

Development of a Sensitive Headspace Gas Chromatography–Mass Spectrometry Method for the Simultaneous Determination of Nitrosamines in Losartan Active Pharmaceutical Ingredients

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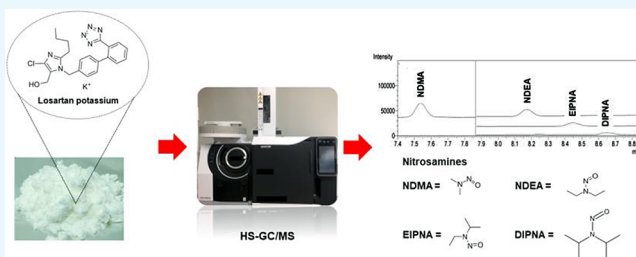


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ABSTRACT: Nitrosamine impurities in angiotensin II receptor antagonists (sartans) containing a tetrazole group represent an urgent concern for active pharmaceutical ingredient (API) manufacturers and global regulators. Regarding safety, API manufacturers must develop methods to monitor the levels of each nitrosamine impurity before individual batch release. In this study, we developed and validated a sensitive, selective, and high-throughput method based on headspace gas chromatography–mass spectrometry (HS-GC–MS) for the simultaneous determination of four nitrosamines in losartan potassium API with simple sample preparation. *N*-Nitrosodimethylamine (NDMA, m/z 74), *N*-nitrosodiethylamine (NDEA, m/z 102), *N*-nitrosoethylisopropylamine (EIPNA, m/z 116), and *N*-nitrosodiisopropylamine (DIPNA, m/z 130) levels were quantified using an electron impact, single quadrupole mass spectrometer under a selected-ion-monitoring acquisition method. The method was validated according to the Q2(R1) ICH guidelines. The calibration curves of the assay ranged from 25 to 5000 ng/mL with limits of quantitation of 25 ppb for NDMA and NDEA and 50 ppb for DIPNA and EIPNA. The accuracy of the developed method ranged from -7.04% to 7.25% , and the precision %CV was ≤ 11.5 . Other validation parameters, including specificity, stability, carryover, and robustness, met the validation criteria. In conclusion, the developed method was successfully applied for the determination of nitrosamines in losartan potassium APIs.



1. INTRODUCTION

Angiotensin II receptor antagonists (sartans) such as valsartan, losartan, irbesartan, and olmesartan have been widely used to treat hypertension, diabetic nephropathy, and congestive heart failure.^{1–3} In July 2018, the United States Food and Drug Administration (US FDA)⁴ and the European Medicines Agency (EMA)⁵ initially announced a recall of valsartan tablets because of contamination by the nitrosamine *N*-nitrosodimethylamine (NDMA). Subsequently, other nitrosamine impurities, including *N*-nitrosodiethylamine (NDEA), *N*-nitrosodiisopropylamine (DIPNA), and *N*-nitrosoethylisopropylamine (EIPNA), have been found in several sartan drugs including losartan, irbesartan, and candesartan.⁶ In addition, other nitrosamines, such as *N*-nitrosomethylphenylamine (NMPHA), *N*-nitrosodibutylamine (NDBA), and *N*-nitromethylbutyric acid (NMBA), have been detected in sartan APIs.⁷ These nitrosamine impurities are classified as probable human carcinogens^{3–7} and are believed to be present in finished sartan products because of the manufacturing process for sartan active pharmaceutical ingredients (APIs). Sartans containing a tetrazole ring carry a potential risk of nitrosamine contamination because the excess amount of sodium nitrite used to quench unreacted sodium azide in the synthesis of

sartan APIs can react with residual secondary amines, which inadvertently generates nitrosamines.^{8,9} One example of contaminating secondary amines is dimethylamine, which may be present in the dimethylformamide solvent. Nitrosative dealkylation of triethylamine and diisopropylethylamine is a possible pathway to generate residual secondary amines.⁹

Because of possible nitrosamine formation during API manufacturing, it is essential to monitor nitrosamine impurities in sartan APIs before individual batch release. Since July 2018, the US FDA, EMA, and other national regulatory agencies have released various methods for the determination of nitrosamine contaminants in sartan APIs based on gas chromatography coupled to mass spectrometry (GC–MS) or tandem mass spectrometry (GC–MS/MS), high-performance liquid chromatography with ultraviolet detection (HPLC–

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UV), and liquid chromatography coupled to mass spectrometry (LC–MS) or tandem mass spectrometry (LC–MS/MS).^{10–16} The French National Agency for Medicines and Health Products Safety developed an HPLC–UV method to determine the NDMA content in valsartan APIs and drug products.¹⁰ The German Official Medicines Control Laboratory published a validated LC–MS/MS method to analyze NDMA levels in valsartan products.¹¹ The technique had the linearity range of 0.1–3.0 ppm and a limit of quantification (LOQ) of 0.2 ppm, which is slightly less than the interim limit at 0.3 ppm for NDMA in valsartan. The Public Analyst's Laboratory (Galway, Ireland) developed a headspace gas chromatography–mass spectrometry (HS–GC–MS) method for the quantitative analysis of NDMA in valsartan APIs and drug products without reporting the detection and quantitation limits.¹² In 2019, the US FDA initially reported an HS–GC–MS method for NDMA determination in valsartan APIs and drug products with an LOQ at the interim limit of NDMA at 0.3 ppm.¹³ The US FDA reported a simultaneous HS–GC–MS analysis method for NDMA and NDEA in valsartan with improved sensitivity (LOQs of 0.10 and 0.05 ppm, respectively).¹⁴ Other nitrosamines, including DIPNA and EIPNA, have recently been identified as impurities in the sartan drug class by the EMA.⁹ In addition to the US FDA and EMA report, at least 12 nitrosamines were reported in an application note published by the Taiwanese FDA.^{15,16} Among the 12 nitrosamines,¹⁶ only three nitrosamines including NDMA, NDEA, and NMBA were found in valsartan, losartan, and irbesartan. The evidence basis of the recall reports of nitrosamine contamination from US FDA and EMA press releases suggests that NDMA, NDEA, NMBA, DIPNA, and EIPNA are most commonly found in sartans. Therefore, it is essential to develop a method for detecting these potential nitrosamine contaminants in sartans. The US FDA recently proposed the simultaneous analysis of NDMA, NDEA, DIPNA, and EIPNA using a GC–MS triple quadrupole coupled with either sample direct injection or a headspace autosampler.^{16–18} Nonetheless, the published GC–MS methods were developed and validated for valsartan APIs and drug products with particular acceptable sensitivity.

In addition to valsartan, losartan is one of the most commonly used sartans,¹⁹ and it has been extensively reported to contain nitrosamine impurities above the US FDA²⁰ and EMA^{6,21,22} limit criteria. The US FDA released an application note on the analytical method for determining six nitrosamines in losartan using liquid chromatography coupled to high-resolution, accurate mass (HRAM) mass spectrometry (LC–HRMS).²³ Although HRAM mass spectrometers provide better resolution and selectivity for NDMA impurity analysis than single-quadrupole, triple-quadrupole, and time-of-flight mass spectrometers,²⁴ the HRAM system suggested by the US FDA is not commonly used in routine quality control laboratories. Another concern is that directly injecting samples into GC or LC–MS systems without sample cleanup may lead to machine contamination from high sample matrix loading, resulting in a loss of sensitivity and robustness. Headspace injection is a simple sample preparation technique for preventing high levels of sample matrix contamination in a mass spectrometer. Therefore, HS–GC–MS was developed and optimized for volatile nitrosamine impurity analysis with high sample matrix loading. Literature reviews, some application notes from the FDA in several countries,^{10–16} and a few published studies²⁵ have revealed methods for

nitrosamine determination, and such methods require further validation for specific purposes by users.

Based on US FDA recall events, there are seven reported nitrosamines, including NDMA, NDEA, NMBA, DIPNA, EIPNA, NDBA, and NMPHA. Three of those nitrosamines, including NDMA, NDEA, and NMBA, have caused most often recalled in losartan, while DIPNA and EIPNA have insufficient evidence. Moreover, the recall caused by NDBA and NMPHA contaminations has rarely occurred. However, the number of nitrosamine contamination may be relevant to the manufacturing process and different sartans. In this work, we focused on determining the potential contamination risk of four nitrosamines, e.g., NDMA, NDEA, DIPNA, and EIPNA, in losartan raw materials using HS–GC–MS. Although NMBA is a potential contaminating nitrosamine, it could not be included in this analytical method because of volatile property limitations. This work is the first report on the development and validation of a robust, simple, sensitive, and specific HS–GC–MS method for the simultaneous determination of nitrosamines in losartan APIs in accordance with the Q2(R1) ICH guidelines.²⁶ The method proved applicable for identifying nitrosamine contamination in different batches of losartan APIs.

2. RESULTS AND DISCUSSION

2.1. Method Development. The stationary phase, column length, film thickness, and column diameter have a large impact on the retention and peak shape of the analytes. Based on the trial-and-error methodology with the literature review, the optimal separation of nitrosamines was achieved when the capillary surface was modified by polyethylene glycol residues at the length of 30 m.^{29–31} The optimal chromatographic separation and retention were achieved using a column dimension of 30 × 0.32 mm² i.d. and a film thickness of 1 μm coated with polyethylene glycol stationary phase. The temperature programming is also a very important parameter to influence peak separation. Yang et al. stated that ¹⁵N-DMF representing residual solvents in API has the same molecular mass of 74, which must be discriminated from NDMA using an HRMS in cases of poor chromatographic separation.²⁴ An initial constant temperature at 40 °C for 2 min followed by an increase at a rate of 20 °C can provide the best separation with a resolution greater than 2 between NDMA and ¹⁵N-DMF at the retention times of 7.47 and 7.67 min, respectively, as presented in Figure S1. The temperature of the sample and transfer lines was set above 200 °C, and the temperature of the transfer line should be slightly higher than that of the sample line to prevent a carryover effect from the previous sample. The optimal temperatures of sample and transfer lines were controlled at 230 and 250 °C, respectively. The filament was turned off after all nitrosamines had entered into the mass spectrometer to prevent filament damage from an excessive dimethyl sulfoxide (DMSO) concentration.

The selection of a suitable sample introduction technique requiring no complicated sample preparation is an essential key for GC–MS analysis. Direct and headspace injection techniques are commonly used because they can be fully and easily automated. The concern of the direct injection technique is that the sample matrices can cause variability and inaccuracy of the GC–MS analysis. The injected matrix, which co-elutes with the analyte, can deteriorate the column shelf life and cause mass spectrometer contamination. Additional sample cleanup can resolve these problems, leading

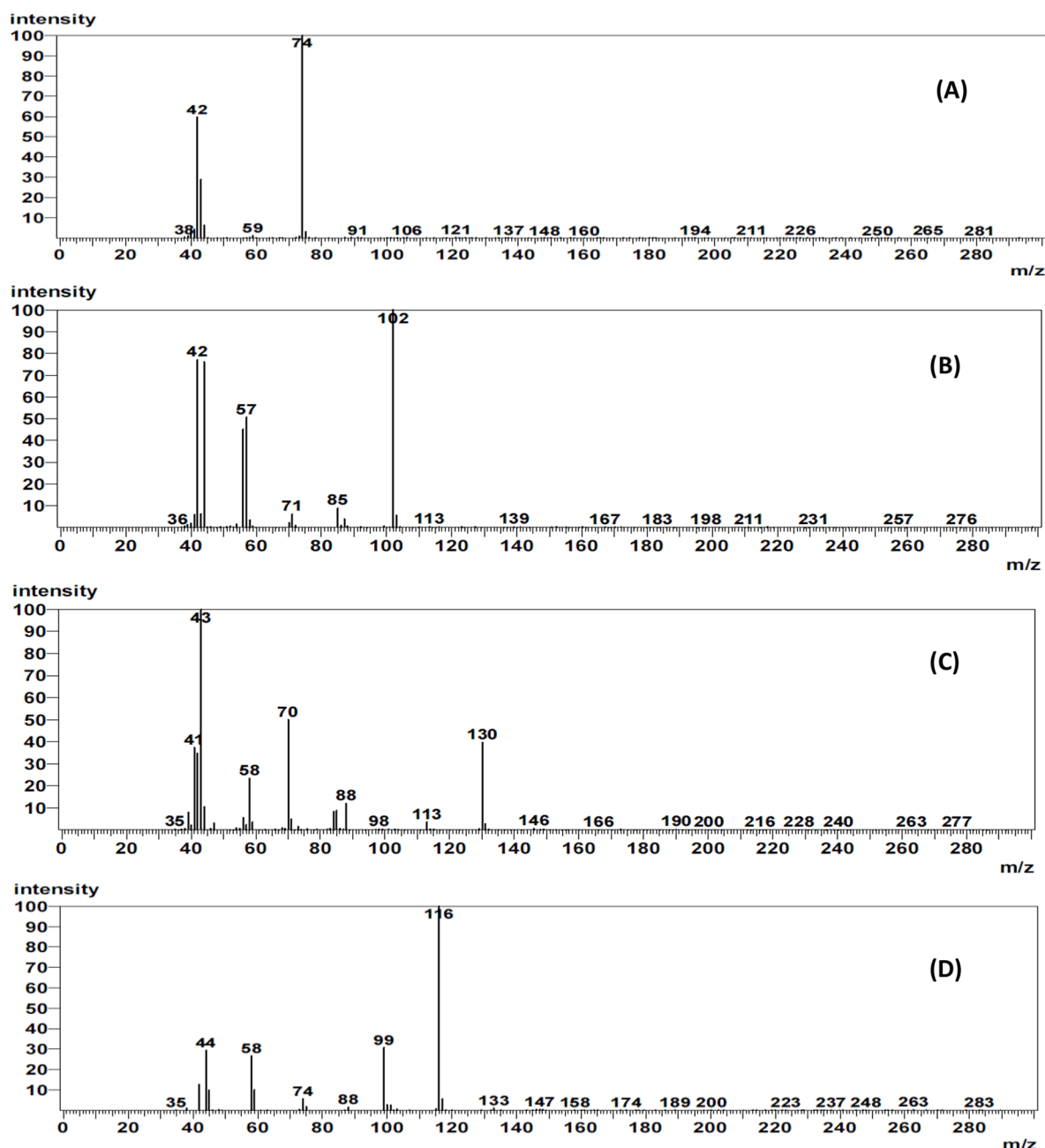


Figure 1. Mass spectra of (A) NDMA, (B) NDEA, (C) DIPNA, and (D) EIPNA.

to a decrease in the matrix interference, an extension of the column lifespan, and a reduction in machine contamination. The samples often need to be preconcentrated before analysis by direct injection due to a deficient level of the target analyte. In this study, the headspace injection technique was chosen because it can avoid loading nonvolatile sample matrices into the column, leading to significantly reduced matrix interference.³² The analyte in the sample solution is vaporized and concentrated during the sample heating process, increasing the sensitivity of the method.³³ Compared to the direct injection, the headspace sampling technique may be inappropriate for thermolabile samples. In addition, reinjection of the pierced sample is not possible if the headspace sampling technique is applied. The headspace condition was optimized to determine the proper headspace oven temperature and incubation period. Typically, the oven temperature should be set below the

boiling point of the solvent. The sensitivity of the method is poor at excessively low incubation temperatures, e.g., 50–100 °C, whereas extremely high incubation temperatures, e.g., 165–180 °C, can generate substantial baseline noise from the solvent. The optimal incubation temperature is achieved at 150 °C. Regarding the incubation time, the highest sensitivity was achieved at an incubation time of 15 ± 1 min. Shorter incubation periods result in lower sensitivity, whereas longer periods do not affect the sensitivity. It is noteworthy that losartan potassium is thermally stable up to 275 °C,³⁴ whereas the autosampler oven was set at 150 °C, indicating that losartan potassium is stable under the autosampler temperature used in this study. Identifying the proper solvent for sample dilution before analysis is challenging during the headspace technique. Low boiling point solvents such as methanol, ethanol, and isopropanol should be avoided because their high

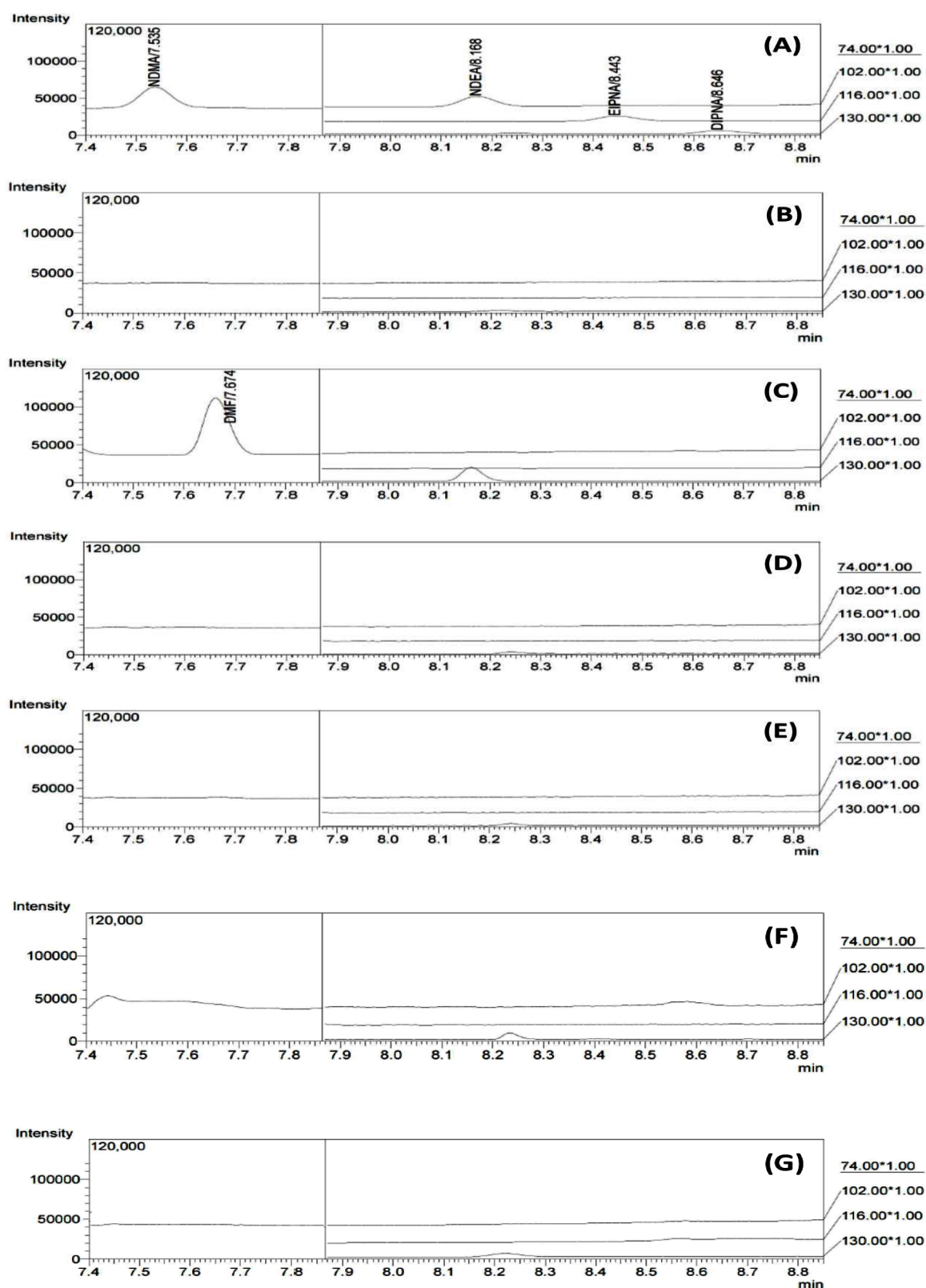


Figure 2. Representative GC chromatograms of (A) the standard mixture of four nitrosamines, (B) DMSO, (C) DMF, (D) methanol, (E) losartan potassium, (F) olmesartan medoxomil, and (G) valsartan sacubitril.

vapor pressure properties can significantly dilute the sample concentration in the headspace. High boiling point solvents such as DMSO, dimethylacetamide, or DMF are often chosen. In this study, DMSO was used, and advantages in terms of volatility and polarity were achieved. More importantly,

losartan and other sartans are soluble in DMSO, and nitrosamine impurities in the APIs can be extracted entirely and dissolved in DMSO. The high boiling point of DMSO also prevents its interference with the nitrosamine peak. The headspace injection technique using DMSO as the sample

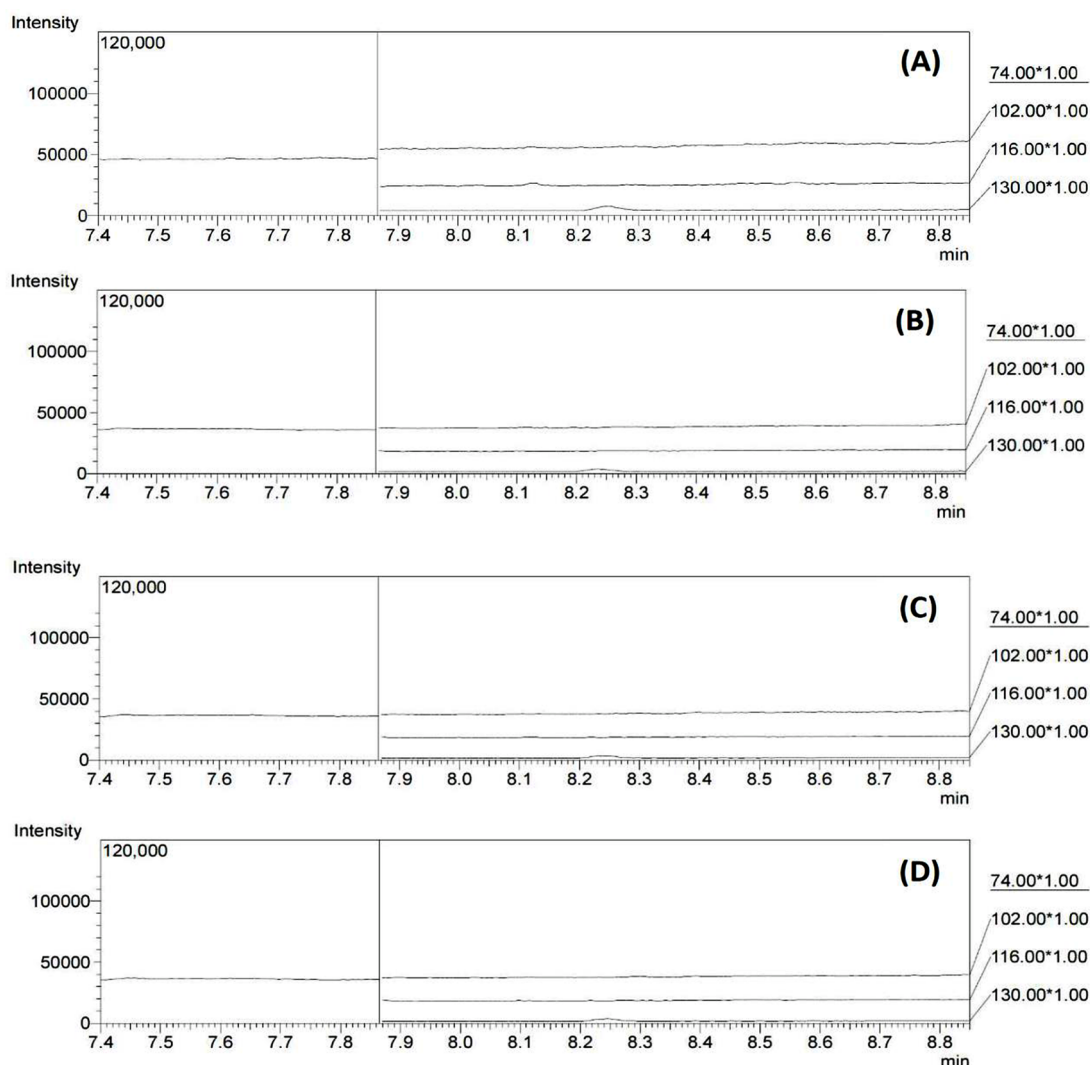


Figure 3. Representative GC chromatograms of the carryover effect on blank solutions after injecting (A) NDMA, (B) NDEA, (C) DIPNA, and (D) EIPNA at a concentration of 5000 ng/mL.

solvent provided sufficient chromatographic performance, and it was, therefore, chosen in this study.

The GC–MS condition for the simultaneous determination of four nitrosamines in losartan was optimized to ensure reliability, selectivity, and sensitivity. Initially, the mass parameters were optimized to achieve the highest sensitivity with a consistent response. NDMA (m/z 74), NDEA (m/z 102), DIPNA (m/z 130), and EIPNA (m/z 116) were tuned on a single-quadrupole mass spectrometer in the electron ionization (EI) mode under the selected ion monitoring (SIM) acquisition method. These particular ions were used for quantification. In addition to the parent ions, the major daughter ions of NDMA (m/z 42), NDEA (m/z 57), DIPNA (m/z 70), and EIPNA (m/z 99) shown in Figure 1 were qualitatively monitored for confirmation. Alternatively, the ion ratio calculation between the parent and daughter ions was also quantitatively used in the confirmatory objective. The GC chromatograms shown in Figure 2A indicate sufficient peak separation with appropriate retention times of 7.535 ± 0.050 , 8.168 ± 0.050 , 8.646 ± 0.050 , and 8.443 ± 0.050 min for NDMA, NDEA, DIPNA, and EIPNA, respectively.

2.2. Method Validation. **2.2.1. System Suitability.** Suitability was evaluated before the validation experiments

and impurity determination to verify the reproducibility and performance of the chromatographic system. A system reproducibility solution containing a mixture of NDMA, NDEA, DIPNA, and EIPNA was prepared at a concentration of 250 ng/mL for each compound by appropriately aliquoting the working standard solution (1000 $\mu\text{g/mL}$). The system suitability solution was injected with five replicates. The system performance was evaluated using the tailing factor of analytes ($T < 2$), and the system reproducibility was assessed by the variation of peak area (%CV < 11%) and retention time (%CV < 2%) to indicate the precision of injections.

2.2.2. Specificity and Selectivity. The specificity of the method for nitrosamine determination focused on the diluents involved in sample preparation, e.g., DMSO and methanol. Contamination of losartan by residual solvents, such as DMF,²⁴ which may affect analysis reliability, was investigated. Furthermore, the selectivity of matrices focused on various sartans. Figure 2A presents a representative GC chromatogram of a standard mixture of four nitrosamines. Figure 2B–D presents the chromatograms of DMSO, DMF, and methanol, respectively. The results revealed the absence of coeluting peaks at the retention times of nitrosamines. The adjacent interfering components from ^{15}N -DMF were observed, but

Table 1. Mean Interday Back-Calculated Standard and Calibration Curve Results ($n = 3$)

compound	nominal concentration (ng/mL)	back-calculated concentration (ng/mL)			mean back-calculated concentration (ng/mL)	% deviation	%CV
		day 1	day 2	day 3			
NDMA	25.0	25.1	24.6	25.9	25.2	0.72	2.58
	50.0	51.5	52.2	48.1	50.6	1.25	4.32
	100.0	98.2	102.6	98.9	99.9	−0.09	2.38
	250.0	250.5	263.4	253.6	255.8	2.32	2.64
	500.0	518.2	485.1	494.3	499.2	−0.17	3.43
	1000.0	995.7	950.9	1040.2	995.6	−0.44	4.48
	5000.0	4864.1	4965.6	4963.1	4931.0	−1.38	1.17
	r^2	0.999	0.998	0.999			
	F_{cal}	12 945.996					
	F_{ANOVA}	2.158×10^{-28}					
	slope	2.158×10^{-28}					
p value	y -intercept	0.425					
NDEA	25.0	24.7	24.6	24.1	24.5	−2.12	1.31
	50.0	49.6	51.0	51.1	50.6	1.17	1.67
	100.0	105.0	102.5	105.9	104.5	4.50	1.69
	250.0	244.8	232.3	264.2	247.1	−1.16	6.51
	500.0	531.4	505.3	489.8	508.8	1.77	4.14
	1000.0	905.8	965.0	999.7	956.9	−4.31	4.96
	5000.0	5075.5	5284.3	4548.5	4969.4	−0.61	7.63
	r^2	0.996	0.997	0.996			
	F_{cal}	17 667.266					
	F_{ANOVA}	1.129×10^{-29}					
	slope	1.129×10^{-29}					
p value	y -intercept	0.941					
DIPNA	50.0	46.2	51.8	51.1	49.7	−0.55	6.11
	100.0	96.7	101.5	99.6	99.3	−0.73	2.45
	250.0	248.1	257.1	241.0	248.7	−0.52	3.25
	500.0	508.4	494.3	520.5	507.7	1.55	2.58
	1000.0	1015.4	965.3	978.2	986.3	−1.37	2.64
	5000.0	5200.9	4954.9	5042.8	5066.2	1.32	2.46
	r^2	0.999	0.999	0.999			
	F_{cal}	794.423					
	F_{ANOVA}	4.573×10^{-15}					
	slope	4.573×10^{-15}					
p value	y -intercept	0.920					
EIPNA	50.0	51.9	49.4	47.3	49.5	−0.93	4.64
	100.0	94.1	97.3	101.6	97.7	−2.33	3.84
	250.0	263.0	255.9	227.2	248.7	−0.53	7.63
	500.0	499.9	485.7	517.1	500.9	0.19	3.14
	1000.0	984.9	993.8	975.2	984.6	−1.54	0.95
	5000.0	4973.3	5200.5	5454.9	5209.6	4.19	4.62
	r^2	0.998	0.999	0.996			
	F_{cal}	1671.131					
	F_{ANOVA}	1.290×10^{-17}					
	slope	1.290×10^{-17}					
p value	y -intercept	0.945					

they were resolved from NDMA with a resolution exceeding 2 (see Figure S1). In addition to the solvents, no interference from the matrices of various sartans, e.g., losartan potassium, olmesartan medoxomil, and valsartan sacubitril, was observed, as demonstrated in Figure 2E–G.

2.2.3. Carryover. Carryover was assessed to ensure that no analyte was present from the previously analyzed sample to prevent contamination of the subsequent sample. In this study, DMSO (blank) was injected and evaluated after injecting the highest concentration of the calibration standard. The experiment was performed in triplicate. The peak response in

the blank sample was compared to that in the lowest calibration sample. The percentage of interference must be less than 20%. The results revealed no peak at the retention times of NDMA, NDEA, DIPNA, and EIPNA, demonstrating that the method has no carryover effect, as shown in Figure 3.

2.2.4. Calibration Curve. The linear relationships between analyte concentrations and signal responses were assessed at concentration ranges of 25–5000 ng/mL for NDMA and NDEA and 50–5000 ng/mL for DIPNA and EIPNA. The calibration curves were constructed by plotting different concentrations of NDMA, NDEA, DIPNA, and EIPNA against

their corresponding peak areas. The regression line and homoscedasticity of the calibration curve were assessed to determine whether the ordinary or weighted least square is appropriate for the robust calibration model. The calibration curve was linear over the tested concentration range ($r^2 > 0.995$) using the weighted-linear least square model with a weighting factor of $1/x^2$. The percent deviations of the mean back-calculated concentration and actual spiked plasma concentrations of NDMA, NDEA, DIPNA, and EIPNA were in the ranges of -1.38 to 2.32 , -4.31 to 4.50 , -1.73 to 1.55 , and -2.33 to 4.19 , respectively. The %CV ($n = 3$) of the back-calculated concentration was less than 7.63%. The calibration curve results are summarized in Table 1.

According to the residual plots and regression generated using one-way analysis of variance, the results demonstrated that the F values (F_{ANOVA}) of all regression lines were significantly less than the calculated F value (F_{cal}), indicating a good linear relationship between the peak response from the instrument (y) and the concentration of the analyte (x). The p values of the slope and y -intercept were also calculated, as summarized in Table 1. The p values of the slope and intercept represent whether the slope and intercept significantly differ from zero.²⁸ The results demonstrated that the p values of the slope were less than 0.05, indicating a significant difference versus zero. However, the p values of the intercept exceeded 0.05, indicating that the intercepts of all regression lines were insignificantly different from zero. Consequently, a single-point calibration standard can be applied for the routine analysis of nitrosamine determination.

2.2.5. Linearity and Range. The range of an analytical procedure evaluates its ability to elicit test results that are directly, or by a well-defined mathematical transformation, proportional to the concentration of an analyte in sample matrices with a given dynamic range. Thus, the linear relationships between the spiked concentration of four nitrosamines in the losartan matrix and the back-calculated concentration results were demonstrated via the dynamic range of analysis. As presented in Table 2, the results were linear over the concentration ranges of 25–5000 ppb for NDMA and NDEA and 50–5000 ppb for DIPNA and EIPNA ($r^2 > 0.995$).

Table 2. Range of Nitrosamines

nitrosamine	dynamic range (ppb)	slope	intercept	r^2
NDMA	25–5000	1.038	−15.178	0.9998
NDEA	25–5000	1.038	−15.791	0.9997
DIPNA	50–5000	0.983	8.498	0.9999
EIPNA	50–5000	0.968	17.140	0.9999

2.2.6. Accuracy and Precision. The intraday and interday accuracy and precision were evaluated at LOQ QC, MCQ, and HQC. The accuracy and precision results are summarized in Table 3. The spiked QC samples of NDMA for intraday and interday accuracy exhibited percent deviation ranging from -4.19 to 4.89 (%CV = 1.3–4.8) and from -2.57 to 5.57 (%CV = 1.3–5.9), respectively. The spiked QC samples of NDEA for the intraday and interday accuracy exhibited % deviation ranging from -6.47 to 7.15 (%CV = 0.4–1.3) and from -7.04 to 5.82 (%CV = 1.4–2.4), respectively. The spiked QC samples of DIPNA for intraday and interday accuracy exhibited % deviation ranging from -0.58 to 6.42 (%CV = 0.9–4.6) and from 2.79 to 3.98 (%CV = 2.9–4.8), respectively. The spiked QC samples of EIPNA for intraday and interday accuracy

exhibited % deviation ranging from 4.34 to 7.25 (%CV = 1.1–2.0) and from -4.19 to 5.13 (%CV = 2.5–11.5), respectively. The accuracy and precision results showed good accuracy and precision of the proposed method.

2.2.7. Limit of Detection and LOQ. The experiments demonstrated the sensitivity of the method via limit of detection (LOD) and LOQ. Compared to the GC–MS from the US FDA application note,¹⁷ the developed method is more sensitive in NDMA and NDEA analysis. Recently, the US FDA has published a more specific HPLC–HRMS method²³ with an LOQ of 50 ppb for nitrosamine analysis. However, the developed method demonstrated a more sensitive LOQ at 25 ppb for NDMA and NDEA, while the sensitivity of DIPNA and EIPNA for both methods is comparable. The quantification limits obtained from the developed method are much lower than the interim limit criteria proposed by the US FDA⁴ (0.96 ppm for NDMA and 0.27 ppm for NDEA, DIPNA, and EIPNA) and the EMA⁶ (0.640 ppm for NDMA and 0.177 ppm for NDEA, DIPNA, and EIPNA), indicating that the method can be applied for nitrosamine analysis in losartan APIs. The LOD and LOQ of the method for nitrosamine determination are summarized in Table 4.

2.2.8. Robustness and Stability. The robustness of the developed method was investigated to determine any impacts on the result in terms of the small variations of chromatographic parameters and the stability profile of the test sample. The headspace equilibrium time was varied in the range of ± 1 min from the proposed method condition. The transfer line temperature was also varied in the range of ± 5 °C from the proposed method condition. The robustness results (see Tables S1–S8) demonstrated the %CV of peak area and retention time of ≤ 7.49 and ≤ 0.03 , respectively, with the tailing factor of ≤ 1.27 , indicating that the efficiency of the method was not affected upon small changes in the investigating parameters. Moreover, there is no significant impact on the separation between ¹⁵N-DMF and NDMA with the resolution of > 2 (see Figure S2) under the variations of the flow rate (± 0.1 mL/min).

In addition to the chromatographic parameter variations, the sample stability in the headspace autosampler at room temperature was studied to estimate the appropriate analysis time after sample preparation. The percent deviation of all nitrosamines at 6, 12, and 24 h was $< -6.54\%$, demonstrating that the sample was stable in the autosampler for up to 24 h after preparation.

2.3. Applications of the Method for Nitrosamine Determination in Losartan Potassium APIs. The validated method was applied to investigate four contaminating nitrosamines, namely, NDMA, NDEA, DIPNA, and EIPNA, in 10 batches of losartan potassium samples from two manufacturers. In this study, triplicate analyses were performed for each production batch. The repeatability (%CV) of the results was found to be less than 1%. The results are summarized in Table 5. Based on the current interim limit criteria of nitrosamines, all samples were found to meet the acceptable levels of the four nitrosamines. It is of note that the EU has proposed a limit for all nitrosamines at 30 ppb with an effective date of April 1, 2021. Therefore, there are five out of 10 batches that contain NDMA exceeding the proposed limit. In contrast, two other batches were detected with NDEA at a level higher than the LOQ of 25 ppb but less than 30 ppb.

Table 3. Intraday and Interday Accuracy and Precision

compound	nominal concentration (ppb)	intraday (<i>n</i> = 3)			interday (<i>n</i> = 6)		
		mean back-calculated concentration (ppb)	accuracy (% deviation)	precision (%CV)	mean back-calculated concentration (ppb)	accuracy (% deviation)	precision (%CV)
NDMA	25.0	24.0	−4.19	4.8	24.4	−2.57	5.9
	250.0	262.2	4.89	1.5	262.9	5.01	1.3
	1000.0	1047.8	4.78	1.3	1063.6	5.57	1.8
NDEA	25.0	23.4	−6.47	0.9	23.2	−7.04	1.4
	250.0	267.9	7.15	0.4	264.5	5.82	1.8
	1000.0	1062.6	6.26	1.3	1045.4	4.54	2.4
DIPNA	50.0	53.2	6.42	0.9	52.0	3.98	2.9
	250.0	247.2	−1.12	4.0	257.0	2.79	4.8
	1000.0	994.2	−0.58	4.6	1030.4	3.04	4.8
EIPNA	50.0	52.2	4.34	2.0	47.9	−4.19	11.5
	250.0	268.1	7.25	1.9	255.4	2.17	6.5
	1000.0	1068.7	6.87	1.1	1051.3	5.13	2.5

Table 4. LOD and LOQ of Nitrosamines

compound	LOD (<i>n</i> = 5)			LOQ (<i>n</i> = 5)			
	nominal concentration (ppb)	%CV	S/N	nominal concentration (ppb)	accuracy (% deviation)	precision (%CV)	S/N
NDMA	5	6.8	6.5	25	3.3	2.9	36.8
NDEA	5	3.7	3.8	25	−5.2	1.0	15.4
DIPNA	25	5.6	4.3	50	3.8	0.8	27.3
EIPNA	25	6.5	3.7	50	−2.4	3.5	39.9

Table 5. Results of Four Nitrosamines in 10 Batches of Losartan Potassium (*n* = 3)

batch no.	nitrosamine content (ppb) (%CV)			
	NDMA	NDEA	DIPNA	EIPNA
1	81.0 (0.75)	<5	<25	<25
2	30.5 (0.38)	<5	<25	<25
3	35.1 (0.16)	<5	<25	<25
4	55.4 (0.28)	<5	<25	<25
5	42.8 (0.54)	<5	<25	<25
6	<5	25.4 (0.82)	<25	<25
7	<5	28.6 (0.35)	<25	<25
8	<5	<5	<25	<25
9	<5	<5	<25	<25
10	<5	<5	<25	<25

3. MATERIALS AND METHODS

3.1. Drugs and Chemicals. An NDMA certified reference standard at a concentration of 1004.0 $\mu\text{g/mL}$ (lot no. A0141467) was purchased from Restek (Bellefonte, PA, USA). NDEA (lot no. H5GMI, purity 100.0%) was purchased from Tokyo Chemical Industry Co, Ltd. (Tokyo, Japan). DIPNA (lot no. 9-AQL-128-2, purity 98.0%) and EIPNA (lot no. 2-AMR-151-1, purity 95.0%) were purchased from Toronto Research Chemicals (Toronto, Canada). Losartan potassium, valsartan sacubitril, and olmesartan medoxomil were provided by Siam Bheasach Co., Ltd. (Bangkok, Thailand). A preliminary screening test for selecting nitrosamine-free sartans was performed for losartan potassium, valsartan sacubitril, and olmesartan medoxomil to ensure the lack of interference in the sartan standards used in the validation experiments. Ten batches of losartan potassium APIs were purchased from two manufacturers (IPCA, Mumbai, India; Hetero Labs Limited, Hyderabad, India). DMSO was

purchased from Thermo Fisher Scientific (Leics, UK). All reagents were of at least analytical grade. The purified water was prepared from a Milli-Q water purification system (Millipore; S.A.S, France).

3.2. Instruments and Chromatographic Conditions.

GC was performed on a Nexis GC-2030 system (Shimadzu Corp., Kyoto, Japan) with an HS-20 headspace autosampler. An analytical separation was performed using an SH-Rtx-Wax (polyethylene glycol) GC column (30 $\times$ 0.32 mm² i.d., 1 μm , Shimadzu America, MD, USA) placed in a column oven built-in GC system. Helium was used as a carrier gas. The temperature programming was set in a gradient mode using an initial constant temperature at 40 $^{\circ}\text{C}$ for 2 min, a ramping temperature of 20 $^{\circ}\text{C}/\text{min}$ to 240 $^{\circ}\text{C}$, and a holding temperature at 240 $^{\circ}\text{C}$ for 3.5 min. The carrier gas flow rate was set at 2.0 mL/min. The oven sample was heated up to 150 $^{\circ}\text{C}$ under a vibration level of 5 with an appropriate sample equilibrating time of 15 min. After equilibration, the sample vapor was pressurized for 1 min before 1 μL of the sample was injected under a gas pressure of 103 kPa with a split ratio of 3:1. The temperatures of the sample and transfer lines were set at 230 and 250 $^{\circ}\text{C}$, respectively. The solvent cut time was set at 1 min. The filament was turned off at 9.0 min. The chromatographic run time was 15.5 min, and the cool down period was 1.7 min. Thus, the total analysis time was 17.2 min. The entire flow was directed into the detector.

Quantitation was achieved via MS detection in the EI mode for nitrosamine detection using a QP2020 NX mass spectrometer (Shimadzu Corp., Kyoto, Japan). The detector voltage was set at +0.6 kV. Both the ion source and the interface temperatures were set at 230 $^{\circ}\text{C}$. NDMA (m/z 74.0), NDEA (m/z 102.0), DIPNA (m/z 130.0), and EIPNA (m/z 116.0) were quantified on a single-quadrupole mass spectrometer in the EI mode under the SIM acquisition method. The dwell time or event time was set at 300 ms.

Quantitation of NDMA, NDEA, DIPNA, and EIPNA was based on their peak areas. Data were processed using Lab Solutions Software (GCMS Solution Version 4.50, Shimadzu Corp., Kyoto, Japan).

3.3. Preparation of Nitrosamine Stock and Working Standard Solutions. The NDMA certified reference standard (1000 $\mu\text{g/mL}$) was used as an NDMA stock standard solution. Other nitrosamine stock standard solutions were separately prepared by dissolving NDEA, DIPNA, or EIPNA standards in methanol at a concentration of 1000 $\mu\text{g/mL}$. Nitrosamine working standard solutions of NDMA, NDEA, DIPNA, and EIPNA were prepared at a concentration of 5 $\mu\text{g/mL}$ by diluting each stock standard solution with an appropriate amount of DMSO.

3.4. Preparation of Calibration Standards and Spiked Quality Control Samples. Calibration standards for NDMA, NDEA, DIPNA, and EIPNA were prepared by aliquoting the working standard solutions in DMSO to yield final concentrations of 25 (LOQ of NDMA and NDEA), 50 (LOQ of DIPNA and EIPNA), 100, 250, 500, 1000, and 5000 ng/mL. Spiked QC samples for NDMA and NDEA were prepared by adding the working standard solutions to the nitrosamine-free losartan API at concentrations of 25, 250, and 1000 ppb, representing QC at LOQ (LOQ QC), medium QC (MQC), and high QC (HQC) samples, respectively. For DIPNA and EIPNA, spiked QC samples were prepared at concentrations of 50 (LOQ), 250, and 1000 ppb, representing LOQ QC, MQC, and HQC, respectively.

3.5. Sample Preparation. Five milliliters of DMSO were transferred to a glass vial for headspace injection containing 1 g losartan potassium. The sample was then gently vortexed using a vortex mixer at the maximum rotation speed for 5 min. The clear sample solution was placed into the HS-20 headspace autosampler. The sample was incubated in the headspace oven at 150 $^{\circ}\text{C}$ under the vibration level of 5, with the appropriate sample equilibrating time of 15 min. The residual nitrosamines were then vaporized into the vial headspace and directly injected into the column.

3.6. Method Validation. The method was validated according to the Q2(R1) ICH guidelines for analytical method validation.²⁵ The validation parameters included system suitability, specificity, carryover, linearity, accuracy, precision, LOD, LOQ, and robustness.

3.6.1. System Suitability. The suitability of the system was evaluated before the validation experiments and impurity determination. A system suitability solution containing a mixture of NDMA, NDEA, DIPNA, and EIPNA was prepared at 250 ng/mL for each compound by appropriately aliquoting the working standard solutions (1000 $\mu\text{g/mL}$). The system suitability solution was injected with five replicates. The system performance was evaluated on the tailing factor of analytes ($T \leq 2$), and the system reproducibility was assessed as the variation of the peak area ($\%CV \leq 11\%$) and retention time ($\%CV \leq 2\%$) to indicate the precision of injections.

3.6.2. Specificity and Selectivity. Specificity is the ability to discriminate the analyte in the presence of interferences such as solvents, matrices, and impurities.²⁶ The specificity was investigated by separately injecting diluents, e.g., DMSO and methanol, an authentic standard of nitrosamine solution (250 ng/5 mL in DMSO), and blank sartan matrices including losartan potassium, olmesartan medoxomil, and valsartan sacubitril (1 g/5 mL in DMSO). According to Yang et al.,²⁴ a ^{15}N isotope of dimethylformamide (DMF) with a molecular

mass per charge (m/z) of 74 may interfere with the analysis of NDMA. The concern of ^{15}N -DMF contamination becomes significant if the chromatographic condition cannot separate DMF from NDMA. Therefore, DMF at a concentration of 500 ng/5 mL in DMSO also caused interference. The specificity was determined as the presence or absence of interfering peaks at the same m/z value and retention time of NDMA, NDEA, DIPNA, and EIPNA on the chromatograms.

3.6.3. Carryover. The carryover effect was evaluated as the percentage of carryover by analyzing a blank DMSO after injecting the upper LOQ concentration of 5000 ng/mL of the calibration standard curve. The percentage of carryover in the blank sample was calculated compared to the LOQ and considered acceptable at $\leq 20\%$. Three independent experiments were performed on three different days.

3.6.4. Calibration Curves. Seven-point calibration curves were constructed in triplicate by plotting the analyte peak areas versus different analyte concentrations of NDMA, NDEA, DIPNA, or EIPNA. The weighted-linear least square model with a $1/x^2$ weighting factor was selected using a statistical test of homoscedasticity.²⁷ The coefficient of determination ($r^2 \geq 0.995$) was determined to verify the linearity of the plot. The percent deviation of the mean back-calculated concentrations of NDMA, NDEA, DIPNA, and EIPNA should be within $\pm 10\%$. The residual plot was demonstrated whether the slope and y-intercept were significantly different from zero at a 95% confidence interval. Statistically, a p value of the slope should be less than 0.05, meaning that it is significantly different from zero, whereas the p value of the y-intercept should be more than 0.05, meaning that it is insignificant from zero. F_{cal} should be greater than F_{ANOVA} demonstrating the linear regression.²⁸

3.6.5. Linearity and Range. The range describes the concentration interval between the lower and upper quantitative levels. The range of the method was established using the spiked calibration samples by constructing a calibration curve using a losartan matrix spiked with the corresponding nitrosamines. The equation model was designed in the same manner as the calibration curve experiment (Section 3.6.4). The slope, intercept, and r^2 were calculated.

3.6.6. Accuracy and Precision. Accuracy and precision were determined by analyzing the spiked losartan samples at 25 (LOQ), 250, and 1000 ppb for NDMA and NDEA and 50 (LOQ), 250, and 1000 ppb for DIPNA and EIPNA. Each concentration level was prepared and analyzed in triplicate on the same day and for two different days for intraday (within-run) and interday (between-run) analyses, respectively. Accuracy was assessed using the percent deviation between the mean back-calculated concentration and nominal (actual) concentration. The percent deviation should be within $\pm 10\%$. The intraday and interday precision values were determined using the $\%CV$ of the back-calculated concentration. The $\%CV$ should be $\leq 11\%$ for all concentrations excluding the LOQ, which should be $\leq 15\%$.

3.6.7. LOD and LOQ. The LOD is the lowest amount of analyte in a sample matrix that could be detected but not necessarily quantitated as an exact value. The LOD can be derived from the signal-to-noise (S/N) ratio, and it is usually expressed as the concentration of an analyte in the sample matrix. It is determined by injecting samples that generate the S/N ratio and $\%CV$ of replicate injections. The LOD was verified in terms of $S/N \geq 3$ and $\%CV$ of five replicate injections of $\leq 15\%$.

The LOQ is the lowest amount of analyte in a sample matrix that can be quantitatively determined with appropriate precision and accuracy. Typically, the LOQ is estimated by determining the S/N ratio and %CV of replicate injections. The LOQ established using spiked samples independent of calibration standards with five replicate injections was satisfied with accuracy and precision of percent deviation within ± 10 and %CV ≤ 15 , respectively.

3.6.8. Robustness and Stability. Robustness is an indicator of method tolerability when small variations occur in method parameters to ensure the reliability of the method during routine analysis. In this study, the robustness of the method was determined as the suitability of the system to permit changes in transfer temperature and headspace equilibration time. The precision values of the peak response and retention time for four nitrosamines were determined after slight variation in the method parameters. The %CV of the peak area and retention time should be ≤ 11 and 2, respectively, whereas the tailing factor of each injection should be lower than 2. Moreover, the influence of the flow rate on the resolution between ^{15}N -DMF and NDMA was studied. The resolution should be > 2 . In addition, the stability of the sample solutions was investigated. To confirm the good stability of the sample solution, the percent deviation of the analyte concentration between the initial time point (0 h) and the storage time of 6, 12, or 24 h should not be greater than 10%.

3.7. Application for Nitrosamine Determination in Losartan Potassium APIs. The developed and validated method was applied to determine residual contamination by NDMA, NDEA, DIPNA, and EIPNA in ten batches of losartan potassium API samples. Five milliliters of DMSO was transferred into a glass vial for headspace injection containing 1 g of losartan potassium, and the mixture was vortexed until it was completely dissolved. The sample solution was placed into the built-in headspace autosampler of the oven. The sample was incubated in the headspace oven at 150 °C under a vibration level of 5, with an appropriate sample equilibrating time of 15 min. The residual nitrosamines were then vaporized into the vial headspace and directly injected into the column.

4. CONCLUSIONS

In this work, the HS-GC–MS method was developed and validated for the quantitation of NDMA, NDEA, DIPNA, and EIPNA in losartan potassium APIs. The headspace injection technique was applied to serve as an online sample preparation in residual impurity analysis, thereby offering benefits concerning the efficient removal of potential interference and high-throughput routine analysis. The complete method validation was performed using the Q2(R1) ICH guidelines.²⁶ The overall results indicated that the developed method is reliable and useful for determining the levels of four nitrosamines in losartan APIs. For other sartans, we confirmed the selectivity of the method based on the absence of interfering matrix co-elution at the retention times of individual nitrosamines. To use the method to analyze nitrosamine contamination in other sartan APIs and drug products, the method should be further fully investigated for its accuracy, precision, and other validation parameters using the acceptable limits of nitrosamines in each substance. The LOQ and LOD of the developed method were far below the US FDA and EMA interim limits of the corresponding nitrosamines in losartan potassium. In addition, the established LOD and LOQ can support a tighter limit of 30 ppb for NDMA and

NDEA, which will be implemented in the forthcoming year.⁷ Although the developed method cannot be applied in the quantitative sense under this stricter limit, it can also serve as a screening test for DIPNA and EIPNA. In a broader sense, the methodology can also be adapted for the simultaneous analysis of additional nitrosamines in other sartans.

■ ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acsomega.1c00982>.

Chromatograms of spiked NDMA (m/z 74), spiked DMF (^{15}N isotopic ion at m/z 74), and co-spiked between NDMA and DMF in losartan (Figure S1); results of method robustness under small variations in the HS equilibration time and transfer line temperature (Tables S1–S8); and chromatograms of co-spiked between NDMA and DMF in losartan under the variations of the flow rate (Figure S2) (PDF)

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Author Contributions

W.W. conceived and designed the project methodology. O.S. managed all resources used in all experiments and performed graphic preparation. P.S. and K.C. conducted the method validation, data acquisition, and evaluation. S.W. wrote the original draft of manuscript. P.R. reviewed the manuscript. All authors have given approval to the final version of the manuscript.

Notes

The authors declare no competing financial interest.

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REFERENCES

- (1) Barreras, A.; Gurk-Turner, C. Angiotensin II receptor blockers. *Baylor Univ. Med. Cent. Proc.* **2003**, *16*, 123–126.
- (2) Burnier, M.; Brunner, H. R. Angiotensin II receptor antagonists. *Lancet* **2000**, *355*, 637–645.
- (3) Park, J.; Seo, J.; Lee, J.; Kwon, H. Distribution of seven *N*-nitrosamines in food. *Toxicol. Res.* **2015**, *31*, 279–288.
- (4) US FDA press announcements. FDA announces voluntary recall of several medicines containing valsartan following detection of an impurity. <http://FDA.gov> (accessed Jul 13, 2018).
- (5) EMA release. CHMP list of questions. EMA/CHMP/467845/2018. https://www.ema.europa.eu/en/documents/referral/angiotensin-ii-receptor-antagonists-sartans-article-31-referral-chmp-list-questions-be-addressed-api_en.pdf (accessed Jul 16, 2018).
- (6) EMA release. Temporary interim limits for NMBA, DIPNA and EIPNA impurities in sartan blood pressure medicines. EMA/351053/2019 rev 1. https://www.ema.europa.eu/en/documents/other/temporary-interim-limits-nmba-dipna-eipna-impurities-sartan-blood-pressure-medicines_en.pdf (accessed Aug 20, 2019).
- (7) EMA release. Overview and recommendation: Lessons learnt from presence of *N*-nitrosamine impurities in sartan medicines. EMA/CHMP/526934/2019. https://www.ema.europa.eu/en/documents/report/lessons-learnt-presence-n-nitrosamine-impurities-sartan-medicines_en.pdf (accessed Jun 23, 2019).
- (8) Oruganti, S. A viewpoint on the presence of *N*-nitrosodimethylamine (NDMA) impurity in certain generic samples of Valsartan, Drils Perspectives; Dr. Reddy's Institute of Life Sciences, 2018.
- (9) EMA release. Assessment report. EMA/CHMP/217823/2019. https://www.ema.europa.eu/en/documents/variation-report/angiotensin-ii-receptor-antagonists-sartans-article-31-referral-chmp-assessment-report_en.pdf (accessed Feb 14, 2019).
- (10) European network of OMCLs, OMCLs Release. *Determination of NDMA in valsartan active substances and finished products by HPLC/UV*; French National Agency for Medicines and Health Products Safety, 2018.
- (11) European network of OMCLs, OMCLs Release. *Test method for the determination of NDMA by LC/MS/MS in Valsartan finished products*; Chemisches und Veterinäruntersuchungsamt: Karlsruhe, Germany, 2018.
- (12) European network of OMCLs, OMCLs Release. *Determination of NDMA (HS-GC-MS) method 3/30, issue no. 1*; Public Analyst's Laboratory: Galway, Ireland, 2018.
- (13) USFDA. *GC/MS Headspace Method for NDMA in Valsartan Drug Substance and Drug Products*; US Food and Drug Administration: Washington, DC, 2019.
- (14) USFDA. *Combined N-Nitrosodimethylamine (NDMA) and N-Nitrosodiethylamine (NDEA) Impurity Assay by GC/MS-Headspace*; US Food and Drug Administration: Washington, DC, 2019.
- (15) Taiwan FDA. *Method of Test for Nitrosamines in Medicines Multiple Analysis (GC-MS/MS Method)*; Taiwan Food and Drug Administration: Nangang, Taipei City, 2020.
- (16) Chang, S.; Chang, C.; Wang, L.; Chen, W.; Fan, S.; Zang, C.; Hsu, Y.; Lin, M.; Tseng, S.; Wang, D. A multi-analyte LC-MS/MS method for screening and quantification of nitrosamines in sartans. *J. Food Drug Anal.* **2020**, *28*, 98–107.
- (17) USFDA. *Combined Direct Injection N-Nitrosodimethylamine (NDMA), N-Nitrosodiethylamine (NDEA), N-Nitrosoethylisopropylamine (NEIPA), N-Nitrosodiisopropylamine (NDIPA), and N-Nitrosodibutylamine (NDBA) Impurity Assay by GC-MS/MS*; US Food and Drug Administration: Washington, DC, 2019.
- (18) USFDA. *Combined headspace N-Nitrosodimethylamine (NDMA), N-Nitrosodiethylamine (NDEA), N-Nitrosoethylisopropylamine (NEIPA), N-Nitrosodiisopropylamine (NDIPA), and N-Nitrosodibutylamine (NDBA) Impurity Assay by GC-MS/MS*; US Food and Drug Administration: Washington, DC, 2019.
- (19) Brian Byrd, J.; Chertow, G. M.; Bhalla, V. Hypertension hot potato—Anatomy of the angiotensin-receptor blocker recalls. *N. Engl. J. Med.* **2019**, *380*, 1589–1591.
- (20) FDA Updates and Press Announcements on Angiotensin II Receptor Blocker (ARB) Recalls (Valsartan, Losartan, and Irbesartan): FDA updates table of interim limits for nitrosamine impurities in ARBS. <https://www.fda.gov/drugs/drug-safety-and-availability/fda-updates-and-press-announcements-angiotensin-ii-receptor-blocker-arb-recalls-valsartan-losartan> (accessed Feb 28, 2019).
- (21) EMA release. Questions and answers for marketing authorization holder/applicants on the CHMP Opinion for the Article 5 (3) of Regulation (EC) No 726/2004 referral on nitrosamine impurities in human medicinal products. EMA/CHMP/409815/2020. https://www.ema.europa.eu/en/documents/referral/nitrosamines-emea-ha53-1490-questions-answers-marketing-authorisation-holders/applicants-chmp-opinion-article-53-regulation-ec-no-726/2004-referral-nitrosamine-impurities-human-medicinal-products_en.pdf (accessed Aug 03, 2020).
- (22) EMA release. Assessment report: Procedure under Article 5(3) of Regulation EC (No) 726/2004 Nitrosamine impurities in human medicinal products. EMA/CHMP/369136/2020. https://www.ema.europa.eu/en/documents/referral/nitrosamines-emea-ha53-1490-assessment-report_en.pdf (accessed June 25, 2020).
- (23) USFDA. *Liquid Chromatography-High Resolution Mass Spectrometry (LC-HRMS) Method for the Determination of Six Nitrosamine Impurities in ARB Drugs*; US Food and Drug Administration: Washington, DC, 2019. <https://www.fda.gov/media/125478/download>
- (24) Yang, J.; Andres Marzan, T.; Ye, W.; Sommers, C. D.; Rodriguez, J. D.; Keire, D. A. A cautionary tale: Quantitative LC-HRMS analytical procedures for the analysis of *N*-Nitrosodimethylamine in metformin. *AAPS J.* **2020**, *20*, No. 89.
- (25) Tsutsumi, T.; Akiyama, H.; Demizu, Y.; Uchiyama, N.; Masada, S.; Tsuji, G.; Arai, R.; Abe, Y.; Hakamatsuka, T.; Izutsu, K.; Goda, Y.; Okuda, H. Analysis of an impurity, *N*-Nitrosodimethylamine, in valsartan drug substances and associated products using GC-MS. *Biol. Pharm. Bull.* **2019**, *42*, 547–551.
- (26) ICH Harmonised tripartite guideline: Validation of Analytical Procedures: Text and methodology Q2(R1). Current Step 4 version (accessed Oct 27, 1994).
- (27) Danzer, K.; Currie, L. A. Guidelines for calibration in analytical chemistry. *Pure Appl. Chem.* **1998**, *70*, 993–1014.
- (28) Barwick, V. A. *Preparation of Calibration Curves: A Guide to Best Practice*. LGC/VAM/2003/032, 2003, 1–27.
- (29) Crews, C. The determination of *N*-nitrosamines in food. *Qual. Assur. Saf. Crops Foods* **2010**, *2*, 2–12.
- (30) Seyler, T. H.; Kim, J. G.; Hodgson, J. A.; Cowan, E. A.; Blount, B. C.; Wang, L. Quantitation of urinary volatile nitrosamines from exposure to tobacco smoke. *J. Anal. Toxicol.* **2013**, *37*, 195–202.
- (31) Munch, J. W.; Bassett, M. V. *Method S21. Determination of Nitrosamines in Drinking Water by Solid-phase Extraction and Capillary Column Gas Chromatography with Large Volume Injection and Chemical Ionization Tandem Mass Spectrometry (MS/MS)*; U.S. Environmental Protection Agency: Cincinnati, Ohio, 2004.
- (32) Parr, M. K.; Joseph, J. F. NDMA impurity in valsartan and other pharmaceutical products: Analytical methods for the determination of *N*-nitrosamines. *J. Pharm. Biomed. Anal.* **2019**, *164*, 536–549.
- (33) Șerban, E. S.; Socaci, S. A.; Tofană, M.; Maier, S. C.; Bojiță, M. Advantages of “headspace” technique for GC/MS analysis of essential oils. *Farmacia* **2012**, *60*, 249–256.
- (34) Ibrahim, M. M. Investigation on thermal stability and purity determination of two antihypertensive drugs, valsartan and losartan potassium. *Int. J. Curr. Pharm. Res.* **2015**, *7*, 64–69.

Supporting information

Development of a Sensitive Headspace Gas Chromatography–Mass Spectrometry Method for the Simultaneous Determination of Nitrosamines in Losartan Active Pharmaceutical Ingredients

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1. Method development

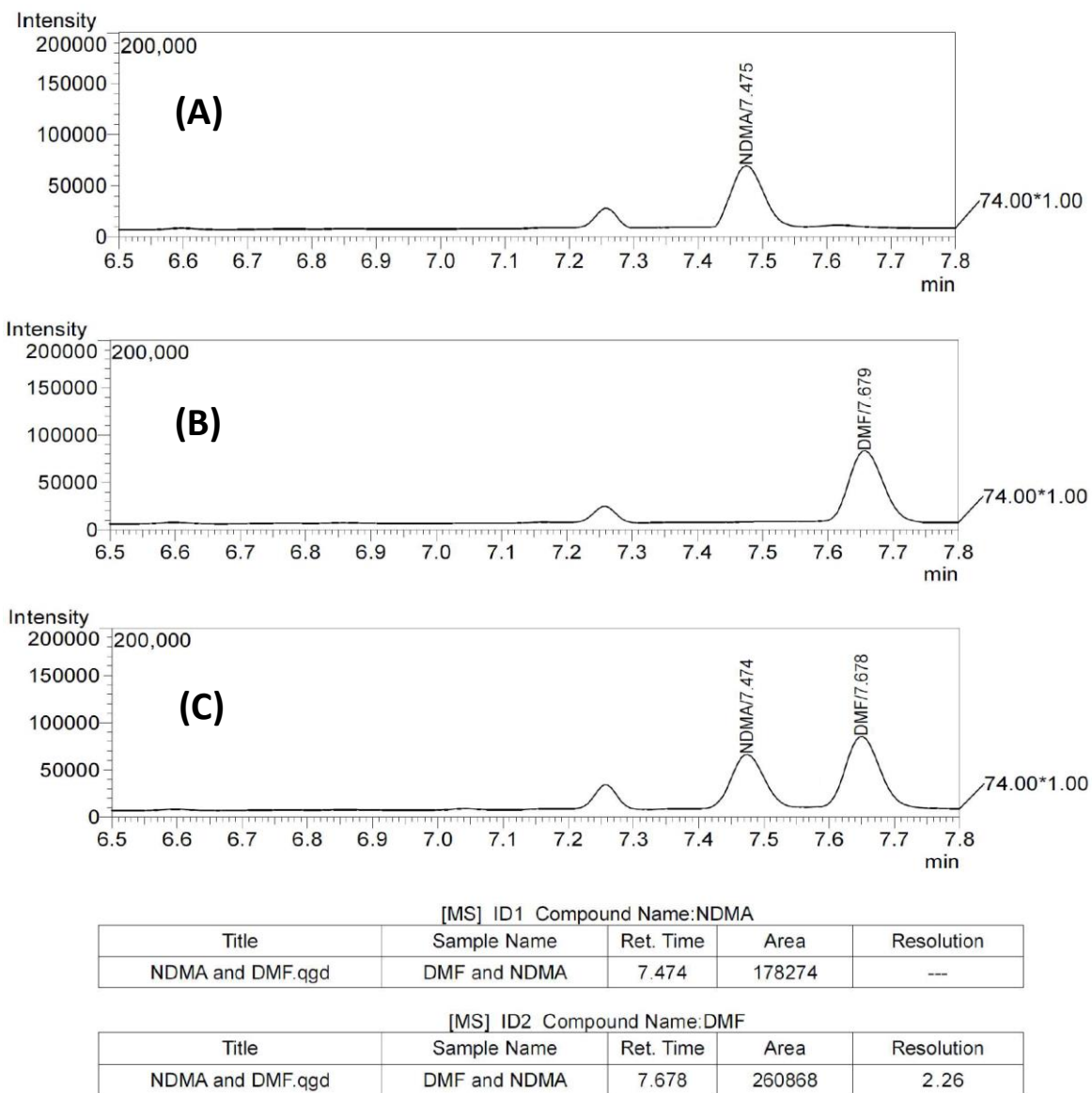


Figure S1. The representative chromatograms of (a) spiked NDMA (m/z 74), (b) spiked DMF (^{15}N isotopic ion at m/z 74) and (c) co-spiked between NDMA and DMF in losartan

2. Robustness

Table S1. System suitability results under HS equilibration time (intra-variation of typical method: 15 min)

Injection no.	NDMA			NDEA			EIPNA			DIPNA		
	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing
#1	140636	7.537	1.106	70612	8.171	1.135	35203	8.446	1.154	24355	8.646	1.317
#2	128975	7.537	1.106	66265	8.170	1.151	32720	8.447	1.143	22677	8.647	1.273
#3	129689	7.535	1.105	65754	8.169	1.135	32043	8.445	1.145	21965	8.646	1.241
#4	132265	7.536	1.103	68565	8.170	1.145	33557	8.446	1.152	22817	8.646	1.267
#5	131122	7.535	1.112	67767	8.168	1.159	33476	8.445	1.127	22380	8.645	1.268
Average	132537	7.536	1.106	67793	8.170	1.145	33400	8.446	1.144	22839	8.646	1.273
SD	4702.897	0.001	0.003	1938.796	0.001	0.010	1181.639	0.001	0.011	908.232	0.001	0.027
%CV	3.55	0.01	0.30	2.86	0.01	0.91	3.54	0.01	0.93	3.98	0.01	2.16

Table S2. System suitability results under HS equilibration time (intra-variation of 14 min)

Injection no.	NDMA			NDEA			EIPNA			DIPNA		
	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing
#1	119970	7.534	1.097	60663	8.168	1.147	29054	8.442	1.164	20142	8.645	1.223
#2	134818	7.539	1.096	68957	8.171	1.160	34396	8.448	1.131	23445	8.649	1.216
#3	127253	7.537	1.094	64574	8.170	1.135	31738	8.445	1.158	21807	8.647	1.213
#4	132658	7.537	1.095	66586	8.171	1.126	33169	8.447	1.149	22813	8.649	1.183
#5	134538	7.540	1.105	69318	8.173	1.144	35076	8.448	1.166	23518	8.650	1.215
Average	129847	7.537	1.097	66020	8.171	1.142	32687	8.446	1.154	22345	8.648	1.210
SD	6302.817	0.002	0.004	3557.088	0.002	0.013	2395.810	0.003	0.014	1409.575	0.002	0.016
%CV	4.85	0.03	0.40	5.39	0.02	1.12	7.33	0.03	1.24	6.31	0.02	1.29

Table S3. System suitability results under HS equilibration time (intra-variation of 16 min)

Injection no.	NDMA			NDEA			EIPNA			DIPNA		
	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing
#1	136310	7.538	1.100	68092	8.172	1.141	34293	8.448	1.124	23424	8.647	1.263
#2	134393	7.541	1.101	69253	8.174	1.140	34025	8.450	1.130	23567	8.651	1.271
#3	133069	7.539	1.099	67883	8.172	1.136	33698	8.449	1.135	23391	8.650	1.215
#4	136804	7.538	1.101	69572	8.171	1.149	33899	8.445	1.174	23293	8.646	1.268
#5	142334	7.539	1.101	72469	8.172	1.133	36004	8.449	1.131	24782	8.649	1.244
Average	136582	7.539	1.100	69454	8.172	1.140	34384	8.448	1.139	23691	8.649	1.252
SD	3547.593	0.001	0.001	1834.914	0.001	0.006	931.045	0.002	0.020	617.525	0.002	0.023
%CV	2.60	0.02	0.08	2.64	0.01	0.53	2.71	0.02	1.76	2.61	0.02	1.86

Table S4. System suitability results under transfer line temperature (intra-variation of Typical method: 250°C)

Injection no.	NDMA			NDEA			EIPNA			DIPNA		
	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing
#1	130505	7.532	1.126	68342	8.166	1.151	34019	8.441	1.174	22763	8.642	1.268
#2	128563	7.532	1.115	66211	8.165	1.148	33529	8.440	1.185	22523	8.643	1.243
#3	121988	7.532	1.122	63913	8.166	1.154	32445	8.441	1.192	21427	8.643	1.241
#4	124316	7.533	1.132	65558	8.167	1.157	33548	8.442	1.216	21427	8.644	1.275
#5	129650	7.534	1.114	67920	8.168	1.156	34665	8.443	1.196	22582	8.645	1.277
Average	127004	7.533	1.122	66389	8.166	1.153	33641	8.441	1.193	22144	8.643	1.261
SD	3676.765	0.001	0.008	1803.538	0.001	0.004	812.876	0.001	0.016	660.837	0.001	0.017
%CV	2.89	0.01	0.67	2.72	0.01	0.32	2.42	0.01	1.30	2.98	0.01	1.39

Table S5. System suitability results under transfer line temperature (intra-variation of 245°C)

Injection no.	NDMA			NDEA			EIPNA			DIPNA		
	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing
#1	127042	7.535	1.129	65466	8.168	1.149	33123	8.443	1.165	21415	8.645	1.245
#2	125627	7.535	1.125	64358	8.169	1.156	32807	8.444	1.188	21507	8.646	1.259
#3	124766	7.535	1.121	64329	8.168	1.141	32702	8.443	1.158	21132	8.644	1.265
#4	127934	7.536	1.129	66488	8.169	1.172	33336	8.445	1.187	22203	8.647	1.258
#5	125748	7.536	1.122	65250	8.169	1.166	32588	8.444	1.210	21952	8.646	1.275
Average	126223	7.535	1.125	65178	8.169	1.157	32911	8.444	1.182	21642	8.646	1.260
SD	1255.186	0.001	0.004	894.068	0.001	0.013	309.998	0.001	0.021	430.312	0.001	0.011
%CV	0.99	0.01	0.33	1.37	0.01	1.08	0.94	0.01	1.75	1.99	0.01	0.87

Table S6. System suitability results under transfer line temperature (intra-variation of 255°C)

Injection no.	NDMA			NDEA			EIPNA			DIPNA		
	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing
#1	114794	7.537	1.114	59203	8.170	1.142	29732	8.444	1.189	19559	8.646	1.236
#2	113182	7.537	1.113	58708	8.170	1.136	29054	8.444	1.185	19116	8.645	1.235
#3	116510	7.536	1.116	59298	8.169	1.145	30375	8.444	1.179	20116	8.646	1.249
#4	132082	7.538	1.127	69710	8.171	1.188	34451	8.444	1.231	22914	8.646	1.336
#5	116531	7.536	1.128	61283	8.170	1.156	30941	8.443	1.229	20165	8.646	1.240
Average	118620	7.537	1.120	61640	8.170	1.153	30911	8.444	1.203	20374	8.646	1.259
SD	7652.640	0.001	0.007	4617.166	0.001	0.021	2101.088	0.000	0.025	1483.967	0.000	0.043
%CV	6.45	0.01	0.65	7.49	0.01	1.79	6.80	0.01	2.10	7.28	0.01	3.44

Table S7 System suitability results under HS equilibration time (inter-variation of 14-16 min)

Variation of HS equilibration time	NDMA			NDEA			EIPNA			DIPNA		
	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing
15 min	132537	7.536	1.106	67793	8.170	1.145	33400	8.446	1.144	22839	8.646	1.273
14 min	129847	7.537	1.097	66020	8.171	1.142	32687	8.446	1.154	22345	8.648	1.210
16 min	136582	7.539	1.100	69454	8.172	1.140	34384	8.448	1.139	23691	8.649	1.252
Average	132989	7.537	1.101	67756	8.171	1.142	33490	8.447	1.146	22958	8.648	1.245
SD	3390.141	0.002	0.005	1717.304	0.001	0.003	852.099	0.001	0.008	680.889	0.002	0.032
%CV	2.55	0.02	0.42	2.53	0.01	0.22	2.54	0.01	0.67	2.97	0.02	2.58

Table S8. System suitability results under transfer line temperature (inter-variation of 245-255°C)

Variation of Transfer line temperature	NDMA			NDEA			EIPNA			DIPNA		
	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing	Peak area	RT	Tailing
250 °C	127004	7.533	1.122	66389	8.166	1.153	33641	8.441	1.193	22144	8.643	1.261
245 °C	126223	7.535	1.125	65178	8.169	1.157	32911	8.444	1.182	21642	8.646	1.260
255 °C	118620	7.537	1.120	61640	8.170	1.153	30911	8.444	1.203	20374	8.646	1.259
Average	123949	7.535	1.122	64402	8.168	1.154	32488	8.443	1.193	21387	8.645	1.260
SD	4631.541	0.002	0.003	2467.690	0.002	0.002	1413.377	0.002	0.011	912.207	0.002	0.001
%CV	3.74	0.03	0.22	3.83	0.03	0.20	4.35	0.02	0.88	4.27	0.02	0.08

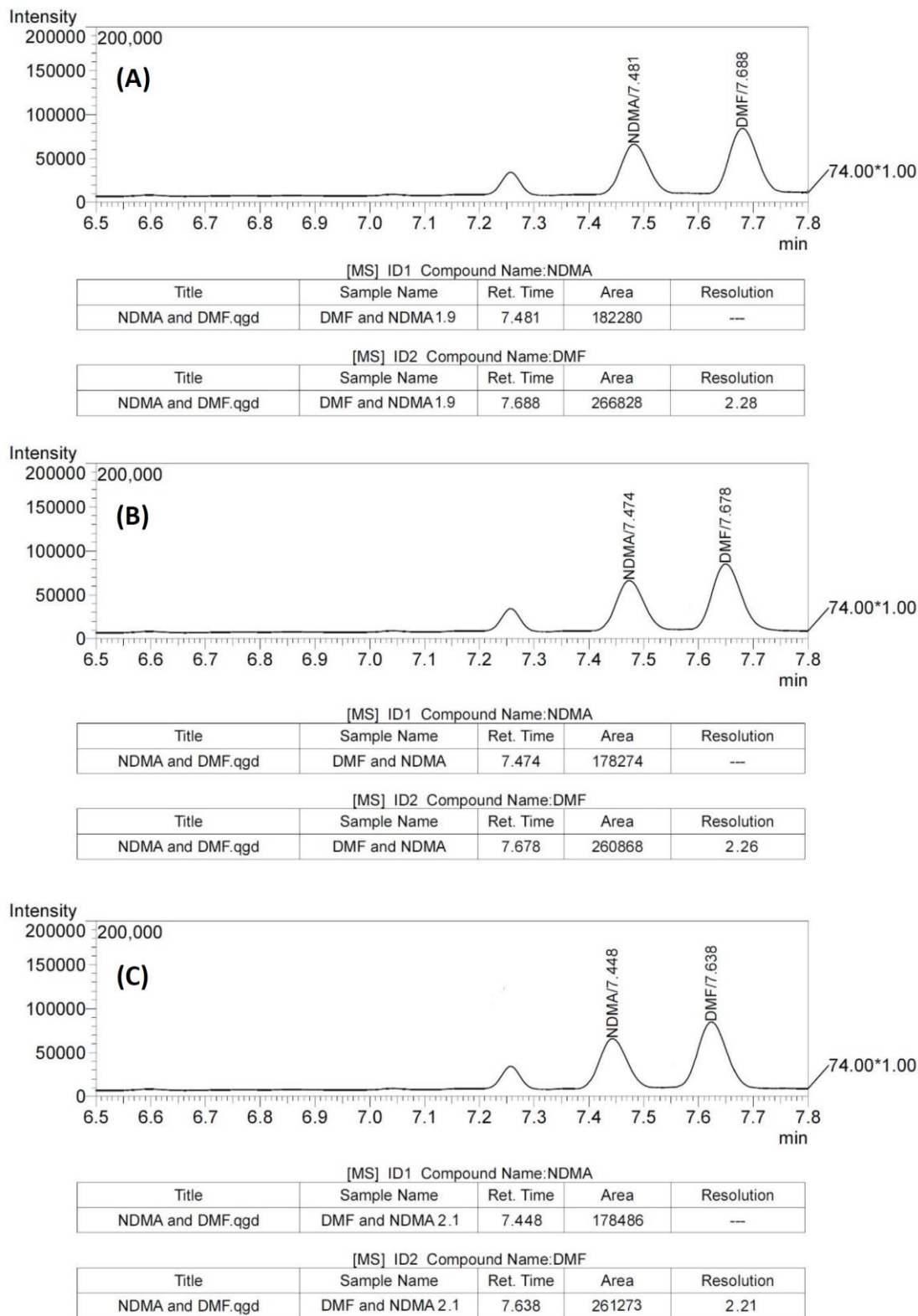


Figure S2. The representative chromatograms of co-spiked between NDMA and DMF in losartan under the variations of flow rate at (a) 1.9 mL/min (b) 2.0 mL/min (method condition) (c) 2.1 mL/min.