

Review Article

Development and Validation of a GC-MS Method for the Simultaneous Quantification of Two Piperazine Designer Drugs Sold in Combination as 'Legal X'

Patrícia Moreira, MSc^{1*}, Diana Dias da Silva, PhD¹, Sara Cunha, PhD², Maria de Lourdes Bastos, PhD¹ and Helena Carmo, PhD^{1*}



¹UCIBIO-REQUIMTE, Laboratory of Toxicology, Department of Biological Sciences, Faculty of Pharmacy, University of Porto, Rua Jorge Viterbo Ferreira, Portugal

²LAQV-REQUIMTE, Laboratory of Bromatology/Hydrology, Department of Chemistry Sciences, Faculty of Pharmacy, University of Porto, Rua Jorge Viterbo Ferreira, Portugal

Abstract

1-Benzylpiperazine (BZP) and 1-(3-trifluoromethylphenyl) piperazine (TFMPP) are piperazine drugs of abuse often taken in combination. Considering the frequency of the co-administration of BZP and TFMPP in the clinical, analytical and forensic contexts and its toxicological impact, we aimed to implement a gas chromatography-mass spectrometry (GC-MS) technique for the detection and simultaneous quantification of BZP and TFMPP in plasma, urine and cell culture medium. The sample preparation involved a protein precipitation (only for plasma), an enzymatic hydrolysis, a liquidliquid extraction (only for cell culture medium), a solid phase extraction, and a perfluoroacylation derivatization. The method was optimized and fully validated, demonstrating linearity over a concentration range of 0-10 μ g/mL for BZP and TFMPP in the three studied matrices. Limits of detection and quantification were 0.004 and 0.016 μ g/mL in plasma, 0.002 and 0.008 μ g/mL in urine, and 0.156-0.312 and 0.312-0.625 μ g/mL in cell culture medium. The method showed high sensitivity, accuracy and precision for the simultaneous detection and quantification of BZP and TFMPP in the three studied matrices and has direct application for the control of drug abuse in humans, post-mortem analysis, and research purposes.

Keywords

Piperazines, Plasma, Urine, Cell culture medium, GC-MS

Introduction

Recently, the synthesis and abuse of new psychoactive substances (NPS), including the so-called designer drugs, have greatly increased [1]. In this context, 1-benzylpiperazine (BZP) and 1-(3-trifluoromethylphenyl)piperazine (TFMPP) are the piperazines most consumed all over the world. These drugs are easily acquired through the Internet [2] and have been abused in association with the purpose of increasing their subjective effects and to promote psychostimulation similar to that elicited by 3,4-methylenedioxy-methamphetamine (MDMA, ecstasy) [3,4]. Also, all too often BZP and TFMPP are misleadingly sold as ecstasy [5-8]. The commercialized forms usually contain larger amounts of BZP than TFMPP, and the most common combination ratio found in seized tablets is 2 BZP:1 TFMPP [9-12].

Although these drugs have been massively consumed by the young attenders of electronic music nightclubs and rave parties in Australia, China and UK [13-17], they have also been a abuse drug of choice in other European countries, with the EMCDDA drug report of 2017 stating that piperazines corre-

*Corresponding author: Patrícia Moreira, MSc, UCIBIO-REQUI-MTE, Laboratory of Toxicology, Department of Biological Sciences, Faculty of Pharmacy, University of Porto, Rua Jorge Viterbo Ferreira, 228, 4050-313, Porto, Portugal; and

Helena Carmo, PhD, UCIBIO-REQUIMTE, Laboratory of Toxicology, Department of Biological Sciences, Faculty of Pharmacy, University of Porto, Rua Jorge Viterbo Ferreira, 228, 4050-313, Porto, Portugal

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sponded to 6% of the NPS reported to the EU Early Warning System [18]. These data indicate that this occurrence cannot be overlooked, particularly because there have been reports of non-fatal and fatal intoxication attributed to BZP and TFMPP [3,8,19,20].

Regarding the pharmacological, toxicological or pharmacokinetic properties of BZP and TFMPP, there is limited information available. Of note, their drug interactions are yet to be completely clarified [21,22]. In this sense, a method for the simultaneous quantification of BZP and TFMPP would be extremely useful to clarify these interactions and to fulfill the lack of data.

Since no analytical methods were previously validated for the concomitant analysis and/or quantification of both drugs, and by virtue of the high frequency of the co-administration of both drugs and the expected toxicological impact, it was deemed of greatest significance to develop and validate a GC-MS methodology that enabled the simultaneous quantification of BZP and TFMPP and that was appropriate for clinical, forensic and research purposes.

Experimental

Materials

All chemicals and reagents were of analytical grade. BZP (99.3% purity) was acquired from Chemos GmbH (Regenstauf, Germany) and TFMPP (98% purity) from Alfa Aesar (Karlsruhe, Germany). 4-Hydroxy-3-methoxybenzylamine hydrochloride (98% purity), phenylpropanolamine, triethylamine (TEA), tert-butyl methyl ether, trifluoroacetic anhydride (TFAA), β-glucuronidase (type 2) from Helix pomatia, bovine insulin, dexamethasone, gentamicin, fungizone and Williams' Medium E with L-glutamine were purchased from Sigma-Aldrich (St. Louis, Missouri). Antibiotic solution (consisting of 10,000 U/mL penicillin and 10,000 µg/mL streptomycin) was purchased from Invitrogen Corporations (Paisley, UK). Methanol (MeOH; HPLC-grade) was supplied by Chem-Lab (Zedelgem, Belgium). Ethyl acetate and hexane (HPLCgrade) were supplied by Fisher Scientific (Loughborough, UK). Hydrochloric acid 37% (HCl), ammonia solution 25% (NH₂OH), sodium acetate trihydrated (C2H3NaO2.3H2O), trichloroacetic acid (TCA) and formic acid were purchased from Merck (Darmstadt, Germany). Sodium hydroxide (NaOH) was obtained from VWR (Leuven, Belgium).

Blank samples

Drug free urine and plasma (blanks) were collected from adult volunteers, healthy and without history of drug use. To obtain plasma, blood was collected into heparin tubes and centrifuged at 1,600 g for 10 min at 4 °C. Blank urine and plasma samples were frozen at -80 °C. The cell culture medium used was Williams' Medium E (no L-glutamine; no sodium bicarbonate) supplemented with 1% antibiotic solution, 5 µg/ mL bovine insulin, 50 µM dexamethasone, 100 µg/mL gentamicin and 2.5 µg/mL fungizone, and stored at 4 °C.

Calibrator samples

Stock solutions of standards and internal standards (IS)

were prepared at 1 mg/mL in MeOH. This solvent was also used for further intermediate solutions. A set of calibrators of the two drugs were prepared for plasma (0.016, 0.08, 0.4, 2 and 10 µg/mL), urine (0.016, 0.08, 0.4, 2 and 10 µg/mL) and cell culture medium (0.625, 1.25, 2.5, 5 and 10 µg/mL) by fortifying 500 µL of blank plasma, urine or cell culture medium. Each urine and plasma specimen was added of 10 µL of 100 µg/mL 4-hydroxi-3-methoxybenzylamine hydrochloride (IS), while cell culture medium specimens were added of 10 µL of 50 µg/mL phenylpropanolamine (IS) before sample preparation.

Sample preparation

A sample preparation protocol was newly developed using as reference the methods previously described for amphetamines [23,24].

Protein precipitation for plasma: MeOH was added to plasma samples at a 1:1 ratio. Samples were vortex mixed for 30 sec, then centrifuged (1,600 g, 15 min at 4 °C), and 500 μ L of the supernatant were collected for subsequent enzymatic hydrolysis.

Enzymatic hydrolysis for plasma, urine and cell culture medium: Five hundred microliters of each sample were transferred into a 5 mL tube containing 500 μ L of 0.2 M sodium acetate buffer (pH 5.2) and 25 μ L of β -glucuronidase, and incubated overnight at 37 °C.

Liquid-liquid extraction for hydrolysed cell culture medium: Three successive liquid-liquid extractions of the hydrolyzed cell culture medium samples were performed by addition of 1 mL of tert-butyl methyl ether preceded by alkalization of the sample (pH 11) with 1M NaOH. After vigorous vortex mixing and centrifugation (1,600 *g*, 5 min), the organic phase was collected and evaporated to dryness under nitrogen flow. The residue obtained was dissolved in 3 mL of 0.2 M sodium acetate buffer (pH 5.2) and subjected to the further extraction.

Solid phase extraction (SPE) for plasma, urine, and cell culture medium: The totality of the sample was applied on to a 1 cm³ (30 mg) OASIS MCX SPE column (Waters, Milford, Massachusetts), immediately followed by 2 mL of 0.1 M HCl and 2 mL of MeOH. The eluates were discarded, and the compounds of interest eluted into a glass tube using 2 mL of a 5% NH₄OH/MeOH solution. The solution was evaporated to dryness under nitrogen flow and the tubes further left open in a desiccator with phosphorus pentoxide (P₂O₅) overnight to eliminate residual water.

Derivatization procedure: To the dry residue, 50 μ L of ethyl acetate and 50 μ L of TFAA were added. After incubation at 70 °C for 30 min, the samples were cooled to room temperature and dried under nitrogen flow. The obtained residue was dissolved in 100 μ L of ethyl acetate and 1 μ L was analyzed by GC-MS.

Chromatographic equipment and conditions

An Agilent 6890N gas chromatograph equipped with an Agilent 5975 mass selective detector and an Agilent MSD Productivity ChemStation for GC and GC-MSD Systems software was used in the GC-MS analysis after automatic injection of 1 µL of final sample extracts. Injection was performed by a Combi-PAL autosampler (CTC Analytics, Zwingen, Switzerland) in the splitless mode with pulse pressure (40 psi) at 250 °C. A capillary column J&W DB-5ms (30 m × 0.25 mm × 0.25 μm) from Agilent was employed for the separation. The carrier gas was helium at a constant flow of 1 mL/min. The oven temperature program was optimized as follows: The initial column temperature of 120 °C was maintained for 1 min and then raised to 150 °C at 10 °C/min and hold at 150 °C for 5 min, followed by a temperature ramp of 7.5 °C/min to 300 °C, with a 2 min hold. Total separation run time was 31 min. The transfer line temperature was 280 °C and the electron energy was 70 eV. A solvent delay time of 3 min was added to avoid solvent overloading. The Full Scan mode was used for identification of the compounds and characterization of the mass spectra, and the Selected Ion Monitoring (SIM) mode was used for quantification of the compounds (scan range 50-600 m/z).

The selected ions used for the quantitative measurements were m/z 140, m/z 117, m/z 203 and m/z 230 for phenylpropanolamine; m/z 345, m/z 69, m/z 216, m/z 232 and m/z 248 for 4-hydroxy-3-methoxybenzylamine hydrochloride; m/z 272, m/z 91, m/z 175 and m/z 181 for 1-benzylpiperazine; m/z 326, m/z 173, m/z 200 and m/z 229for 1-(3-trifluoromethylphenyl)piperazine.

Method validation

Linearity: Working calibrators for plasma and urine (0.016, 0.08, 0.4, 2 and 10 μ g/mL) and for cell culture medium (0.625, 1.25, 2.5, 5 and 10 μ g/mL) were injected, along with a blank sample (IS, no analytes). The linearity of the method was evaluated by the square correlation coefficient (R²) of the regression curves obtained by plotting the peak-area ratio between each analyte and the IS against analyte concentration. Five calibration curves were independently obtained, and the mean slopes calculated.

Limits of detection and quantification (LOD and LOQ): The sensitivity of the method was determined by the limit of detection (LOD) and the limit of quantification (LOQ). Briefly, the 0.016 μ g/mL (plasma and urine) and 0.625 μ g/mL (cell culture medium) working calibrations were progressively diluted with blank plasma, urine or cell culture medium. A signal-to-noise ratio of 3 and 10 were considered adequate for estimating the LOD and LOQ, respectively. The lowest concentration found to fit these criteria was injected at least five times and the coefficient of variation (CV%) was calculated.

Sample stability: The sample stability was evaluated at three levels of concentration in plasma and urine samples (0.016 μ g/mL, 0.4 μ g/mL and 10 μ g/mL) and in cell culture medium samples (0.625 μ g/mL, 2.5 μ g/mL and 10 μ g/mL). The working calibrators were prepared and extracted on the same day (day 0) or stored at 4 °C and -20 °C and extracted after 7 days (plasma, urine and culture medium) or 30 days (plasma and urine). The stability was evaluated comparing the signal of samples extracted and analysed 7 and 30 days after being stored with the signal of the same freshly processed samples (day 0). The CV% was calculated.

Derivatized samples stability: The stability of the derivatized samples was evaluated at three levels of concentration in derivatized plasma and urine extracts (0.016 μ g/mL, 0.4 μ g/mL and 10 μ g/mL) and in derivatized cell culture medium extracts (0.625 μ g/mL, 2.5 μ g/mL and 10 μ g/mL). These samples were stored at 4 °C and injected for five days. The stability was evaluated comparing the signal of samples injected during five days after being derivatized, with the signal of the same freshly derivatized samples. The CV% was calculated.

Precision: Inter-day and intra-day precision were evaluated at three levels of concentration in plasma, urine and cell culture medium samples, and expressed as the CV%.

For all matrices, the inter-day precision was determined by the analysis, at five different days, of five independent samples (0.016 μ g/mL, 0.4 μ g/mL and 10 μ g/mL for plasma and urine; 0.625 μ g/mL, 2.5 μ g/mL and 10 μ g/mL for cell culture medium).

The intra-day precision of the method was performed by extracting five times the same plasma and urine sample $(0.016 \ \mu g/mL, 0.4 \ \mu g/mL and 10 \ \mu g/mL)$ and the same cell culture medium sample $(0.625 \ \mu g/mL, 2.5 \ \mu g/mL and 10 \ \mu g/mL)$. The intra-day precision of the equipment was determined by analyzing five times, on the same day, one sample at three different concentrations $(0.016 \ \mu g/mL, 0.4 \ \mu g/mL and 10 \ \mu g/mL)$ mL for plasma and urine; $0.625 \ \mu g/mL, 2.5 \ \mu g/mL and 10 \ \mu g/mL)$ mL for cell culture medium).

Accuracy: The accuracy of the method was assessed five times for BZP and TFMPP at three different concentrations 0.016 μ g/mL, 0.4 μ g/mL and 10 μ g/mL for plasma and urine, and 0.625 μ g/mL, 2.5 μ g/mL and 10 μ g/mL for cell culture medium, by calculating the percent deviation between the calculated value and the nominal value [accuracy (%) = (mean calculated concentration/nominal concentration) × 100].

Extraction efficiency: The extraction efficiency for each analyte was assessed at three different concentrations 0.016 μ g/mL, 0.4 μ g/mL and 10 μ g/mL for plasma and urine, and 0.625 μ g/mL, 2.5 μ g/mL and 10 μ g/mL for cell culture medium, using five replicates for each concentration.

Samples were prepared by spiking blank plasma, urine and cell culture medium with each analyte, extracted, and the IS added to each eluate before evaporation. The control samples were prepared by extracting blanks of each matrix and the analyte and IS solutions were added to each eluate before evaporation. For each analyte at each concentration, extraction efficiency was assessed by comparison of the peak area ratios of analyte to IS from extraction samples and control samples.

The recovery of the IS was independently tested at the 2 μ g/mL and 1 μ g/mL concentrations for 4-hydroxi-3-methoxybenzylamine hydrochloride and phenylpropanolamine, respectively.

Evaluation of interferences and specificity: Several blank samples of each matrix were processed and injected to evaluate the presence of chromatographic interferences at the retention times of the studied compounds. The GC-MS chromatograms of spiked samples and blank samples were then compared.

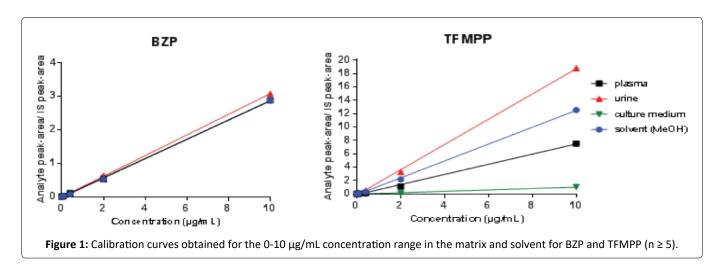


Table 1: Monitoring conditions of the test compounds in SIM mode.

Compound	Time interval (min)	Retention time (min)	Quantification ion (m/z)	Confirmation ions (m/z)
Phenylpropanolamine	3-6	3.5	140	117; 203; 230
4-hydroxy-3-methoxybenzylamine	6-9.5	6.85	345	69; 216; 232; 248
BZP	9.5-31	11.2	272	91; 175; 181
TFMPP	9.5-31	11.4	326	173; 200; 229

Proof of applicability

The applicability of the method could not be tested with real human samples since blood and/or urine samples from intoxicated individuals were not available for testing. To overcome this difficulty, we applied the method for the quantification of BZP and TFMPP in the extracellular exposure medium of primary rat hepatocytes after single and combined incubation with both drugs for 24 h. After the incubation period, the extracellular media was removed for the quantification of BZP and TFMPP using the validated method. Additionally, potential metabolites that could be formed due to the metabolic activity of the primary hepatocytes, were searched for using the Full Scan mode.

Results and Discussion

Sample preparation

In order to decrease the complexity of plasma samples, the protein precipitation before SPE extraction was considered necessary. In urine samples, the hydrolysis of conjugates before SPE extraction and GC-MS analysis was necessary, since glucuronide and sulfate conjugates are not volatile, precluding the direct analysis. Furthermore, it has been previously described the elimination of BZP and TFMPP as conjugates both in humans [21,25] and animals [26-28]. The major challenge of this work was the low sensitivity of the method for TFMPP in the cell culture medium and its consequent lack of linearity. A liquid-liquid extraction before SPE extraction was optimized in order to increase the method sensitivity for TFMPP in cell culture medium and solve this challenge.

Matrix effect

The matrix effect is one of the main sources of error in

analytical methods. Therefore, it is an important aspect to consider in the analysis of analytes in matrices as varied as those studied in this work, as it might impact all stages of chromatographic analysis (injection, separation, detection) and impair the quality of the results: Masking the peaks of the compounds of interest (false negatives); falsely identifying substances as impurities (false positives); increasing or decreasing the analytic signal of the detector [29-31].

Taking into consideration the diversity of matrices that we intended to validate, the matrix effect was evaluated by comparing the different calibration curves obtained in the matrix and solvent (MeOH). As observed in Figure 1, a significant matrix effect was noted, especially for TFMPP. Therefore, the method was validated for each matrix separately. As can be depicted in Figure 1, the calibration curve obtained with the cell culture medium presented a lower slope for TFMPP, and the analytical signal appears to be decreased, which can be explained by the high lipid content of the cell culture medium. This also explains the success of the liquid-liquid extraction.

Gas chromatographic separation

The 4-hydroxy-3-methoxybenzylamine hydrochloride (IS) showed an extractive and chromatographic behavior similar to the analytes and adequate retention time in plasma and urine. However, in cell culture medium, the liquid-liquid extraction drastically decreased the sensitivity of this IS, so another compound, phenylpropanolamine, was tested as IS to be used in the cell culture medium. This compound showed an extractive and chromatographic behavior similar to the analytes and adequate retention time and for these reasons was selected as the IS to be used in this matrix.

The SIM ions were selected taking into account their

specificity and their abundance in the mass spectra. For each substance at least four ions were chosen, one for quantification and the others for confirmation (Table 1). The selected

quantification ions and the retention times for phenylpropanolamine, 4-hydroxy-3-methoxybenzylamine, BZP and TFMPP are shown in Table 1. The chromatograms in Figure 2

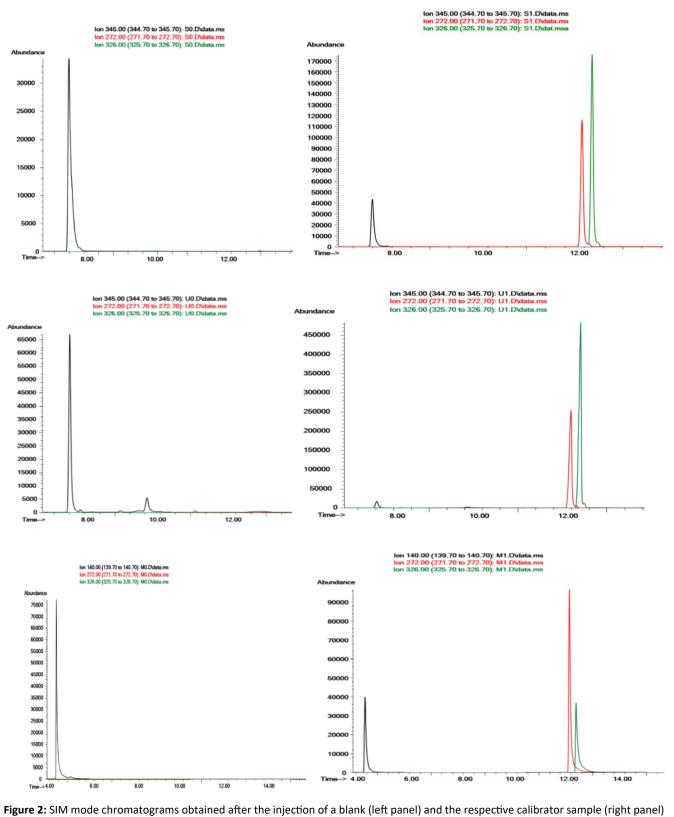


Figure 2: SIM mode chromatograms obtained after the injection of a blank (left panel) and the respective calibrator sample (right panel) spiked with a mixture of phenylpropanolamine (IS), 4-hydroxy-3-methoxybenzylamine (IS), BZP, and TFMPP at a concentration of 10 μ g/mL, of plasma (A), urine (B) and cell culture medium (C). The chromatograms were obtained with the following ions: m/z 140 for phenylpropanolamine (3.5 min), m/z 345 for 4-hydroxy-3-methoxybenzylamine (6.85 min), m/z 272 for BZP (11.2 min) and m/z 326 for TFMPP (11.4 min).

	Equation	Range (µg/mL)	R ²
Plasma			
BZP y = 0.1391x - 0.00635		0-10	0.9996
TFMPP	y = 0.2147x - 0.01388	0-10	0.9993
Urine			
BZP	y = 0.3233x + 0.00010	0-10	1.0000
TFMPP	TFMPP y = 0.5118x - 0.01997		0.9996
Cell culture medium			
BZP	y = 0.1970x - 0.03478	0-10	0.9983
TFMPP	y = 0.2140x - 0.03845	0-10	0.9986

Table 2: Calibration curves for BZP and TFMPP in plasma, urine and cell culture medium.

(n ≥ 5)

Table 3: LOD and LOQ (µg/mL) for BZP and TFMPP in plasma, urine and cell culture medium.

	LOD (µg/mL)	LOD (µg/mL)		
	BZP	TFMPP	BZP	TFMPP
Plasma	0.004 (11%)	0.004 (12%)	0.016 (13%)	0.016 (12%)
Urine	0.002 (12%)	0.002 (11%)	0.008 (4%)	0.008 (6%)
Cell culture medium	0.156 (18%)	0.312 (21%)	0.312 (20%)	0.625 (15%)

(CV%)

correspond to SIM mode chromatograms, where it is possible to see a good separation between the compounds and a low effect of bleeding, which proves that the column is adequate for this analysis.

Method validation

Linearity: Calibration curves were linear for all analytes in the range of 0-10 μ g/mL. This range of linearity was observed for the three matrices studied. The acceptable criteria for linearity is a calibration curve with a square correlation coefficient of 0.99 or higher [32], and in this study the square correlation coefficients were always greater than 0.99 (Table 2).

Limits of detection and quantification: The LOD for both analytes was 0.004 μ g/mL in plasma and 0.002 μ g/mL in urine. In cell culture medium, the LOD for BZP was 0.156 μ g/mL and for TFMPP was 0.312 μ g/mL. The LOQ for both analytes was 0.016 μ g/mL in plasma and 0.008 μ g/mL in urine. In cell culture medium, the LOQ for BZP was 0.312 μ g/ mL and for TFMPP was 0.625 μ g/mL. As depicted in Table 3, the coefficient of variation was equal or below 20% for both BZP and TFMPP in plasma, urine and cell culture medium.

Samples stability: The samples stability was evaluated by comparing the signal of the analytes in samples processed at day 0 with the signal of samples stored at different temperatures and processed 7 and 30 days later. A coefficient of variation of 20% was considered acceptable. The samples stability data can be depicted in Table 4. After 7 days in plasma and urine, the two compounds are stable for the three concentrations tested at two temperatures (4 °C and -20 °C). After 30 days in plasma, the two compounds are stable for

the three concentrations at -20 °C, with the exception of the lowest TFMPP concentration at 4 °C. Marginal increases in CV% values were also noted for BZP (0.016 μ g/mL at 4 °C) and for TFMPP (10 μ g/mL at 4 °C and 0.016 μ g/mL at -20 °C). In urine, both substances are stable for the three concentrations at -20 °C after 30 days, with only a slight increase over 20% CV for the lowest concentration. However, at 4 °C, the calculated CV% is generally much higher. Thus, it is possible to conclude that the two substances are stable in plasma and urine even after 30 days of storage, especially if stored at -20 °C. In cell culture medium, the samples stability was only assessed after 7 days, and it is possible to conclude that, overall, the two compounds are stable for all concentrations at both temperatures, with the exception of the highest BZP concentration (10 μ g/mL) when stored at 4 °C.

Derivatized samples stability: The derivatized samples stability was evaluated by comparing the signal of the analytes in samples injected on day 0 with the signal of samples injected in the following 5 days. The coefficient of variation was determined and values below 20% were considered acceptable. The stability of the derivatized samples is shown in Table 5. At the lowest concentration, BZP and TFMPP are stable in plasma derivatized samples until 2 days after derivatization. For the intermediate (0.4 μ g/mL) and the high (10 µg/mL) concentrations, BZP and TFMPP are stable until 4 days after derivatization. In urine derivatized samples, BZP and TFMPP are stable for all the concentrations until 2 days after derivatization. In cell culture medium, BZP and TFMPP are stable for all the concentrations, at least 5 days after derivatization. Based upon these data, it is recommended that the extracts are analyzed within a relatively short period after derivatization.

		Concentration (µg/mL)	Plasma : (CV%)	Plasma sample stability CV%)		Urine sample stability (CV%)		Cell culture medium sample stability (CV%)	
			BZP	TFMPP	BZP	TFMPP	BZP	TFMPP	
Day 7	4 °C	0.016	6	9	9	11	-	-	
		0.4	16	7	12	16	-	-	
		0.625	-	-	-	-	15	4	
		2.5	-	-	-	-	17	15	
		10	6	6	15	14	30	15	
-20 °C	0.016	6	6	7	12	-	-		
	0.4	9	4	9	15	-	-		
	0.625	-	-	-	-	8	8		
	2.5	-	-	-	-	15	16		
		10	10	13	15	13	23	12	
Day 30	4 °C	0.016	22	112	11	53	-	-	
		0.4	3	16	25	10	-	-	
	10	6	22	45	33	-	-		
	-20 °C	0.016	7	24	26	23	-	-	
		0.4	1	5	9	9	-	-	
		10	1	9	20	8	-	-	

 Table 4: Stability of BZP and TFMPP in plasma, urine and cell culture medium.

"-": Not determined.

 Table 5: Stability of BZP and TFMPP in derivatized plasma, urine and cell culture medium.

	Concentration (µg/mL)	Derivatized plasma sample stability (CV%)		Derivatized urine sample stability (CV%)		Derivatized cell culture medium sample stability (CV%)	
		BZP	TFMPP	BZP	TFMPP	BZP	TFMPP
Day 1	0.016	8	4	6	2	-	-
	0.4	0	7	7	6	-	-
	0.625	-	-	-	-	4	7
	2.5	-	-	-	-	2	1
	10	2	3	7	7	2	1
Day 2	0.016	18	9	13	14	-	-
	0.4	3	4	15	15	-	-
	0.625	-	-	-	-	3	2
	2.5	-	-	-	-	5	7
	10	5	3	14	15	1	2
Day 3	0.016	61	54	93	72	-	-
	0.4	12	2	28	26	-	-
	0.625	-	-	-	-	0	1
	2.5	-	-	-	-	3	9
	10	9	6	24	27	8	4

Day 4	0.016	126	134	92	117	-	-
	0.4	22	18	54	53	-	-
	0.625	-	-	-	-	4	8
	2.5	-	-	-	-	1	3
	10	11	8	52	72	0	3
Day 5	0.016	139	139	-	-	-	-
	0.4	43	39	-	-	-	-
	0.625	-	-	-	-	12	15
	2.5	-	-	-	-	1	9
	10	17	15	-	-	2	0

Table 6: Inter-day precision for BZP and TFMPP in plasma, urine and cell culture medium.

	Concentration (µg/mL)	Inter-day precis	sion (CV%)
		BZP	TFMPP
Plasma	0.016	6	11
	0.4	9	11
	10	9	11
Urine	0.016	11	14
	0.4	12	8
	10	13	8
Cell culture medium	0.625	11	11
	2.5	12	8
	10	11	8

Table 7: Intra-day precision for BZP and TFMPP in plasma, urine and cell culture medium.

	Concentration (µg/mL)	Intra-da	y precision of the method (CV%)	Intra-day p	recision of the equipment (CV%)
		BZP	TFMPP	BZP	TFMPP
Plasma	0.016	5	6	2	0.8
	0.4	4	13	2	2
	10	9	15	0.6	1.3
Urine	0.016	6	3	2	3
	0.4	6	8	3	3
	10	9	10	7	7
Cell culture	0.625	1	7	4	3
medium	2.5	8	12	1	4
	10	5	11	2	4

Precision: The precision, expressed as coefficient of variation, should not exceed 15% [32]. This method showed satisfactory precision (inter-day and inter-day), since the coefficient of variation never exceeded 15%. The inter-day precision was less than or equal to 14% for all concentrations of the two substances in all matrices (Table 6). In plasma, the coefficient of variation varied between 6% and 11%, in urine between 8% and 14% and in cell culture medium between

8% and 12%. The intra-day precision of the method was also less than or equal to 15% for all concentrations of the two analytes in all matrices (Table 7). In plasma, the coefficient of variation ranged between 4% and 5%, in urine from 3% to 10% and in cell culture medium between 1% and 12%. The intraday precision of the equipment was less than or equal to 7% for all concentrations of the two substances in all matrices. In plasma, the coefficient of variation varied between 0.6% and **Table 8:** Accuracy for BZP and TFMPP in plasma, urine and cell culture medium.

	Concentration	Accuracy (%)	
	(µg/mL)	BZP	TFMPP
Plasma	0.016	91	127
	0.4	101	96
	10	100	99
Urine	0.016	86	82
	0.4	101	107
	10	100	104
Cell culture medium	0.625	100	95
	2.5	96	94
	10	101	95

Table 9: Extraction efficiency of BZP and TFMPP in plasma, urine and cell culture medium.

	Concentration	Extraction e	fficiency (%)
	(µg/mL)	BZP	TFMPP
Plasma	0.016	95	85
	0.4	84	79
	10	96	89
Urine	0.016	101	108
	0.4	97	93
	10	90	102
Cell culture	0.625	91	99
medium	2.5	81	101
	10	76	85

2%, in urine from 2% to 7%, and in cell culture medium from 1% to 4%.

Accuracy: The accuracy, calculated as the percentage of target concentration, was 91-127% for plasma, 82-107% for urine and 94-101% for cell culture medium (Table 8). The accepted limit for this parameter is 100 ± 20% [32], which was met by all test conditions, except for lowest TFMPP concentration (0.016 μ g/mL) in plasma, for which the calculated accuracy was 127%.

Extraction efficiency: The calculated recovery percentage for 4-hydroxi-3-methoxybenzylamine hydrochloride, after five independent determinations, was of $84\% \pm 9.5\%$ (CV% = 11) in plasma, $90\% \pm 9.7\%$ (CV% = 11) in urine. The calculated recovery percentage for phenylpropanolamine, after three independent determinations, was of $63\% \pm 9.6\%$ (CV% = 15) in cell culture medium. The extraction efficiency for BZP and TFMPP at the three concentrations in all matrices is described in Table 9. In plasma, the extraction efficiencies were between 79% and 96%, in urine between 90% and 108%, and in cell culture medium between 76% and 101%. The accepted limit for this parameter is $100 \pm 20\%$, which was always met, except for the highest BZP concentration ($10 \ \mu g/mL$) in cell culture medium, for which the extraction efficiency was 76%.

Evaluation of interferences and specificity: Several blank samples of each matrix were prepared and injected to evaluate chromatographic interference. In all three matrices studied no interfering peaks at the retention times of analytes or IS were detected as can be depicted in Figure 2.

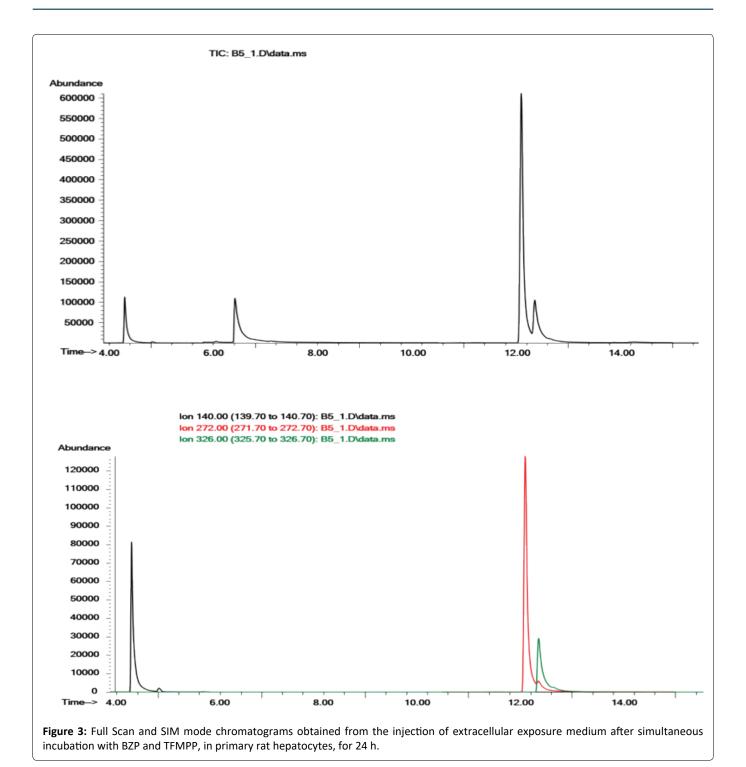
Proof of applicability

For the proof of applicability, primary rat hepatocytes were incubated with BZP and TFMPP, alone and in combination, for 24 h. Primary hepatocytes have considerable metabolic capacity and are widely accepted as an *in vitro* model suitable for *in vivo* extrapolations. The toxicokinetic data available for both BZP and TFMPP indicate that humans and rats share the same metabolic pathways. The main metabolites produced by both species are ring-hydroxylated metabolites that can be subsequently conjugated and excreted as glucuronides and/or sulfates [21,25,26,28,33,34]. For both species, BZP and TFMPP are largely excreted unchanged or as glucuronide conjugates, and the quantitative analysis of the parent drugs has been used in clinical and forensic settings [21,25,33]. With our successful quantifications, the metabolic interaction between these substances in primary hepatocytes was established [35].

Under these experimental conditions, no metabolites were observed for either BZP or TFMPP. In Figure 3, representative Full Scan and SIM mode chromatograms obtained from the injection of extracellular exposure medium after incubation with BZP and TFMPP, in primary rat hepatocytes, for 24 h can be depicted. No interfering peaks were observed in these samples. In humans, the main metabolites of both BZP and TFMPP correspond to the ring 4-hydroxylated and 3-hydroxylated metabolites that are excreted alongside with the unchanged parent drugs, after conjugation [21,25,26,28,33]. In the case of TFMPP, for forensic and clinical purposes, the analytical data must be carefully interpreted since some of the excreted metabolites are common to other drugs such as leflunomide [26]. Therefore, the detection of the parent drug is in this case preferred. Additionally, in cases of acute intoxications high drug concentrations can be achieved in the liver and metabolic enzyme saturation often occurs, thus emphasizing the interest of the detection of the unchanged parent drugs for unequivocally proving the intake of the drug.

Conclusion

This GC-MS method was developed, optimized and validated successfully for the concurrent analysis of BZP and TFMPP. The method showed high sensitivity and an adequate performance for the simultaneous detection and quantification of these drugs in the three studied matrices (plasma, urine and cell culture medium) using only an IS in each matrix. Thus, this method proved to be an alternative to methods previously published in the literature, which are not validated or only rely on the validation for the single drugs. Linearity, detection and quantification limits, precision and accuracy of the method were all suitable for scientific, clinical and forensic use. In addition, the method proved to have quite



acceptable extraction efficiency and absence of interferences. The selected IS proved to be suitable and easy to obtain, which avoids the use of synthesis or acquisition of multiple deuterated standards. Using a commercially available IS, the method showed a high sensitivity and linearity for a wide range of concentrations, which is particularly relevant for the clinical and forensic applications of the method. A proof of applicability of the present method was its successful use to demonstrate metabolic interactions between BZP and TFMPP involved in the *in vitro* hepatotoxicity of the combination of these two drugs, commercialized under the name "Legal X" as a legal alternative to 3,4-methylenedioxymethamphetamine (MDMA or ecstasy). Therefore, the developed analyti-

cal methodology has direct application not only for research purposes but also for the control of drug abuse in humans.

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Conflict of Interest

All authors declared no conflict of interest.

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