



Future-proof LC-MS solution for nitrosamine analysis: quantitative fidelity meets sustainability

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This technical note demonstrates a method for quantitation of 6 nitrosamines in a model API, esomeprazole, using the novus V55 system [Figure 1].¹ Limits of quantitation [LOQs] as low as 0.08 ng/mL were achieved with reliable quantitative performance offered by the novus V55 system in a new, compact design.

Nitrosamines are a major concern in the pharmaceutical industry, given their potential for carcinogenic health effects.² LC-MS has been widely used for nitrosamine testing, given the sensitivity, reproducibility, and selectivity it provides. Moreover, LC-MS methods have offered adequate separation between the nitrosamines and API, thereby increasing quantitative confidence when measuring multiple impurities.

Quality control [QC], process, and manufacturing laboratories often require multiple MS systems to conduct routine, high-volume testing daily. As a result, sensitive, selective, reliable, and sustainable analytical methods are essential for accurate quantitation of nitrosamines to ensure drug product safety throughout the development lifecycle.

Key benefits for nitrosamine analysis using the novus V55 system

- **Low level of quantitation:** Reach LOQs as low as 0.08 ng/mL for the quantitation of nitrosamines in esomeprazole API.
- **Robust analytical performance:** Achieve accurate and highly reproducible [%CV <15] quantitative performance at all concentration levels.
- **Baseline separation:** Good chromatographic separation of the esomeprazole API and nitrosamines was demonstrated, enabling reliable quantitation.
- **Small footprint without compromising quantitative fidelity:** Reach optimal bioanalytical quantitative performance using the most compact triple quadrupole mass spectrometer in its class.

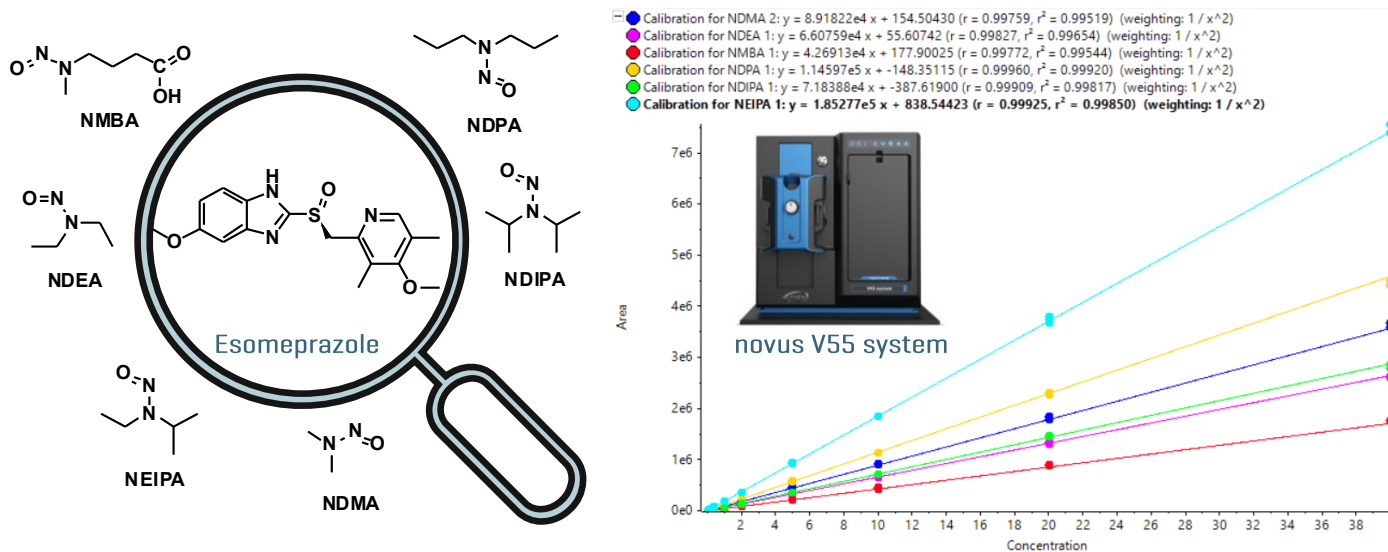


Figure 1. LC-MS/MS method for the determination of 6 nitrosamines in esomeprazole API. Calibration curves demonstrate the measurement of 6 nitrosamines across a wide range of concentrations using the novus V55 system. Good linearity was observed with a coefficient of determination (r^2) of >0.995 for all measured nitrosamines. A weighing factor of $1/x^2$ was applied for all 6 nitrosamines.

- **Streamlined data management:** Easily acquire, manage, and process data using SCIEX OS software, a 21 CFR Part 11-compliant platform.

Introduction

Nitrosamine analysis has commonly relied on mass spectrometry-based methods. The regulatory thresholds for nitrosamine testing are stringent, necessitating sensitive and reliable detection. In addition to threshold requirements, QC and analytical laboratories are continually expanding efforts to reduce energy consumption, instrument footprint, and overall operational costs.

In this study, 6 nitrosamines including NDMA, NDEA, NMBA, NDPA, NDIPA, and NEIP were evaluated in esomeprazole API. Analysis was performed using the novus V55 system, which enables reliable monitoring of impurities and quantitative performance. Additionally, the novus V55 system empowers pharmaceutical laboratories with enhanced energy efficiency solutions.

Methods

Standard preparation: Nitrosamine mix in methanol was purchased from Accustandard [Catalog # FDA-001S]. Calibration points ranging from 0.08 to 40 ng/mL were prepared in a 95:5 [v/v] water/methanol mixture.

Sample preparation: A concentrated esomeprazole API stock was prepared in methanol and subsequently diluted to 80 mg/mL in water. The solution was vortexed, briefly centrifuged, and transferred to a 1.5 mL centrifugal filter unit. The sample was centrifuged at 12,000 rpm for 5 minutes. Supernatant was transferred to a new tube. For the recovery experiment, API stock was diluted to 40 mg/mL using 95:5 [v/v] water/methanol and used as API blank. Spiked samples were prepared by spiking 1 ng/mL nitrosamine mix in API blank.

Chromatography: Sample separation was performed using a Shimadzu X3 system at a flow rate of 1.2 mL/min on a [Phenomenex Biphenyl column \(4.6 x 150 mm, 2.6 μm, 100 Å\)](#). The column temperature was maintained at 40°C. A 26-minute gradient was run using 0.1% formic acid in water as mobile phase A and 0.1% formic acid in methanol as mobile phase B [Table 1]. A high organic wash was applied when running the API

samples. An injection volume of 15 μL was used for analysis. A 90:10 [v/v] methanol/water mixture was used as the needle wash solvent.

Table 1. LC gradient conditions.

Time [min]	Mobile phase A [%]	Mobile phase B [%]
0.0	95	5
1	95	5
5	60	40
17	60	40
23	20	80
24.5	20	80
24.6	2	5
26	90	5

UV chromatography: UV data were collected using a Shimadzu UV-SPD-40 equipped with a D2 lamp. The API detection wavelength was set to 300 nm.

Mass spectrometry: Analysis was performed on the [novus V55 system](#) [SCIEX]. The optimized source and gas parameters are listed in Table 2, and the MRM parameters are discussed in Table 3. For all transitions, the dwell time and EP values were set to 40 msec and 10 V, respectively.

Table 2. Source and gas parameters.

Parameter	Value
Polarity	Positive
Source	OptiFlow APCI
Ion source gas 1	45 psi
Curtain gas	30 psi
Source temperature	375°C
Nebulizer current	5.0 μA
CAD gas	8

Table 3. MRM parameters applied for quantitation.

ID	Precursor ion [m/z]	Fragment ion [m/z]	CE [V]	CXP [V]	DP [V]
NDMA 1	75.0	58.0	17	6	70
NDMA 2	75.0	43.1	21	7	70
NDEA 1	103	75.1	20	9	70
NDEA 2	103	47.1	22	7	70
NMBA 1	147	117	9	14	60
NMBA 2	147	86.9	15	11	60
NDPA 1	131.1	89.0	12	10	65
NDPA 2	131.1	43.1	20	10	65
NDIPA 1	131.1	89.0	10	10	65
NDIPA 2	131.1	43.1	25	7	65
NEIPA 1	117	75.0	15	12	75
NEIPA 2	117	43.1	19	10	75
API	346.1	198.1	14	7	70

Data processing: Data collection and analysis were performed using SCIEX OS software, version 4.3. Peaks were integrated using the MQ4 algorithm, and a weighting of $1/x^2$ was used for nitrosamine quantitation.

Quantitative performance on the novus V55 system

Baseline separation was achieved between the esomeprazole API and the 6 nitrosamines [Figure 2]. The UV chromatogram shows the analysis of esomeprazole API at 300 nm, while the XIC traces show the elution of the nitrosamine compounds. The retention time of the esomeprazole API was 18.99 min, indicating baseline separation relative to the nitrosamine peaks. Three impurity peaks were observed in the UV chromatogram of the esomeprazole API between 7 and 8.5 min. None of the nitrosamine compounds were detected at that retention timeframe. Therefore, no impact on the quantitation of nitrosamine compounds was observed.

Figure 3 shows the XICs of the blank and the LOQ levels for all 6 nitrosamines. An LOQ of 0.2 ng/mL was reached for NDMA, NDPA, and NDIPA, while for NDEA, NMBA, and NEIPA, an LOQ of 0.08 ng/mL was achieved. The blank XICs did not indicate any interferences at the retention times of the nitrosamine analytes.

The evaluated calibration curve range for NDMA, NDPA, and NDIPA was 0.2 ng/mL to 40 ng/mL, while NDEA, NMBA, and NEIPA were evaluated from 0.08 ng/mL to 40 ng/mL [Figure 1]. Replicate analysis ($N = 3$) was performed at each concentration level. Overall, excellent linearity was observed with r^2 values of >0.995 and a weighting factor of $1/x^2$ [Figure 1].

Analytical performance was evaluated for accuracy and precision. The accuracy of the calculated mean was expected to be between 80% and 120% at the LOQ and between 85% and 115% at higher concentrations. The %CV of the calculated mean for each concentration was expected to be $<20\%$ at the LOQ.³

Assay accuracy at the LOQ levels was within $\pm 4\%$ of the actual concentration, and %CV was <15 . The calculated percentage accuracy and %CV values were within the acceptance criteria at each concentration level [Table 4].

Recovery was evaluated at 1 ng/mL in 6 replicates. Overall, recovery across 6 nitrosamines was $>98\%$ with a %CV <7 , indicating a highly reliable and reproducible assay for nitrosamine analysis in esomeprazole API [Table 5].

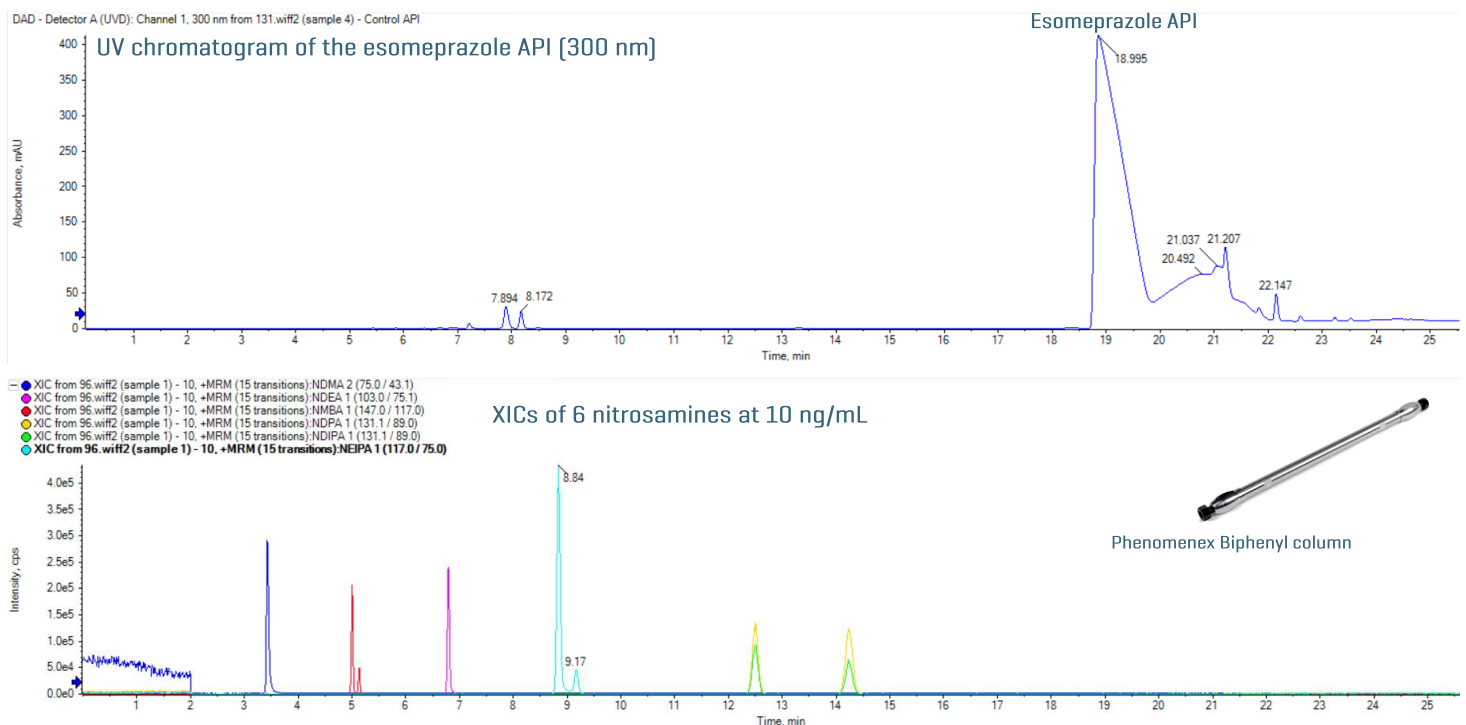


Figure 2. Baseline separation was achieved between the 6 nitrosamines and the esomeprazole API. UV chromatogram of the esomeprazole API (top; collected at 300 nm) and the extracted ion chromatograms (XICs) of the 6 nitrosamines (bottom; collected at 10 ng/mL) are displayed.

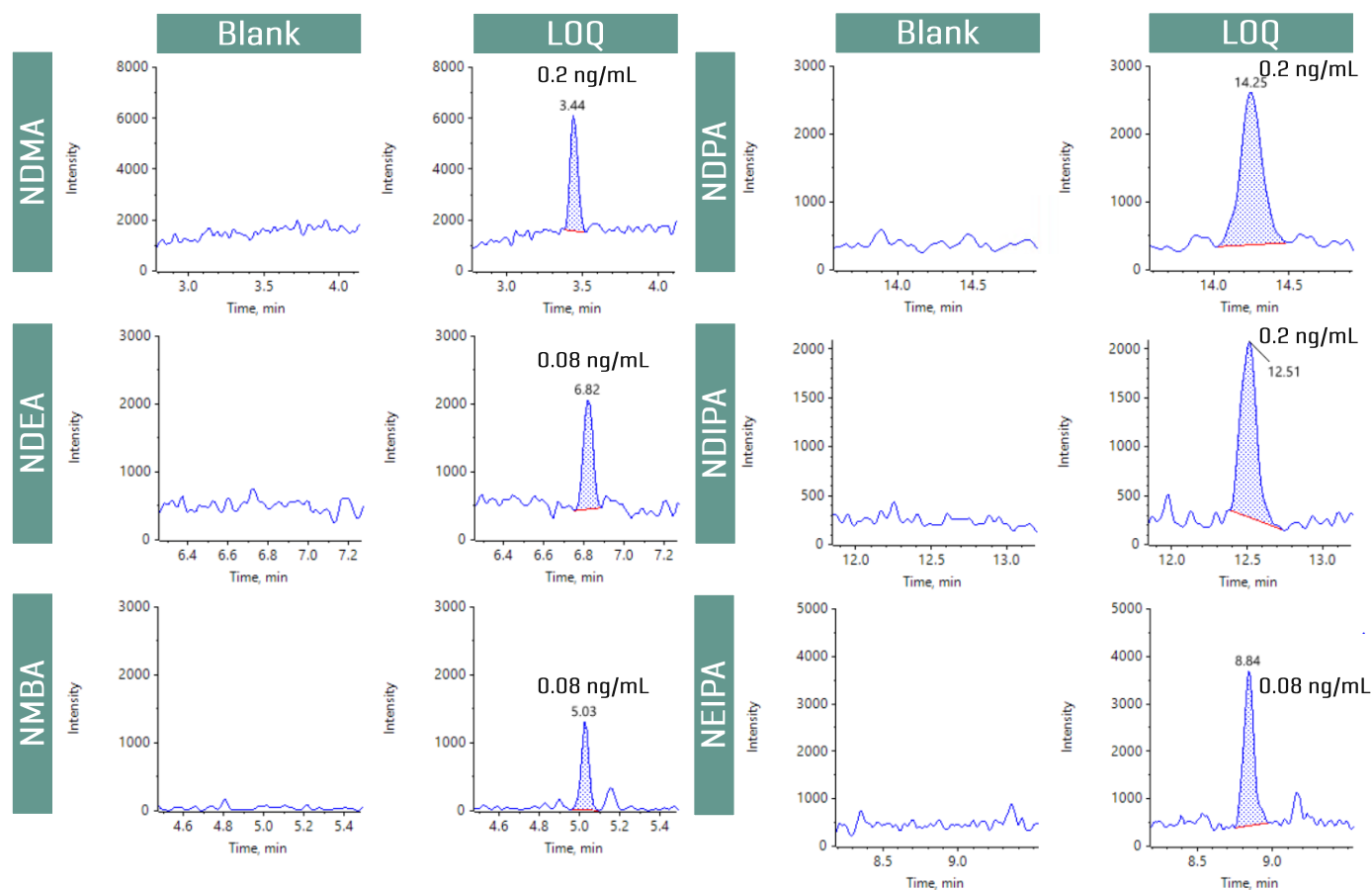


Figure 3. Representative XICs of the blank and LOQ levels. LOQ of 0.2 ng/mL was reached for NDMA, NDPA, and NDIPA, while an LOQ of 0.08 ng/mL was achieved for NDEA, NMBA, and NEIPA. No interferences were observed in the blank samples.

Table 4. Quantitative performance of the evaluated nitrosamines. Average accuracy and precision were calculated across all concentration levels in triplicate.

Concentration [ng/mL]	NDMA		NDEA		NDPA		NDIPA		NMBA		NEIPA	
	Avg. Acc.	Avg. %CV	Avg. Acc.	Avg. %CV	Avg. Acc.	Avg. %CV	Avg. Acc.	Avg. %CV	Avg. Acc.	Avg. %CV	Avg. Acc.	Avg. %CV
0.08			99.7	2.47					102.	7.67	100.	2.83
0.2	104.	11.8	101.	12.7	4.09	99.3	9.87	98.2	96.0	8.06	98.7	10.1
0.4	92.7	6.51	98.8	8.71	3.68	101.	3.32	104.	103.	8.60	101.	3.46
1	95.7	2.66	101.	4.08	1.72	99.5	3.00	101.	91.7	3.08	100.	1.99
2	101.	0.685	100.	1.51	1.46	101.	3.41	98.4	100.	2.12	98.4	0.143
5	100.	2.31	98.7	1.78	1.54	101.	1.74	99.8	99.4	2.22	100.	0.569
10	102.	0.899	101.	2.66	0.233	98.9	1.13	99.1	102.	3.97	99.7	0.507
20	102.	1.77	100.	1.83	1.06	100.	1.04	101.	104.	1.56	101.	1.66
40	102.	1.77	99.0	0.961	1.40	97.3	1.22	98.4	104.	0.04	101.	1.57

Blue font is used to denote LOQ levels

Table 5. Recovery was evaluated at 1 ng/mL in 6 replicates.

Compound	Recovery [%]	%CV
NDMA	98.2	6.19
NDEA	99.8	3.22
NMBA	109	4.30
NDPA	100	2.93
NDIPA	101	2.25
NEIPA	103	3.53


Compliance-ready SCIEX OS software

Equivalent SCIEX OS software capabilities for nitrosamine analysis can be executed on the novus V55 system, ensuring high fidelity when performing method transfers while retaining critical compliance features.

SCIEX OS software is a closed system and requires records and signatures to be stored electronically, meeting the regulations outlined by 21 CFR Part 11. SCIEX OS software can open raw data files from any visible storage location within a closed network by using designated processing workstations.

Figure 4 illustrates the features of SCIEX OS software used to monitor the audit trail, acquire and process data, and configure user access. The audit trail feature enables users to audit critical user actions and locks in data integrity.

The Central Administrator Console [CAC] feature allows users to centralize acquisition and processing using a single platform to maximize efficiency for multi-instrument laboratories, independent of compliance standards. The configuration



Audit Trail

Easily search and filter for specific high-risk events in audit trail viewer. Built-in data integrity features allow you to tailor each functionality specifically to meet compliance needs and data security requirements.

Filter Audit Trail

Find instances where


Sample Name Clear

Reason Contains Standard

<No Filter> Contains manual integration

And where time and date are: From To

Ok Cancel



Central Administration

Users can manage groups, role definitions, workstations and projects across all systems using the SCIEX OS software Central Administrator Console (CAC). It supports all regulated and non-regulated compliance standards.

Centralized Management

Manage Workgroups and Assignments

Method Developers

Reviewers

Use Central Settings for Projects


Assign or Unassign Project Role from Project Pool

Project Role Directory

Confirm Change Events

Change Events

E-Signature



Configuration

Assign users and access to administrator, method developer, analyst and reviewer roles under the audit trail module. Easily customize the role and specify level of access.

Users Roles Security

User Roles and Permission Categories

Permission	Administrator	Method Developer	Analyst	Reviewer
Batch				
Submit unlocked methods	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Open	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Save as	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Submit	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Save	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Save lock reference table	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Add data sub-folders	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Configure Decision Rules	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Configuration				
General tab	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
General: change regional setting	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
General: full screen mode	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
LIMS communication tab	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Queue tab	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Queue: instrument life time	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Queue: max. number of acquired samples	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Queue: other queue settings	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>

Duplicate a User Role

Create a new user role based on an existing role.

Existing user role: Administrator

New role name:

New role description:

Ok Cancel

Figure 4. Features of the SCIEX OS software for monitoring user access and evaluating the audit trail. The audit trail view allows users to filter for high-risk events easily and enables data integrity features to meet compliance requirements. The software features a Central Administrator Console [CAC] to manage users and groups, role definitions, workstations, and projects across all systems. The CAC feature supports both regulated and non-regulated compliance standards. The configuration module enables users to quickly set up roles and levels of access for the administrator, method developer, analyst, and reviewer levels.

module allows users to assign roles and access as the administrator, method developer, analyst, and reviewer.

Conclusions

- An LOQ of 0.2 ng/mL was demonstrated for NDMA, NDPA, and NDIPA, while an LOQ of 0.08 ng/mL was reached for NDEA, NMBA, and NEIPA.
- Quantitative performance was demonstrated with accurate and highly reproducible [%CV <15] results on the novus V55 system.
- Excellent linearity was achieved with $r^2 > 0.995$. NDMA, NDPA, and NDIPA were evaluated between 0.2 ng/mL to 40 ng/mL, while NDEA, NMBA, and NEIPA were evaluated from 0.08 ng/mL to 40 ng/mL.
- Baseline separation was achieved between the 6 measured nitrosamine compounds and the esomeprazole API, enabling accurate quantitation.
- Overall, recovery was >98% when evaluating all 6 nitrosamines at 1 ng/mL with a %CV <7, demonstrating high quantitative fidelity.
- Maintain quantitative rigor and lower operational costs with the novus V55 system, the most compact triple quadrupole mass spectrometer in its class.
- Data management and compliance-readiness (21 CFR Part 11) features were shown using the SCIEX OS software to support nitrosamine quantitation on the novus V55 system.

References

1. The SCIEX novus V55 system. [SCIEX brochure, MKT-38393-A](#).
2. Control of Nitrosamine Impurities in Human Drugs. [US Food and Drug Administration Guidance for Industry, September 2025](#).
3. [Bioanalytical Method Validation, May 2018](#).

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