

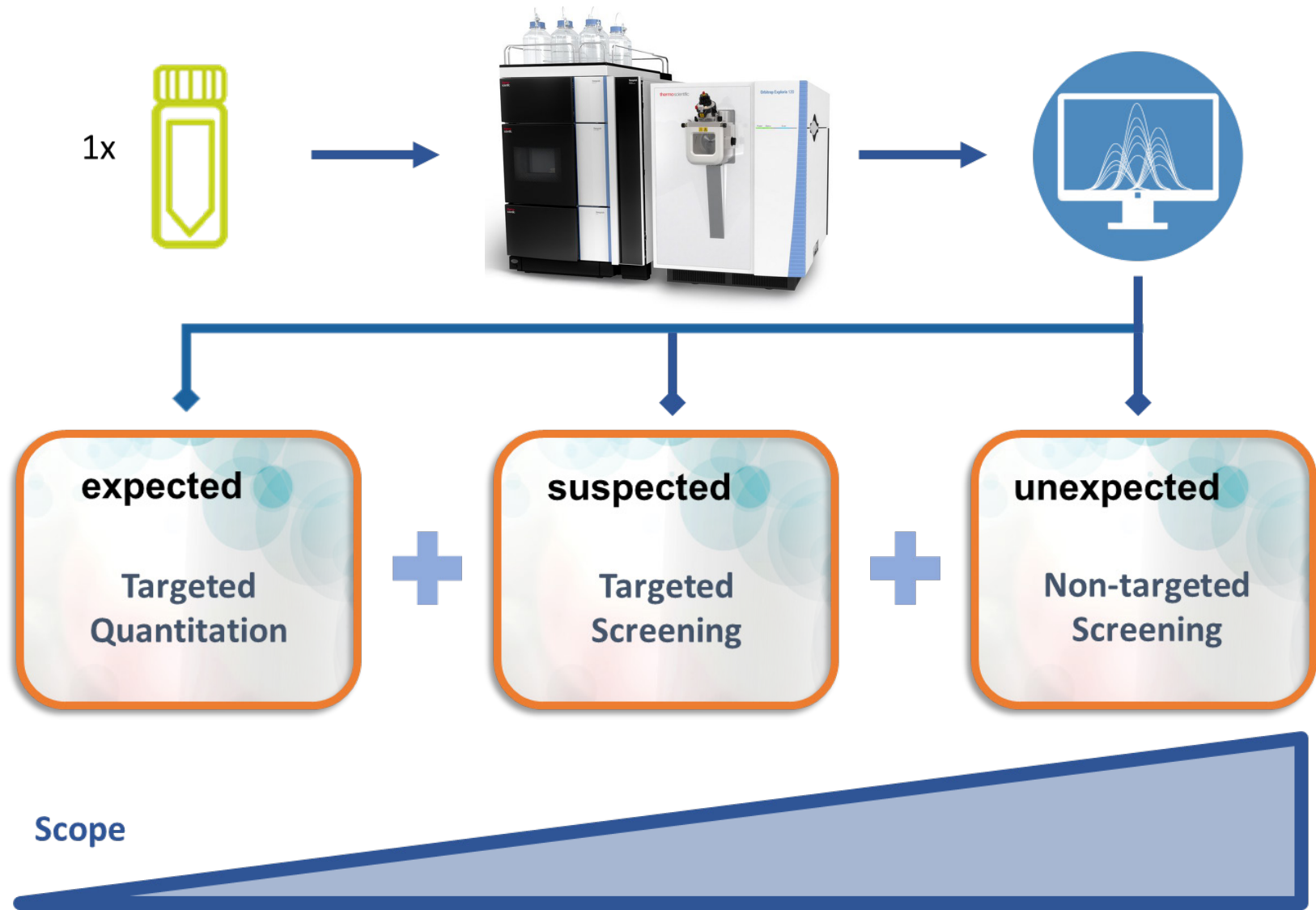


Comprehensive Solutions to Pharmaceutical Analysis

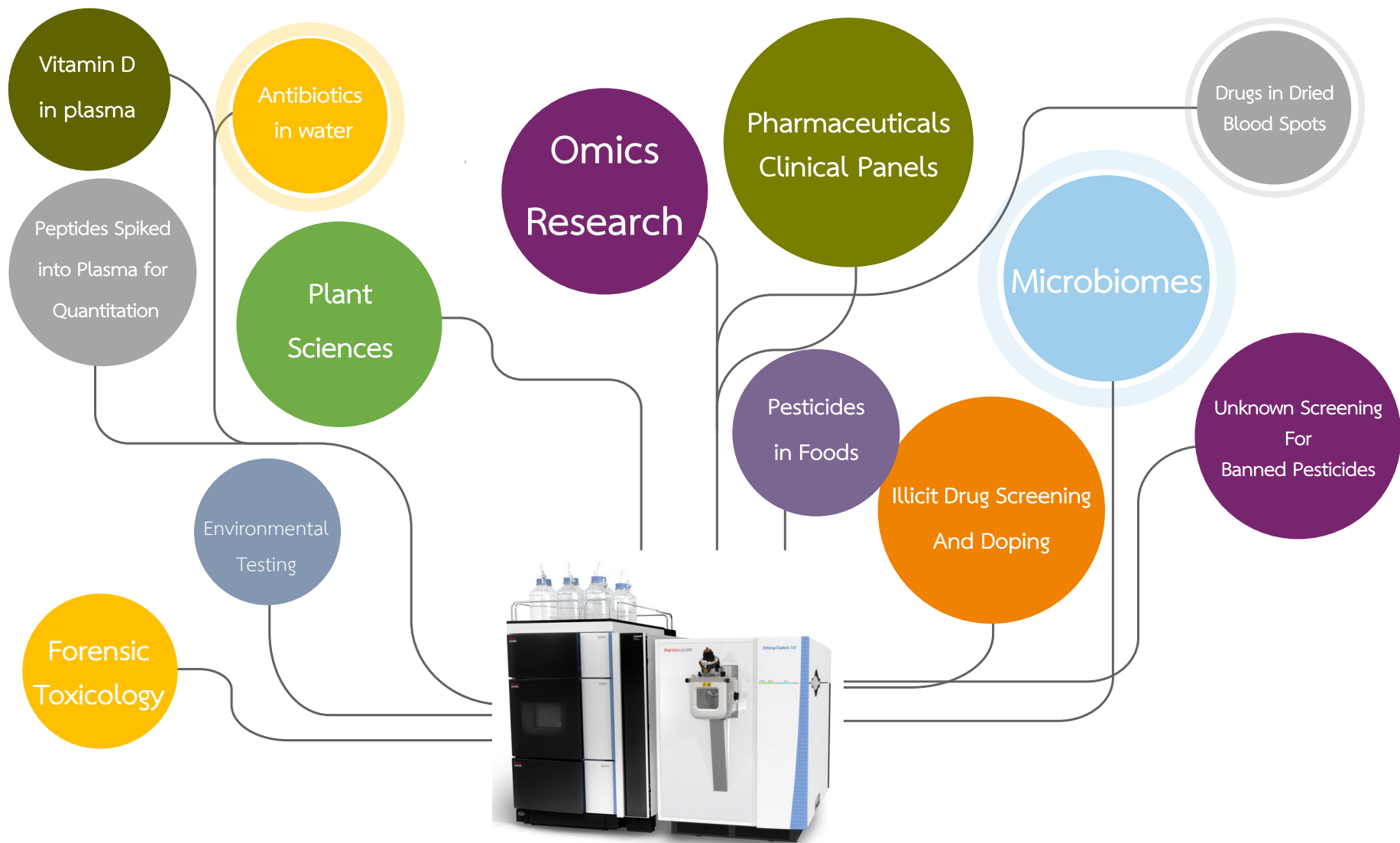
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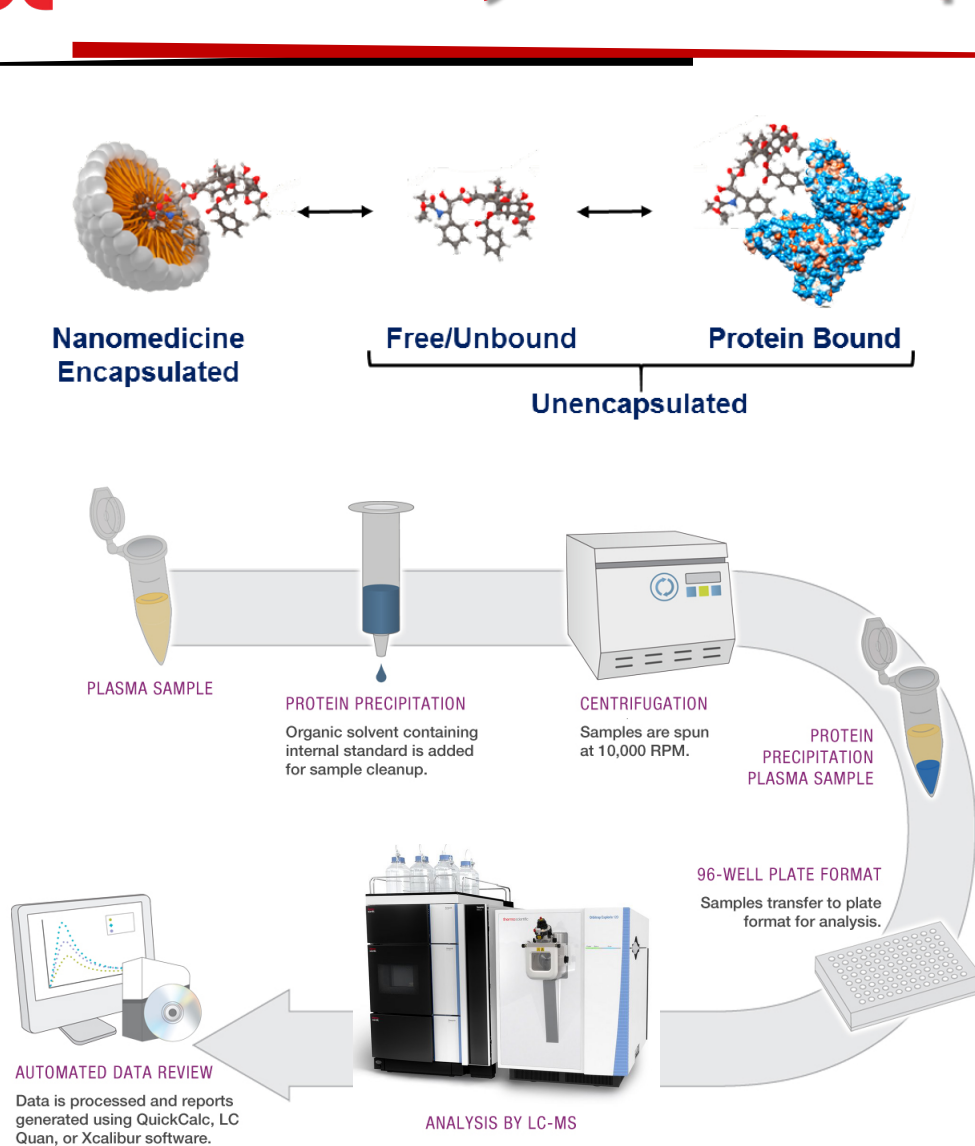
System Main Workflows



Orbitrap Applications Universe



Bioanalysis: Bioequivalence (BE)



Novel Method to Determine Bioequivalence of Nanomedicines
 summary of work conducted for
 U.S. Food & Drug Administration
 (Inter-Agency Award 224-16-3001S)

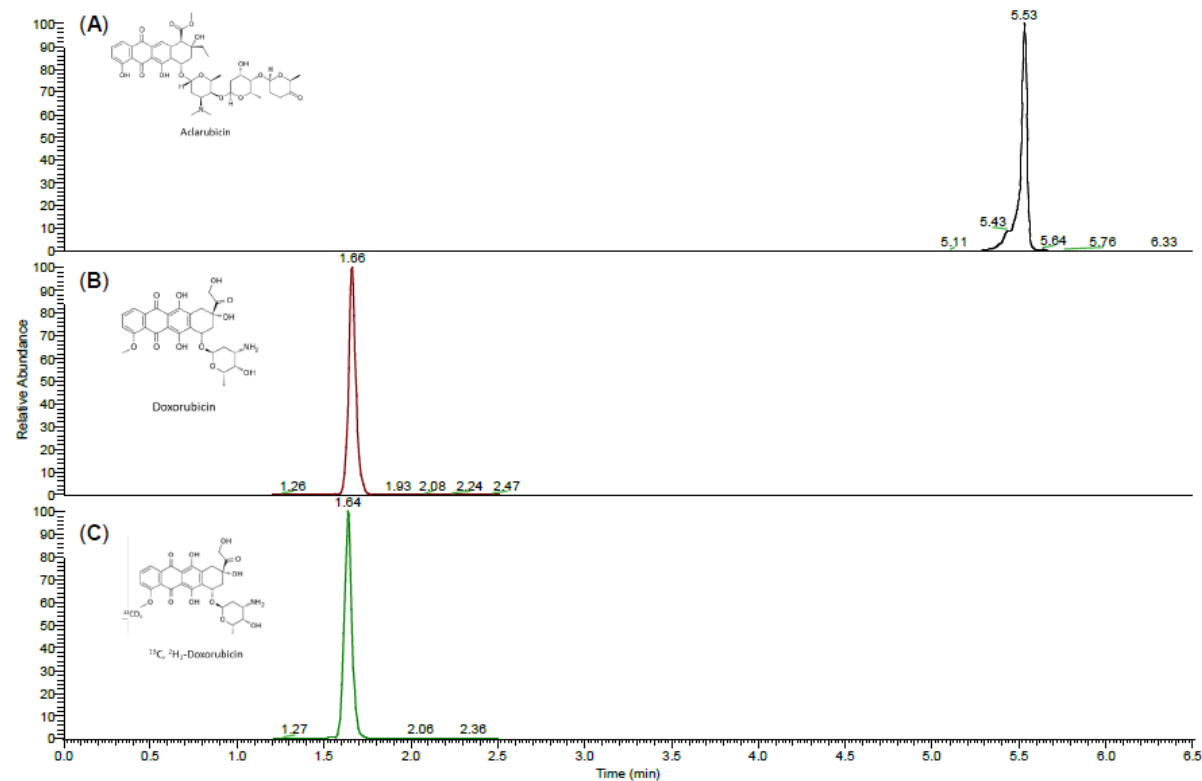


Figure B-2. LC-Q Orbitrap Parallel Reaction Monitoring of Analytes and Internal Standard. Displayed are chromatograms of (A) ISTD, (B) DXR and (C) DXR_C13. The transitions used were: DXR 544.18 → 130.08, 379.07, 397.08; DXR_C13 548.20 → 130.08, 383.09, 401.11; and Aclarubicin 812.34 → 570.23

Extractables and Leachables

Comparison of Soxhlet and Accelerated Solvent Extraction for Leachable and Extractable Analysis of Packing Material

Hua Ying,¹ Kate Cornstock,² and Linda Lopez²
¹Thermo Fisher Scientific, Sunnyvale, California, USA
²Thermo Fisher Scientific, San Jose, California, USA

Application Note 1108

Key Words

Pharmaceutical, Biopharma, U.S. FDA, Sample preparation, Packaging, Automated Extraction, HPLC, Pressurized fluid extraction

Introduction

According to the regulations and guidelines set by regulatory agencies, including the U.S. Food and Drug Administration (US FDA),¹ the European Medicines Agency (EMA),² extractable and leachable information must be included in applications for medical devices and container closure systems packaging for human drugs and biologics. The specific applications include the New Drug Application (NDA), Abbreviated New Drug Application (ANDA) and Biologics License Application (BLA).

Leachables are compounds that can migrate from components of medical devices or container closure systems into the drug product.³ They present safety concerns and may influence the effectiveness of a drug product. Extractables are compounds that can be extracted from components and are determined through Controlled Extraction Studies.⁴ The main purpose of



The accelerated solvent extraction technique is an automated technique with several advantages, including efficient extraction, reduced extraction time (<0.5 h/sample), reduced solvent use (<30 mL/sample), and flexibility in solvent selection. Using the method and sequence editor in the Dionex ASE 350 system, three

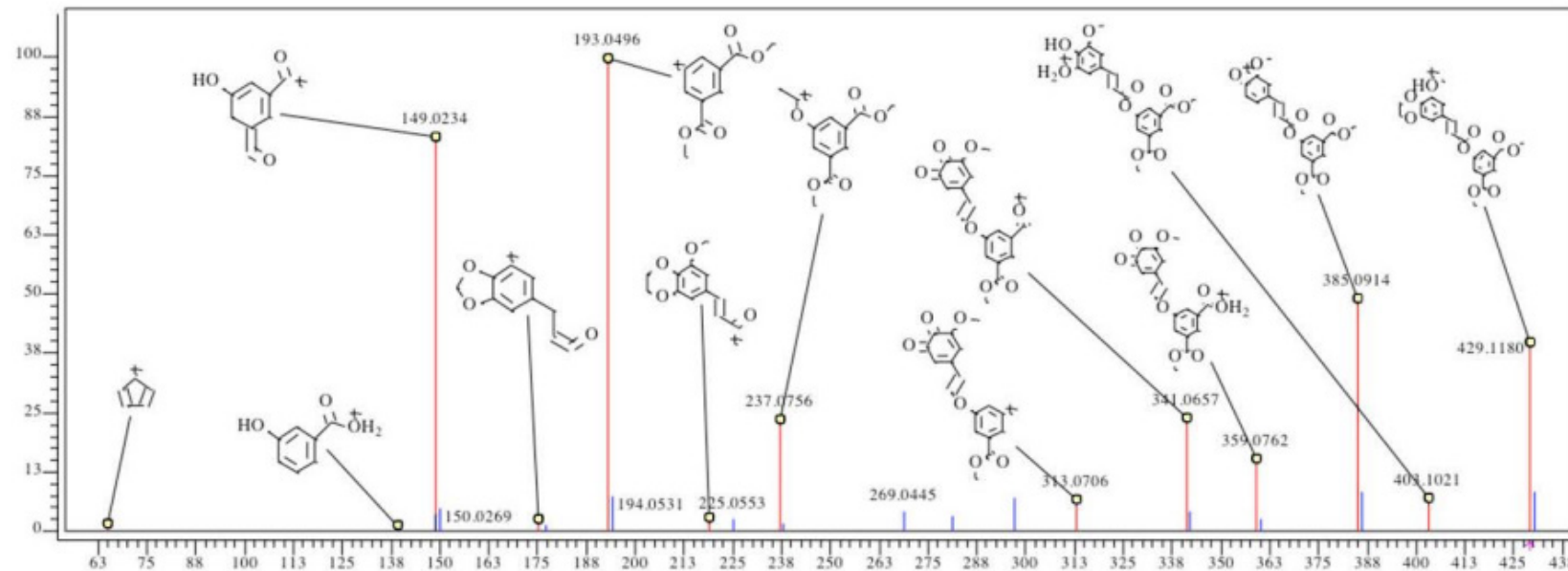
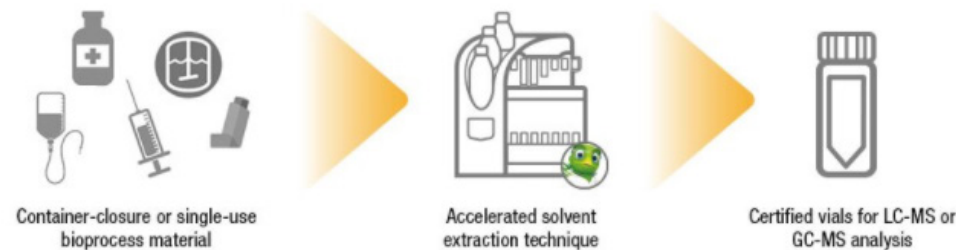
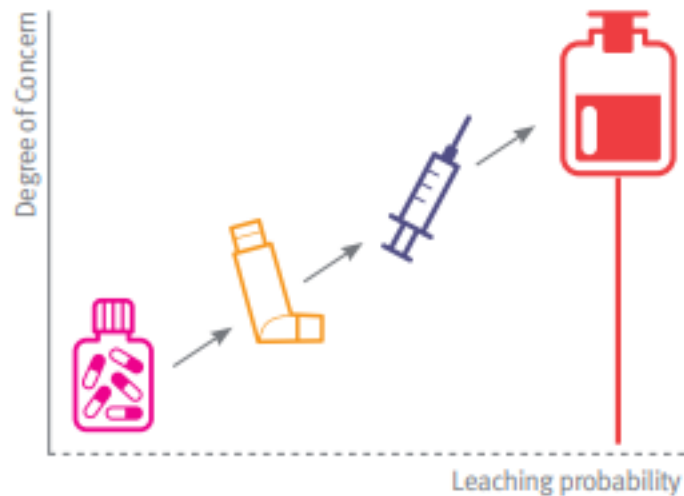
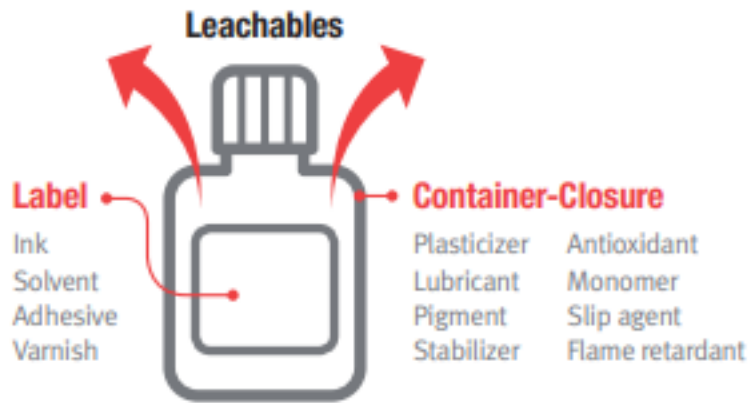
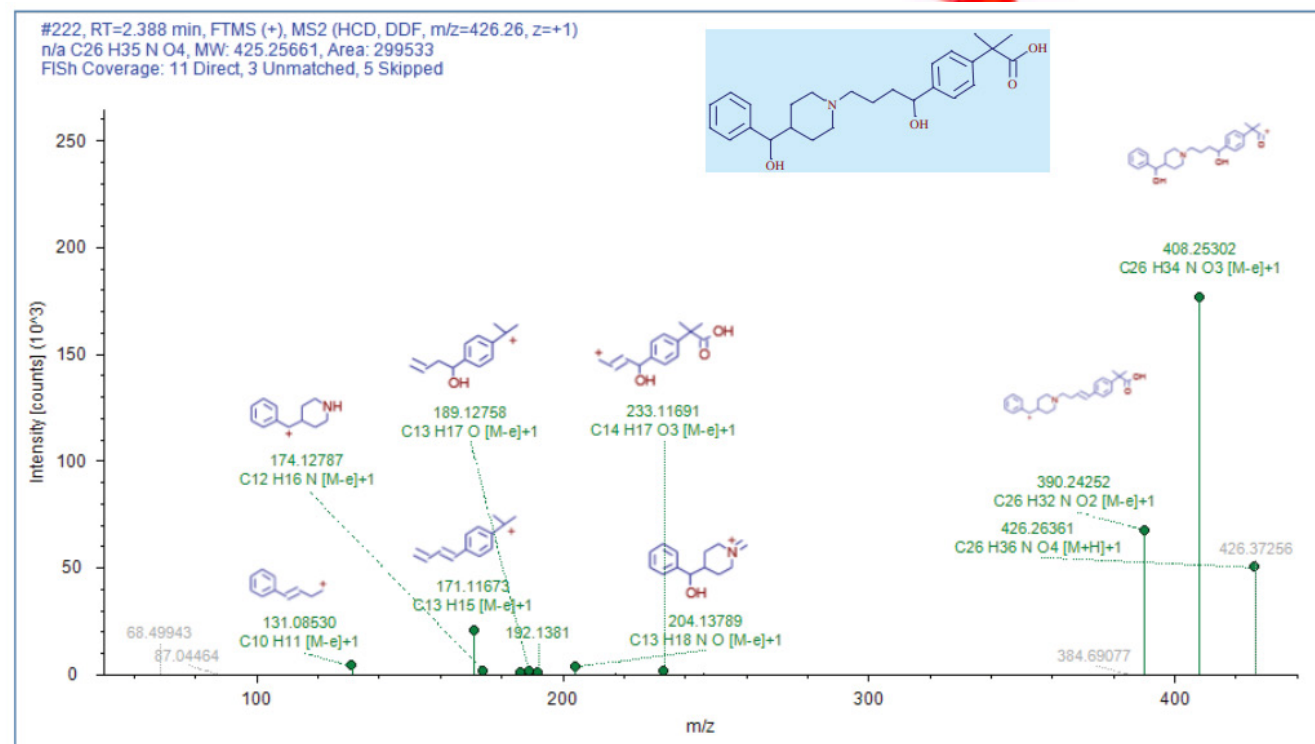
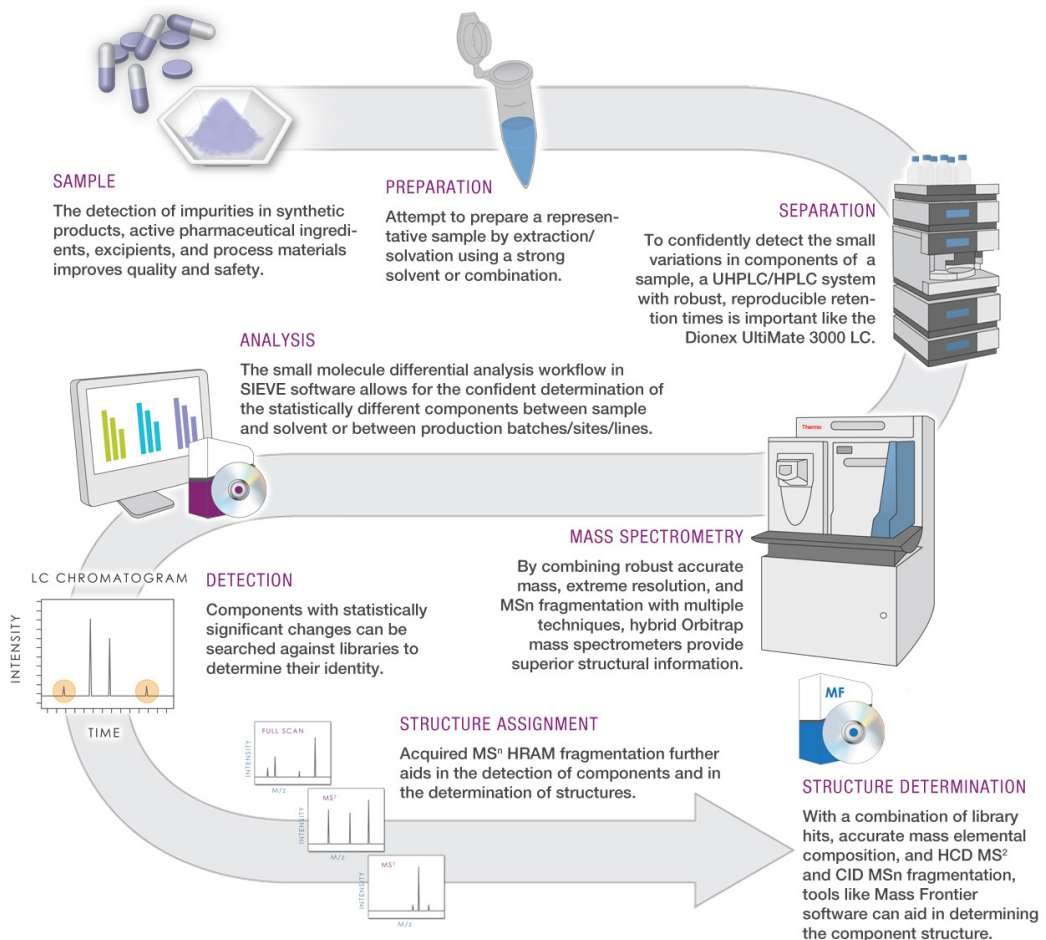


Figure 9. Auto-annotated MS/MS spectra of the proposed structure from 2-propanol extraction of the transdermal patch pouch.

Drug Impurities Analysis



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http://pca.fda.moph.go.th/public_media_detail.php?id=2&cat=50&content_id=1722



https://www.khaosod.co.th/around-thailand/news_2926836

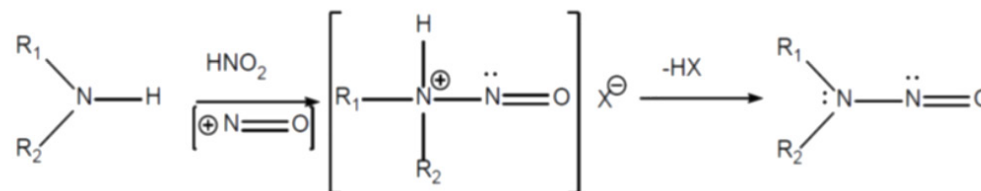
What are Nitrosamines ?

Nitrosamines (NAs) in APIs and drug products



- Classified as genotoxic impurities, proven as probable human carcinogen
- Detected elevated levels of NDMA and other impurities in several drug products, announced several recalls
- United States Food and Drug Administration (US FDA) published validated GC/LCMS/MS and GC/LC-HRAM methods, and interim acceptable limits for several NAs
- US FDA recently released guidance for industry to
 - 1) conduct risk assessments on approved or marketed drugs and pending applications,
 - 2) Implement control strategy to reduce and prevent formation of NAs in all API and drug

Figure 1. Representative Reaction to Form Nitrosamines





Selectivity

- High background, formulation matrices and other impurities from drug substances and solvents
- Chemical Interferences (e.g., DMF and NDMA)
- Structural isomers (e.g., NDIPA and NDPA)



Sensitivity

- Trace level of impurities in presence of high concentration of API
- Total nitrosamine contents should be less than 30 ppb or 26.5 ng/day based on daily maximum dosage intake of 880 mg/day

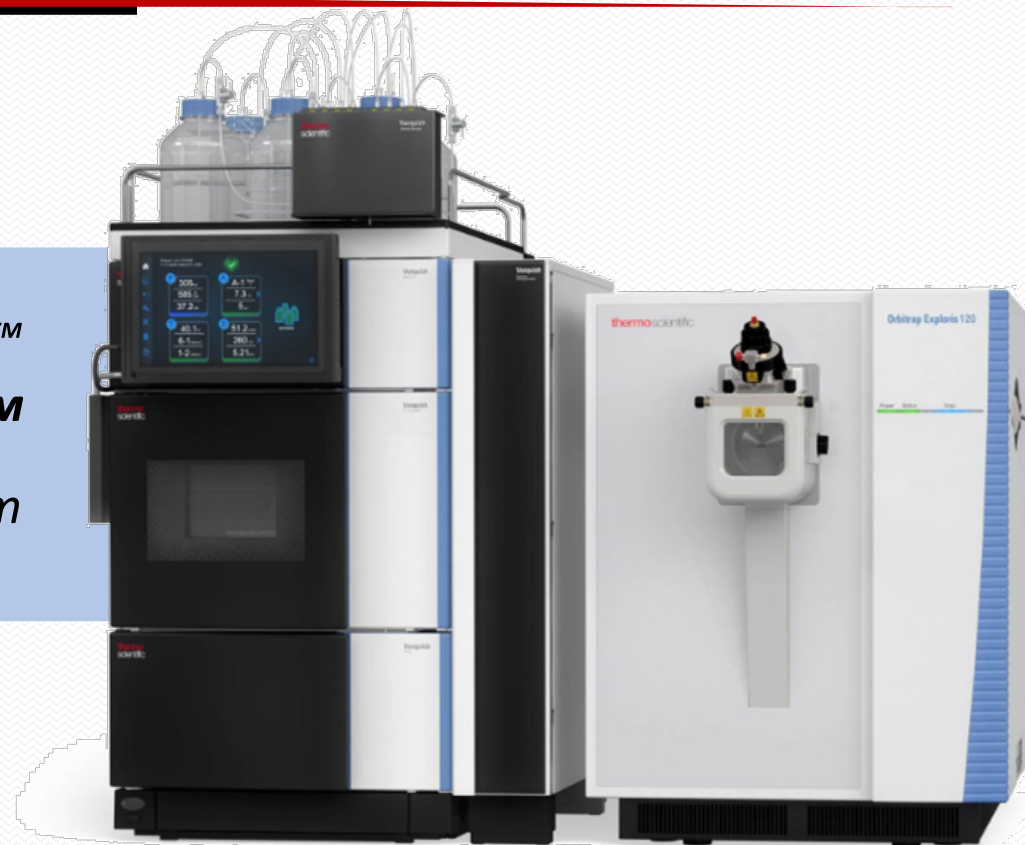


Regulatory Expectation and Compliance

- Meet US FDA requirements in terms of sensitivity, linearity, and reproducibility
- Data collection and processing in compliant software to ensure data integrity and security

Analytical method needs to demonstrate high selectivity, sensitivity, and meets US FDA regulatory guidelines

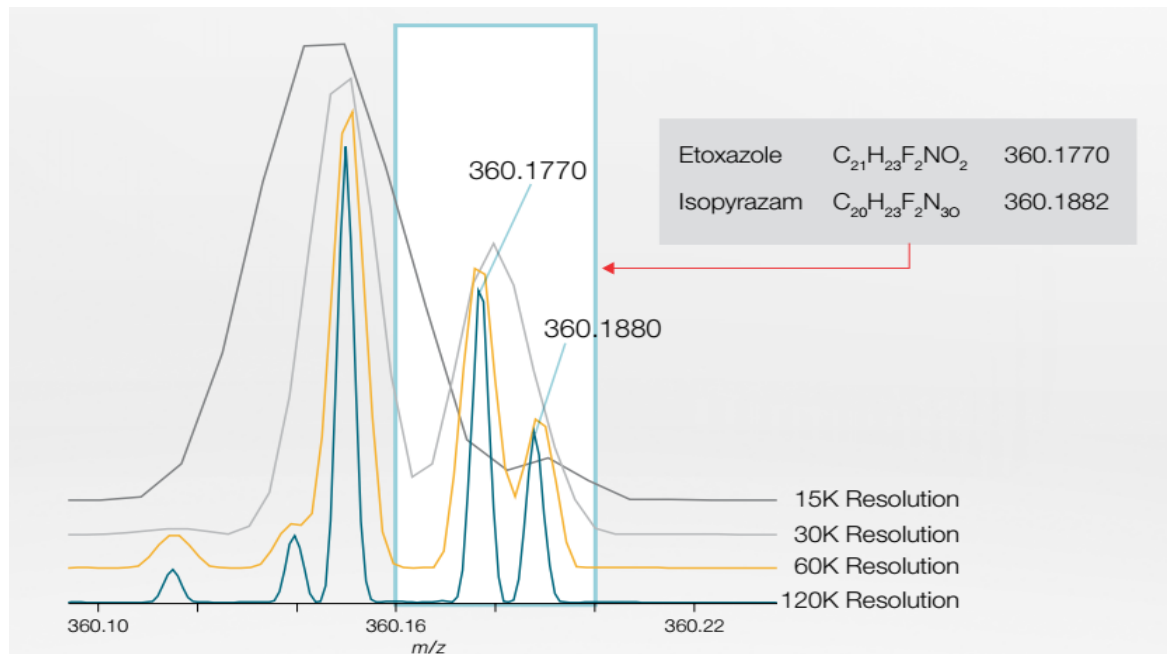
Thermo Scientific™
Vanquish™
UHPLC System



Thermo Scientific™
Exploris™ 120
HRAM-MS System

A rapid, highly selective and sensitive LC-HRAM-MS method was developed for detection and quantitation of 9 nitrosamines in commercial ranitidine drug product

Product Specification

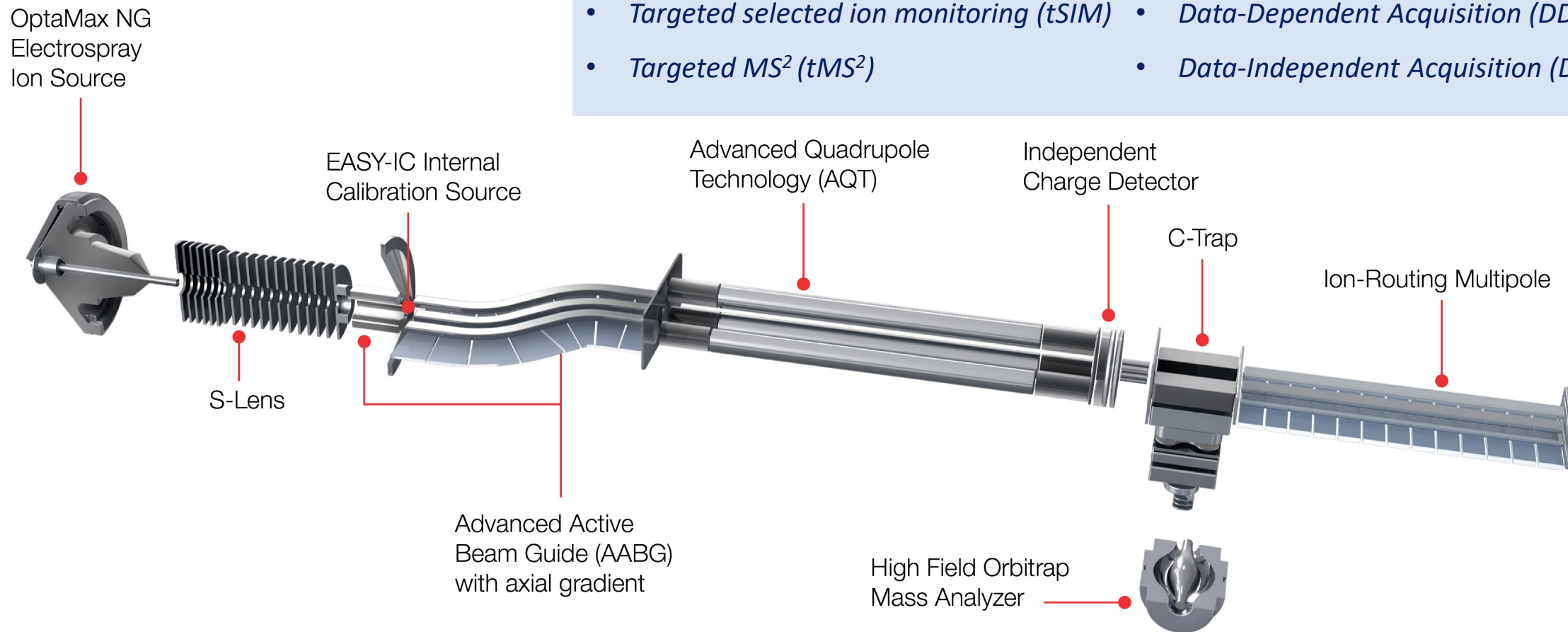


Orbitrap Exploris 120	
Resolution @ m/z 200	up to 120,000 FWHM
Mass Range	40 – 3,000
Scan Rate	up to 22 Hz
Sensitivity	MS/MS: 200 fg reserpine on column S/N 100:1 tSIM: 200 fg reserpine on column S/N 250:1



Data Acquisition Mode

- Full MS scan
- Targeted selected ion monitoring (tSIM)
- Targeted MS² (tMS²)
- All Ion Fragmentation (AIF)
- Data-Dependent Acquisition (DDA)
- Data-Independent Acquisition (DIA)

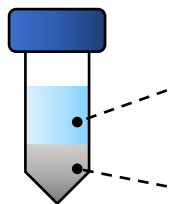


Analyte	Internal standard (ISTD)
N-Nitrosodiethylamine (NDEA)	NDEA-D ₁₀
N-Nitrosodimethylamine (NDMA)	NDMA-D ₆
N-Nitrosodi-n-butylamine (NDBA)	NDBA-D ₁₈
N-Nitroso-di-n-propylamine (NDPA)	NDPA-D ₁₄
N-Nitrosomethylethylamine (NMEA)	NMEA-D ₃
N-Nitrosopiperidine (NPIP)	NPIP-D ₁₀
N-Nitrosopyrrolidine (NPYR)	NPYR-D ₈
N-Ethyl-N-nitroso-2-propanamine (NEIPA)	NEIPA-D ₅
N-Nitroso-di-isopropylamine (NDIPA)	NDIPA-D ₁₄

Sample Preparation

Ranitidine substance

Dissolution



1 ml MeOH
+ 10 μ l 500 ppb ISTD
30 mg ranitidine hydrochloride

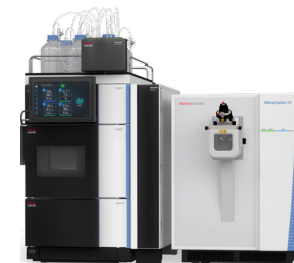
Total concentration = 30 mg/mL API



Filter 750 μ l solution
using 0.2 μ m PVDF
syringe filter



Inject 5 μ l
to LC-HRMS



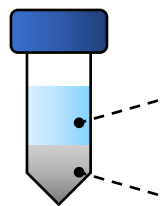
Drug Product



Ranitidine tablet



Extraction



10 ml MeOH
+ 100 μ l 500 ppb ISTD
300 mg API, ground ranitidine tablet

Total concentration = 30 mg/mL API



Shake for 40 min



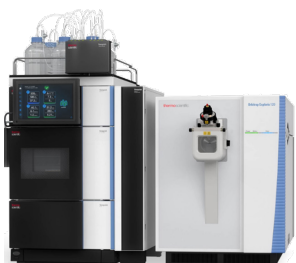
Centrifuge for 15 min at
4000 RPM



Filter 750 μ l supernatant using
0.2 μ m PVDF syringe filter



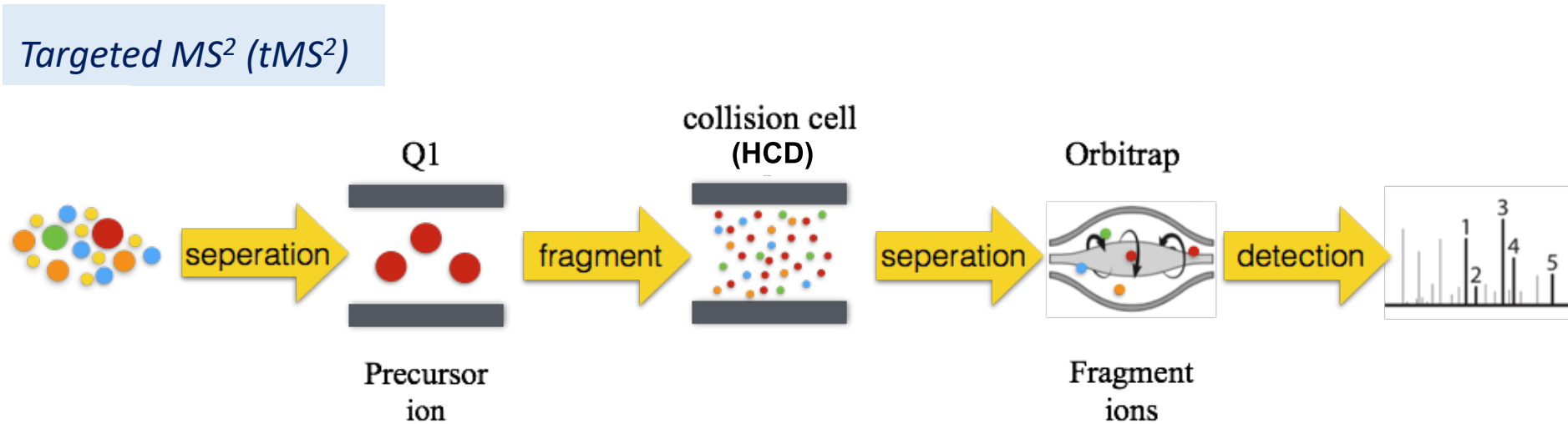
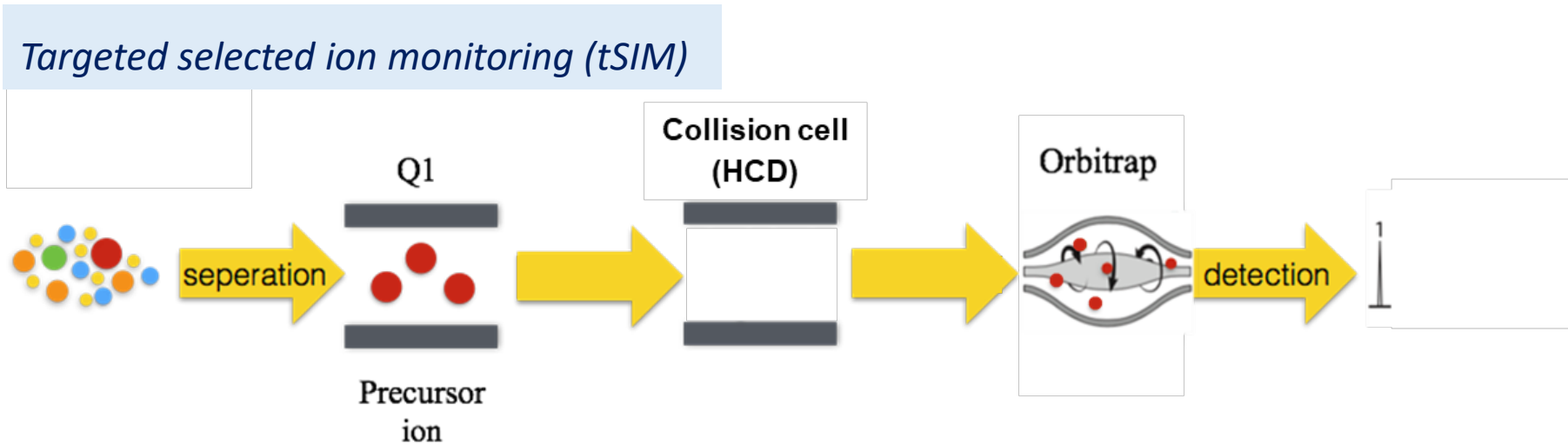
Inject 5 μ l
to LC-HRMS



Parameters	Value		
HPLC column	Acclaim Polar Advantage II, 100 × 2.1 mm, 2.2 μm		
Column temperature	40 °C		
Flow rate	0.5 mL/min		
Mobile phase A	Water + 0.1% formic acid		
Mobile phase B	Methanol + 0.1% formic acid		
Gradient	Time (min)	%Mobile phase A	%Mobile phase B
	0.0	95	5
	0.5	95	5
	8.0	5	95
	9.0	5	95
	9.1	95	5
	12.0	95	5
Injection volume	5 μL		

Parameters	Value	
Ionization	APCI	
Polarity	Positive	
Spray current	2 μA	
Sheath gas	45	
Auxiliary gas	10	
Sweep gas	0.5	
Ion transfer tube temp.	200 °C	
Vaporizer temp.	300 °C	
Scan mode	tSIM	tMS ²
Polarity	Positive	Positive
Resolution	120,000	120,000
AGC target	1e5	1e5
Maximum IT	Auto	Auto
Isolation window	m/z 2.0	m/z 2.0

Acquisition Mode (Target Quantitation)



Optimized MS Condition for Target Nitrosamines

	Scan type	Scan Start – End (min)	Polarity	<i>m/z</i> of Quan. ion	<i>m/z</i> of Qual. Ion	Normalized CE (%)
NDMA	tMS ²	0.25–1.75	Positive	75.0552	43.0290	60
NDMA-D ₆	tMS ²	0.25–1.75	Positive	81.0928	46.0480	60
NMEA	tMS ²	0.75–2.25	Positive	61.0397	89.0708	15
NMEA-D ₃	tMS ²	0.75–2.25	Positive	64.0585	92.0898	30
NPYR	tMS ²	0.75–2.25	Positive	101.0709	55.0540	60
NPYR-D ₈	tMS ²	0.75–2.25	Positive	109.1212	62.0980	30
NDEA	tMS ²	1.95–3.45	Positive	103.0866	75.0550	45
NDEA-D ₁₀	tMS ²	1.95–3.45	Positive	113.1493	81.0930	45
NPIP	tMS ²	2.35–3.85	Positive	115.0866	69.0699	60
NPIP-D ₁₀	tMS ²	2.35–3.85	Positive	125.1494	78.1260	60
NEIPA	tMS ²	2.85–4.35	Positive	75.0553	117.1022	15
NEIPA-D ₅	tMS ²	2.85–4.35	Positive	80.0866	122.1336	15
NDIPA	tSIM	3.75–5.25	Positive	131.1179	-	-
NDIPA-D ₁₄	tSIM	3.75–5.25	Positive	145.2058	-	-
NDPA	tSIM	4.25–5.75	Positive	131.1179	-	-
NDPA-D ₁₄	tSIM	4.25–5.75	Positive	145.2058	-	-
NDBA	tMS ²	5.75–7.25	Positive	159.1492	103.0866	15
NDBA-D ₁₈	tMS ²	5.75–7.25	Positive	177.2622	66.1264	15



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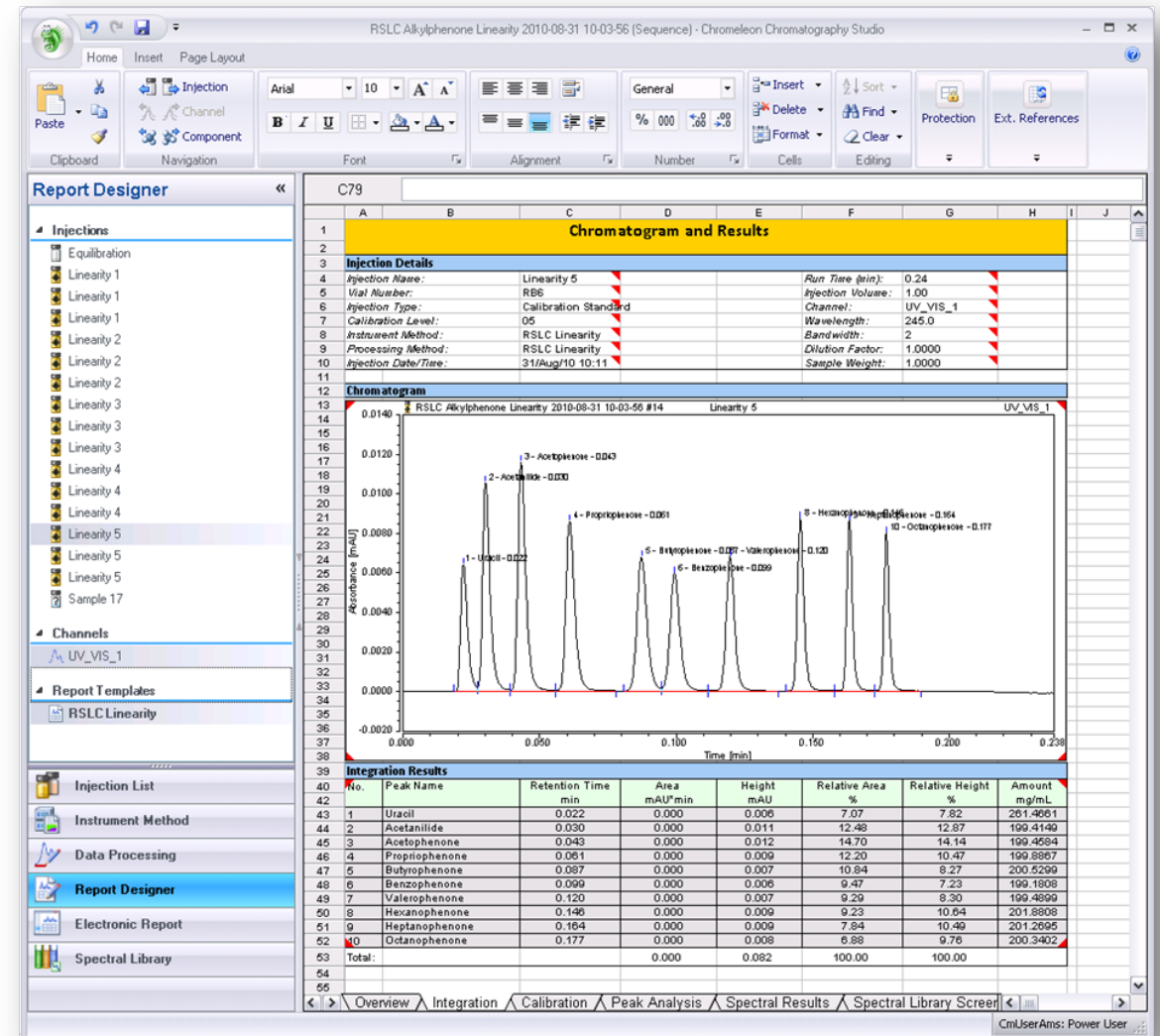
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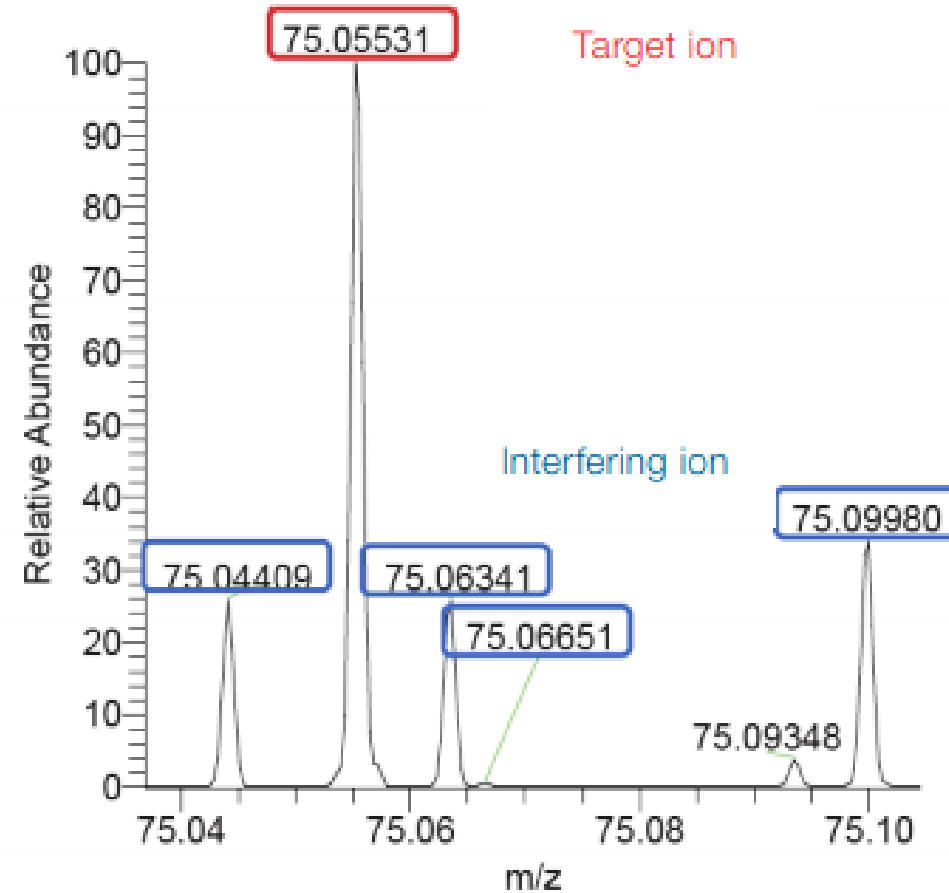
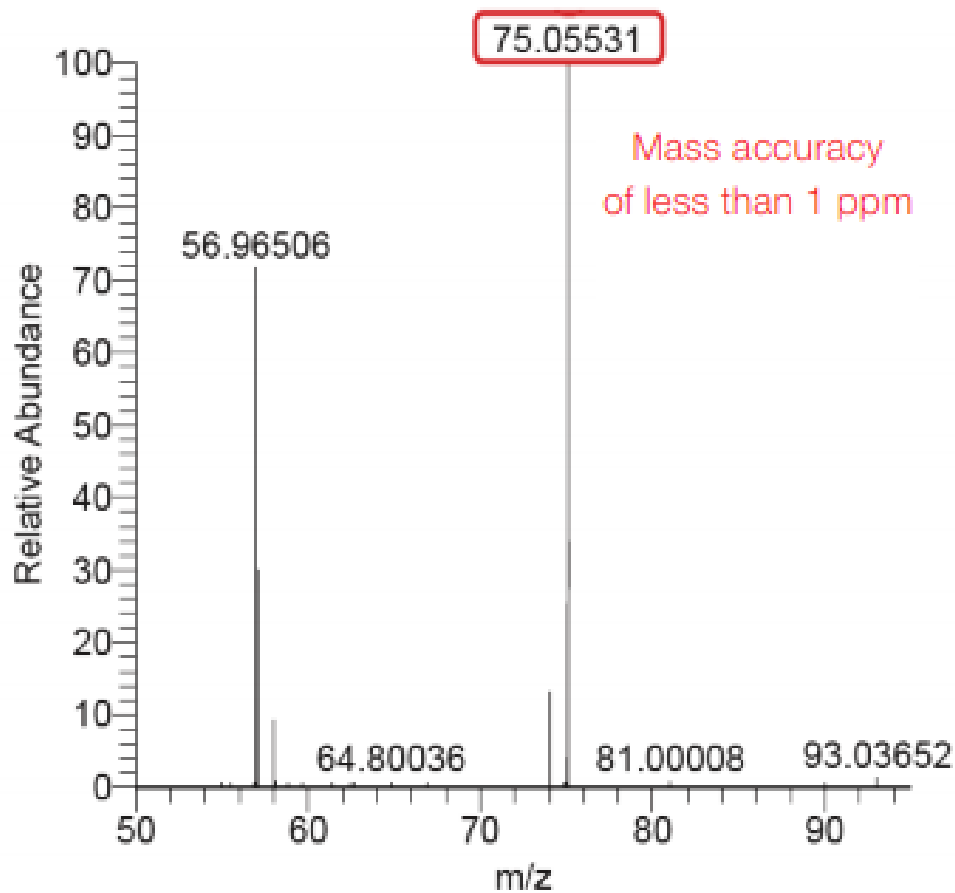
Using **Chromeleon 7** Chromatography Data System to Comply with **21 CFR Part 11**



Selective Determination of Nitrosamine Impurities

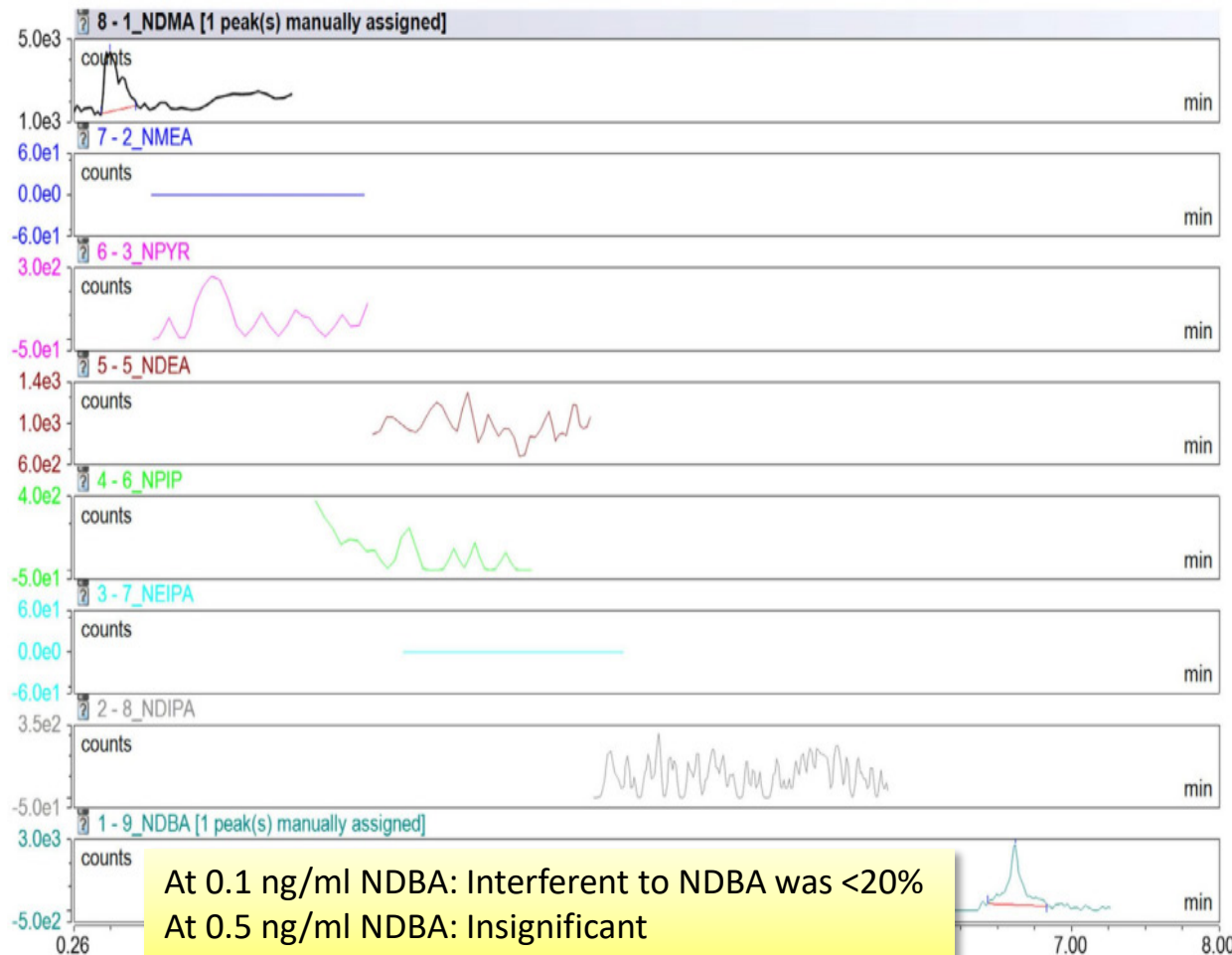
Mass spectrum of 1 ng/ml NDMA standard solution

Target monoisotopic mass of NDMA = 75.0552



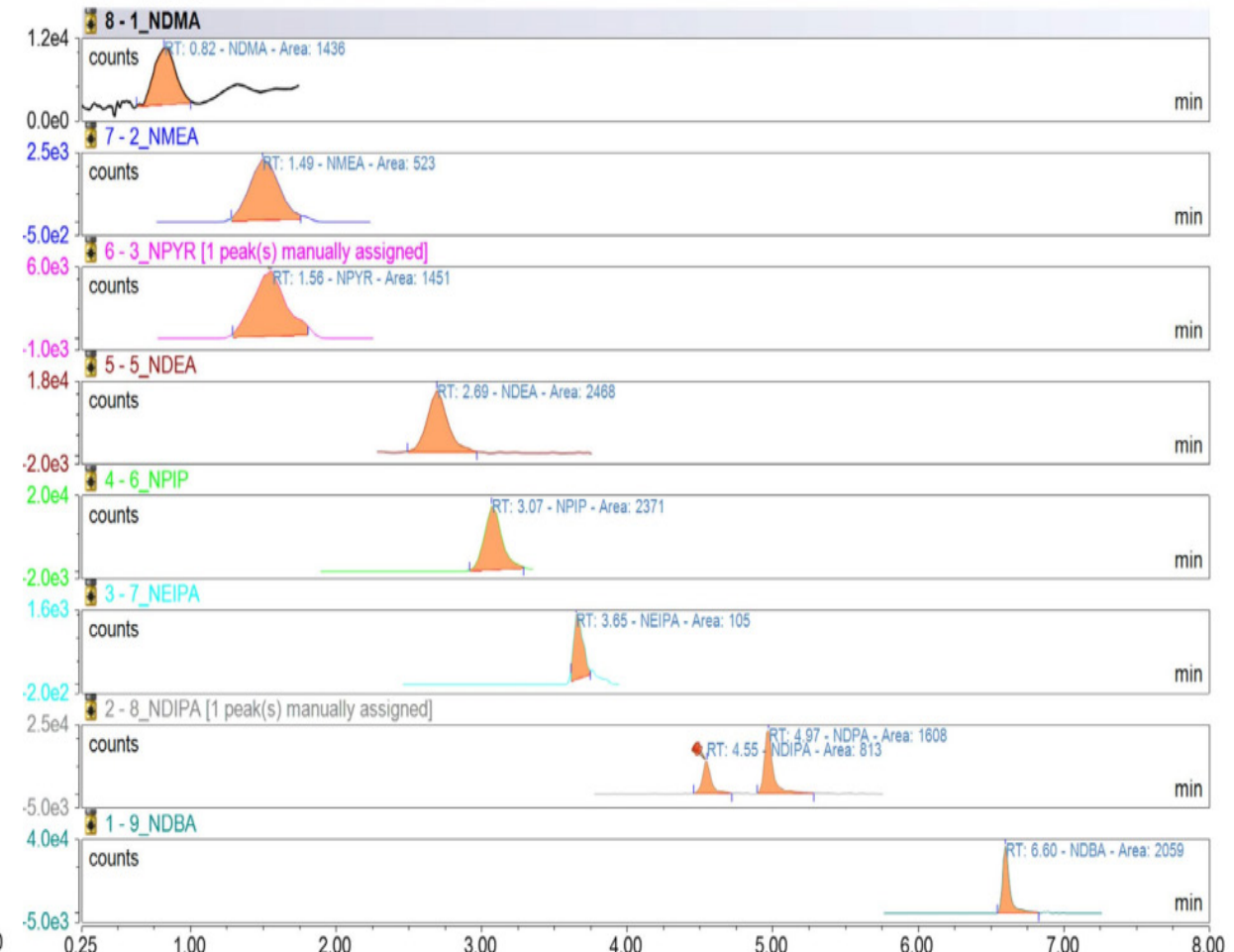
Selective Determination of Nitrosamine Impurities

Blank (Excipient)



At 0.1 ng/ml NDBA: Interferent to NDBA was <20%
 At 0.5 ng/ml NDBA: Insignificant

Standard at 0.5 ng/ml or 17 ppb (ng/g API)



Calibration Curves for All Compounds in Excipient

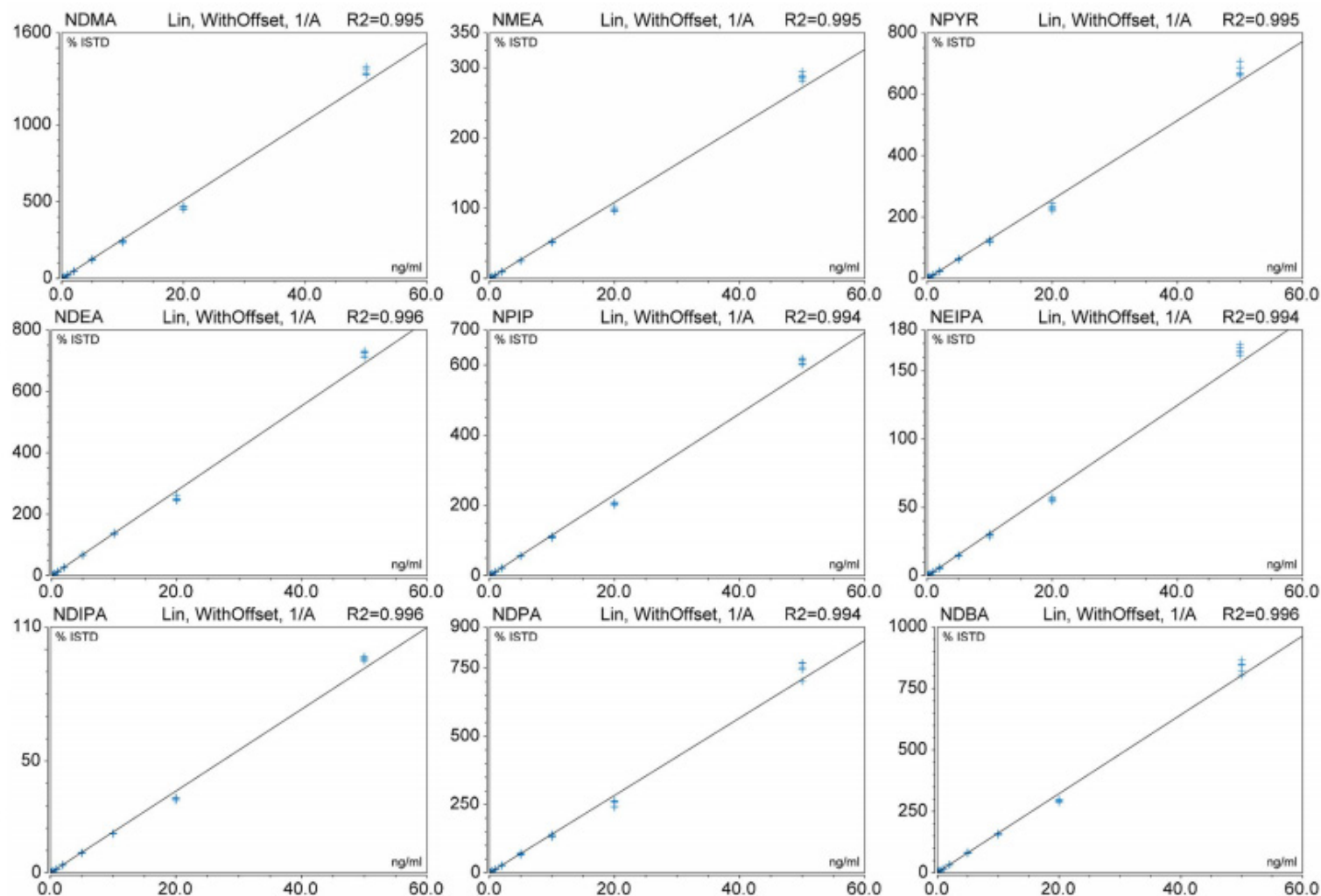


Figure 2. Calibration curves for all compounds in excipient

The newly recommended detection limits from the US FDA : < 30 ppb for a total of 7 nitrosamines

	LOD		LLOQ		Linearity
	ng/mL	PPB	ng/mL	PPB	
NDMA	0.2	6.8	0.2	6.8	LLOQ – 50
NMEA	0.2	6.8	0.2	6.8	
NPYR	0.2	6.8	0.2	6.8	
NDEA	0.1	3.4	0.1	3.4	
NPIP	0.2	6.8	0.2	6.8	
NEIPA	0.5	17.0	0.5	17.0	
NDIPA	0.1	3.4	0.1	3.4	
NDPA	0.1	3.4	0.1	3.4	
NDBA	0.1	3.4	0.5	3.4	

LOD defined as within 20% accuracy, and 15% RSD.
 LOQ defined as within 15% accuracy, and 15% RSD.
 PPB is calculated based on 30 mg/mL of drug substance and product extract.

Recovery, Accuracy, and Precision

Sample recovery and reproducibility

Spiked 2 and 5 ng/mL standard in blank excipient matrix before and after the extraction process (n=5)

	2 ng/mL		5 ng/mL	
	% Recovery	% RSD	% Recovery	% RSD
NDMA	95	5.5	99	3.2
NMEA	96	6.3	98	1.2
NPYR	99	7.4	100	3.9
NDEA	97	2.4	98	3.2
NPIP	96	2.6	99	5.5
NEIPA	99	8.4	98	2.2
NDIPA	99	2.8	98	2.2
NDPA	95	4.4	96	8.4
NDBA	99	2.9	101	1.5

The recovery for all nitrosamines during the extraction process was between 95 and 101%, and the reproducibility of the replicate injections was within 10% RSD

Accuracy and precision

0.5 ng/mL (17 ppb) check standard in excipient (n=5)

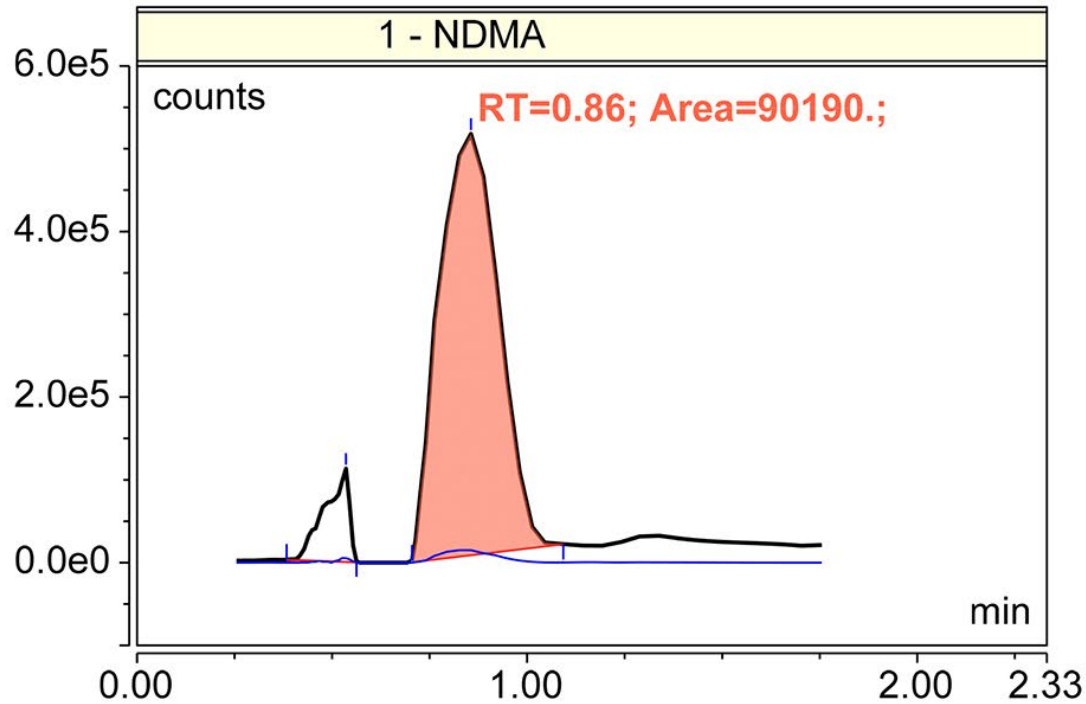
	%Accuracy	%RSD
NDMA	92	10.9
NMEA	95	4.2
NPYR	92	2.7
NDEA	95	3.4
NPIP	98	5.1
NEIPA	93	7.8
NDIPA	97	2.9
NDPA	96	3.0
NDBA	97	4.1

All target nitrosamines at 17 ppb (0.5 ng/mL) could be detected and quantified with high accuracy

Nitrosamine Impurity Levels in Ranitidine Drug Product

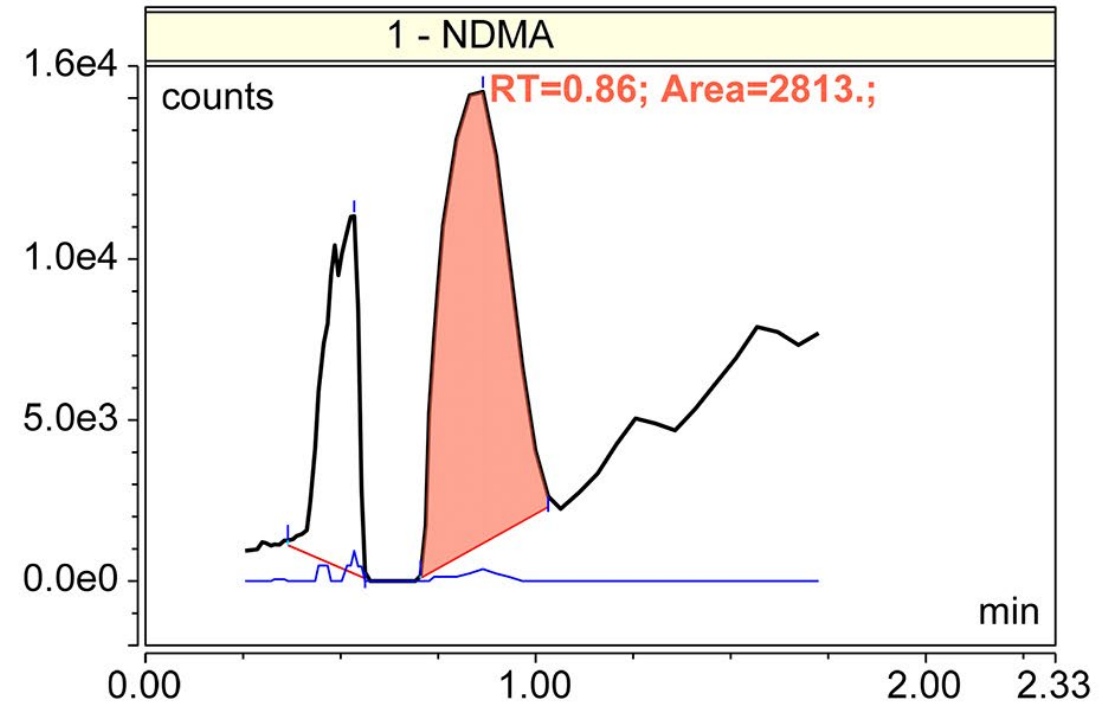
NDMA in ranitidine drug substance and tablet. Data processed with a mass tolerance setting of 3 ppm.

a) Ranitidine drug substance



The measured amount of **NDMA** in 30 mg/mL ranitidine drug substance exceeded the upper limit of calibration and was estimated to be more than 7 ppm

b) Ranitidine drug tablet



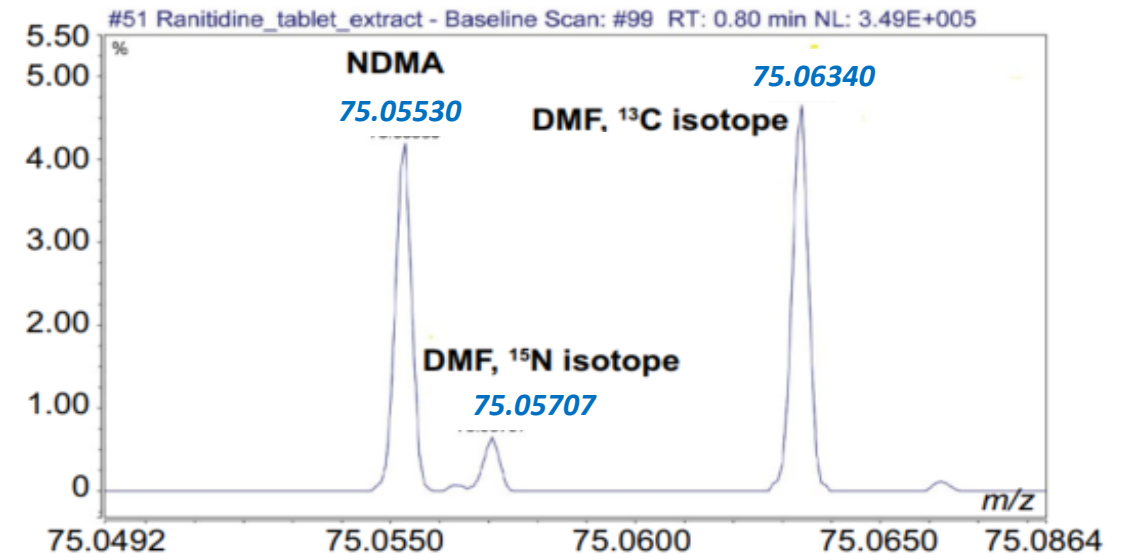
the measured amount of **NDMA** in 300 mg ranitidine tablet was 82 ppb

Both exceeded the acceptable regulatory limit !

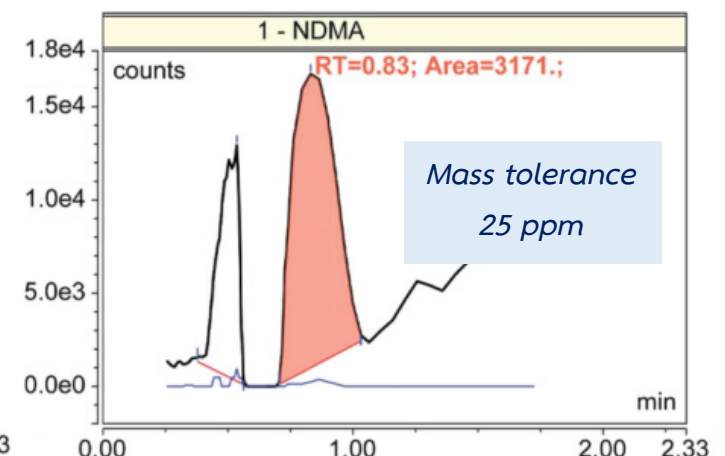
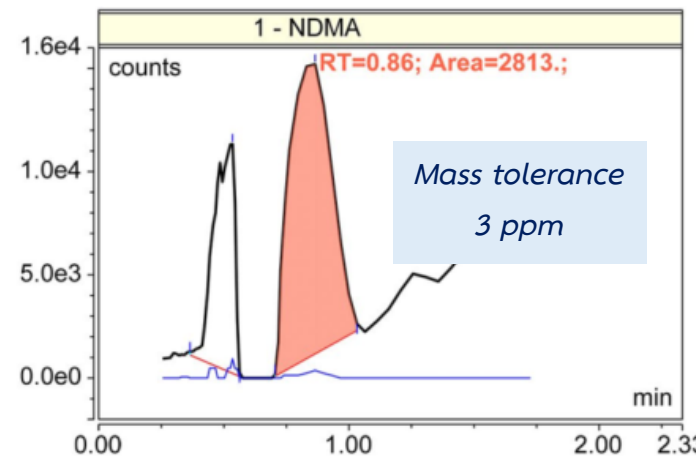
Nitrosamine Impurity Levels in Ranitidine Drug Product

- Mass spectrum of ranitidine drug tablet showing co-elution of NDMA and N,N-dimethylformamide (DMF), which could cause overestimation of NDMA when inadequate mass tolerance setting was used for data processing
- The mass difference between NDMA and DMF ¹⁵N isotope is only 21 ppm
- DMF in pharmaceuticals is allowed to be up to 880 ppm as per the ICH Q3C(R6) guideline.
- A minimum resolution setting of 45,000 and a maximum mass tolerance of 15 ppm are required to prevent overestimation of NDMA when quantifying NDMA using the monoisotopic ion.

Spectrum of ranitidine drug tablet sample containing NDMA and DMF with a resolution setting of 120,000



Overestimation of NDMA when processing the data with mass tolerance set at 25 ppm; the resultant quantitation is 13% higher as compared with results obtained at 3 ppm



- A rapid, highly selective, and sensitive method was developed using the Acclaim Polar Advantage II column, Vanquish Horizon UHPLC system and Orbitrap Exploris 120 mass spectrometer for detection and quantitation of nine nitrosamines in commercially available ranitidine drug products.
- By combining the robust and reproducible chromatography with the 120,000 mass resolving power, fast scanning speed, and sub-ppm mass accuracy of the Orbitrap Exploris 120 system, the resultant method can provide reliable and confident quantitation of nine nitrosamine impurities to meet the September 2020 US FDA regulatory acceptance limits.

thermoscientific

APPLICATION NOTE

73814

HRAM LC-MS method for the determination of nitrosamine impurities in drugs

Authors: Hao Yang, Thermo Fisher Scientific, San Jose, CA, US

Jon Bardsley, Thermo Fisher Scientific, Hemel Hempstead, UK

Min Du, Thermo Fisher Scientific, Boston, MA, US

Olaf Scheibner, Thermo Fisher Scientific, Dreieich, Germany



Keywords: Nitrosamines, NDMA, APCI, high resolution accurate mass, mass spectrometry, Orbitrap Exploris 120, Chromatography Data System, compliance-ready, generic drugs, impurities, genotoxic impurities, ranitidine, excipient, tSIM, tMS²

Application benefits

- Detection and quantification of nine nitrosamines with a single liquid chromatography-high resolution accurate mass (HRAM) mass spectrometry method
- Quantitation of nitrosamine impurities in ranitidine drug substance and product below the daily acceptable intake level, that meets the requirements of FDA regulatory guidelines

- Use of Thermo Scientific™ Chromeleon™ Chromatography Data System (CDS) software for both data collection and processing in a 21 CFR 11 compliant environment with full data integrity and security capabilities for cGMP facilities

Goal

To demonstrate fast, highly sensitive quantitation of nine nitrosamines with a Thermo Scientific™ Orbitrap Exploris™ 120 mass spectrometer, and the use of the LC-MS method to measure nitrosamine impurities in commercially available ranitidine drug substances and products

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