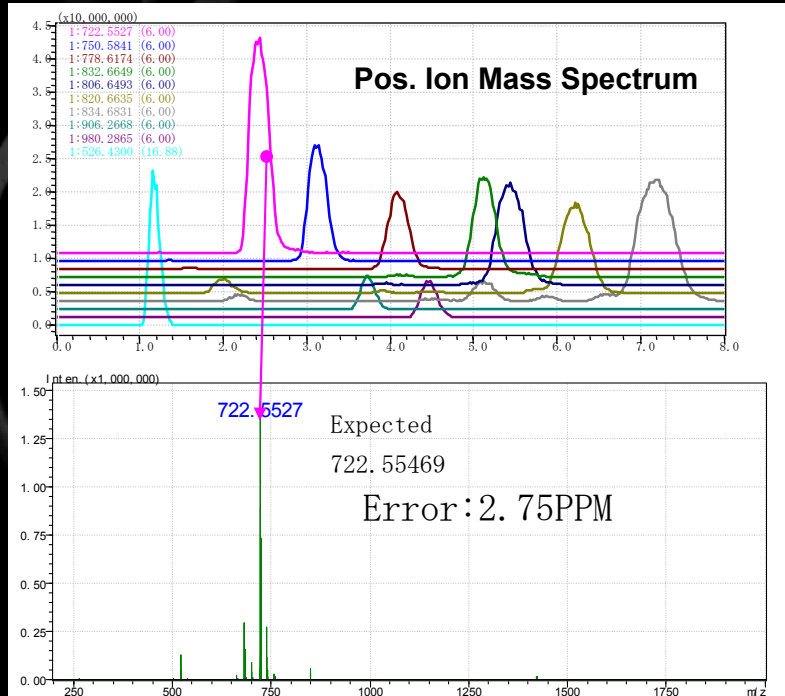
Flow Rate: 0.2ml/min
isocratic



With Pos/Neg Ion TIC

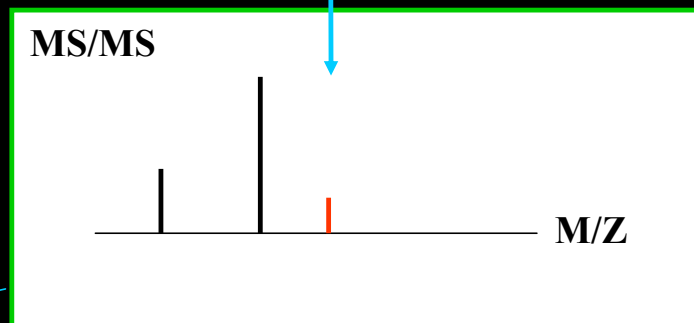
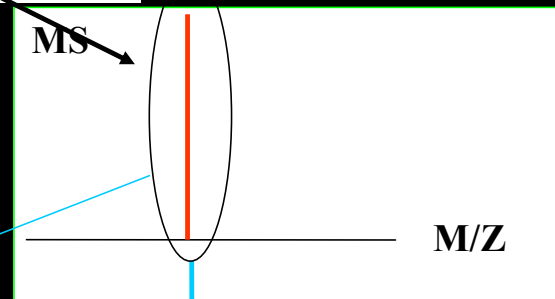
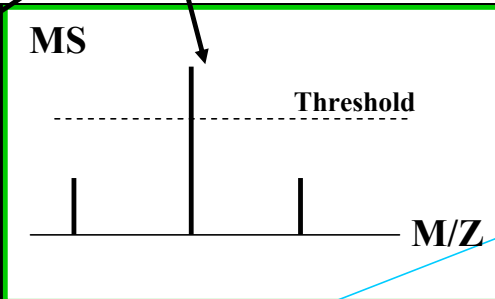
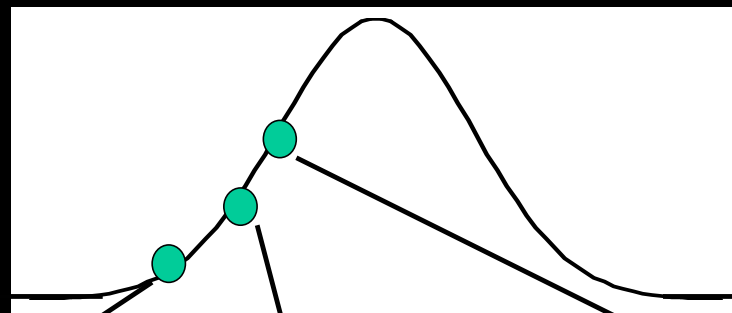
Neg. Ion TIC

Pos. Ion TIC



Auto MS/MS

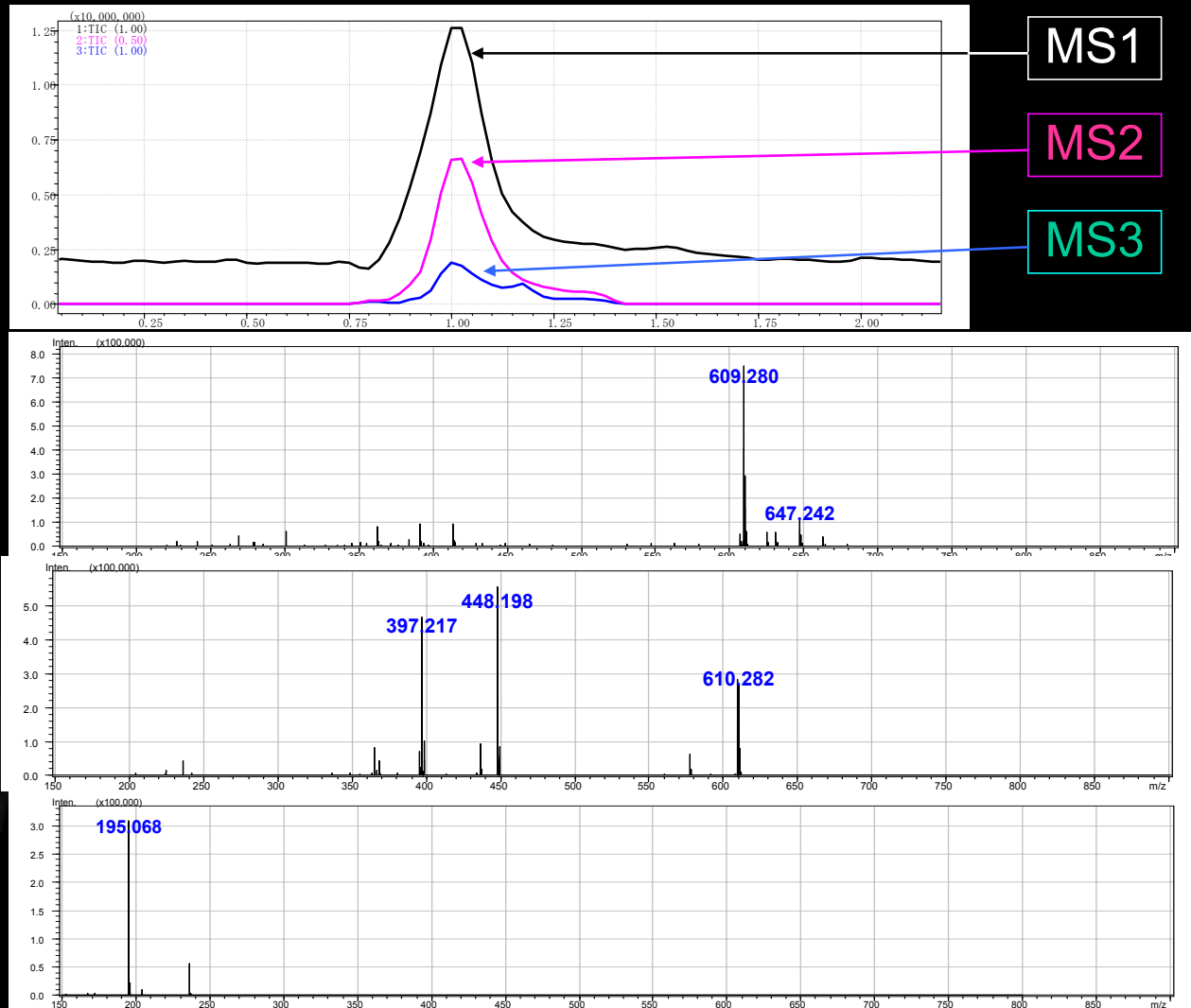
TIC



Peaks are selected automatically

Data after MS/MS analysis

Auto MS/MS



MS1

MS2

MS3

Summary of New Technologies

Compressed Ion Introduction Method

(Continuous ionization -> Pulsed ionization)

Ballistic Ion Ejection

(Transfers ions to TOF all at the same time with similar velocities for higher mass accuracy)

Ion Cooling using Argon Gas

(High resolution analysis of multivalent ions)

Dual Stage Reflectron Improvements

(High sensitivity and high resolution for QIT-TOF)

Temperature controlled interior of instrument housing

(Stability of high mass precision)