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Posted Date: 25 December 2023

doi: 10.20944/preprints202312.1801.v1

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## Article

# Development and Validation of an Analytical Method for the Simultaneous Detection of Sulfonamides, Trimethoprim and Dapsone in Honey by Liquid Chromatography-Tandem Mass Spectrometry

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**Abstract:** The presence of antibiotic residues in honey has recently become a growing public health concern due to the overuse of these products in beekeeping and their transfer to the food industry. The aim of this study was to develop and validate a simple, selective and sensitive method for the simultaneous identification and quantification of residues of twenty-two sulfonamides, trimethoprim and dapsone in honey using a rapid and reliable liquid chromatography–tandem mass spectrometry (LC–MS/MS) method. The developed method showed good linearity, specificity, precision (repeatability and intra-laboratory reproducibility) and recovery. The decision limit (CC $\alpha$ ) ranged between 2.16–22.12  $\mu\text{g/kg}$  for CC $\alpha$ , while the detection capability (CC $\beta$ ) was between 2.16– 2.67–27.36  $\mu\text{g/kg}$ . The method proved to be rapid, reliable and effective for the determination of sulfonamides, trimethoprim and dapsone from honey samples. The average recoveries were estimated to be between 70 and 106%. The relative standard deviations (RSDs) were between 6–18%. In practice, the method is essential to check the honey sold on the market and ensure its safety before it reaches the consumer.

**Keywords:** analytical method; antibiotic residues; sulfonamides; trimethoprim; dapsone; honey; liquid chromatography-tandem mass spectrometry

## 1. Introduction

Sulfonamides, sometimes also known as sulfa drugs, are antibiotics specifically targeting bacteria that can cause infections. These antibiotics are usually broad-spectrum antibiotics that act against a wide range of bacteria and are used to treat a variety of bacterial infections. Sulfonamides account for 11% of total sales of antimicrobial veterinary medicines in Europe in 2011 [1]. Trimethoprim and dapsone, which have a similar effect to sulfonamides, are often administered together with some sulfonamides in pharmaceutical products. When used in combination with trimethoprim and dapsone, their bacteriostatic effect is synergistic and is enhanced by the inhibition

of bacterial dihy-dropteroate synthetase and dihydrofolate reductase. It is known that high levels of sulfonamides, dapsone and trimethoprim in food can have various adverse effects on human health, including increased bacterial resistance to antimicrobial agents, allergic reactions or possible carcinogenicity [2–4].

Honey is produced and processed by honeybees from the nectar and honeydew of plants and contains natural sugars such as glucose and fructose, amino acids, enzymes, minerals and antioxidants [3]. It has been used as a food and medicine for millennia, honeybee products being considered to be natural and healthy, able to prevent infections and even used in wound therapy [4]. However, bee products are produced in a natural environment, which itself may be contaminated. Sources of contamination include the immediate environment and beekeeping itself. Xenobiotics from the environment come from industry, agriculture, or individual treatment. These include organic contaminants (polychlorinated biphenyls – PCBs), heavy metals (Pb, Cd), insecticides (organochlorines - OC, organophosphorus pesticides -OP, carbamates), herbicides (asulam), bactericides (streptomycin), fungicides (vinclozolin, iprodione and methyl tiophanate, captan and difenoconazole), and radioactive isotopes e (40K and 137Cs [5]. The contamination caused by beekeeping practices is more targeted and varies from country to country or even region to region. The active ingredients of these local treatments can spread quickly and are carried into the hives by the bees. Of the many treatments, antibiotic therapy is commonly used in beekeeping to treat the major bacterial diseases, American Foolbrood (AFB) [6] and European Foolbrood (EFB), the protozoan disease Nosema and Varroa [7]. To control foolbrood diseases, the tetracycline family of antibiotics has mainly been used to treat hives for decades, but the bacteria responsible for AFB and EFB have gradually become resistant. Aminoglycosides, macrolides, quinolones, nitrofurans and the sulfonamide family are therefore also used in various countries for the treatment of infested hives or for prophylaxis [8]. Sulfonamides (SA) such as sulfathiazole or sulfamonomethoxine are preferred in some regions of the world. Therefore, it is necessary to routinely check whether honey contains residues of these drugs in order to protect consumers from a possible allergy risk and to avoid the development of bacterial resistance. To date, no maximum residue limits for SAs in honey have been set in the European Union. This lack of regulation for the honey matrix could change in the future, especially if studies showing the presence of veterinary drug residues in beehives complete the picture [9] and lead to the establishment of MRLs for honey as for other foodstuffs. In the European Union, only recommended concentrations for minimum performance limits for confirmatory methods have been proposed, set at 50 µg/kg for trimethoprim and all sulfonamides. Dapsone is banned for use in agricultural food production, but no maximum residue limits (MRLs) have been set for sulfonamides and trimethoprim, although this family of veterinary medicines is covered by Commission Regulation (EU) No 37/2010 and 470/2009/EC (formerly 2377/90/EC) [10–12].

A receptor test called Sulfasensor can detect SA residues in honey or other spices [13] and a rapid screening method for sulfonamides in honey uses a flow injection system coupled with a liquid waveguide capillary cell [14]. However, reliable and sensitive methods are needed to unequivocally detect and confirm residues in honey. Several analytical methods have been described to determine one or more SAs, using high-performance liquid chromatography with UV [7,15,16] or fluorescence detection [17–20] or liquid chromatography combined with mass spectrometry [21–24]. One or more SAs are sometimes detected and quantified together with other molecules in multifamily methods [25–29]. To our knowledge, there is no published LC-MS/MS confirmatory method validated according to Commission Decision No. 2002/657/EC [30] that considers the 22 SAs proposed in our method.

The aim of this work was to develop and validate a simple, selective, reliable and sensitive method for the simultaneous determination - identification and quantification - of residues of twenty-two sulfonamides, trimethoprim and dapsone in honey using LC-MS / MS. This is indeed the first work in which a validation has been performed for this combination of target substances/matrix according to the requirements of Decision 2002/657/ EC [31] and its amending directive SANCO 2726/2004 [32] for the determination of residues of "banned and unauthorized" substances.

In this context, the determination of antibiotic residues in honey is considered important, while the development of accurate, selective, rapid and multi-screening methods for the determination of such residues is still urgently needed.

No such analytical method has been previously reported in Romania. Moreover, the method has been implemented and is currently successfully applied to simultaneously analyse the above-mentioned veterinary drug residues in Romanian veterinary laboratories within the framework of the national residue control plan for honey products.

## 2. Materials and Methods

### 2.1. Chemicals and Reagents

Certified standards of sulfacetamide, sulfaguanidine, sulfanilamide, sulfisomidine, sulfadiazine, sulfathiazole, sulfapyridine, sulfamerazine, sulfamethazine, sulfamethizole, sulfamethoxypyridazine, sulfamethoxydiazine, sulfamonomethoxine, sulfamoxol, sulfadoxine, sulfamethoxazole, sulfisoxazole, sulfabenzamide, sulfadimethoxine, sulfaquinoxaline, sulfanitran, dapsone and trimethoprim and the internal standards sulfadimidine-13C6 and sulfamethoxazole-13C6 were purchased from Sigma-Aldrich (St. Quentin Fallavier, France), while sulfachloropyridazine and sulfaphenazole were obtained from Dr. Ehrenstorfer (Augsburg, Germany).

LC-MS grade acetonitrile, n-hexane, methanol, formic acid and acetone together with hydrochloric acid (0.1 N) and ammonium acetate buffer (0.2 mol/l, pH 5.3) were obtained from Sigma-Aldrich (St. Quentin Fallavier, France). A Millipore Simplicity 185 purification system (Millipore, USA) was used in order to obtain de-ionized double distilled water.

All chemicals and reagents used were of analytical grade, unless otherwise specified.

For the solid-phase extraction Strata<sup>TM</sup>-X (500mg/ 6ml) cartridges from Phenomenex were used.

### 2.2. Standards Solutions

Individual standard stock solutions of the analytes and of the internal standards-sulfadimidine-13C6 and sulfamethoxazole-13C6 (100 µg/ml) were prepared in acetonitrile and further stored in glass ambered bottles. Individual working solutions of the analytes dapsone and trimethoprim (1 µg/ml), as well as a mixed working solution of the analysed sulfonamides (1 µg/ml) were prepared by the appropriate dilution of the standard stock solutions with hydrochloric acid. Individual working solutions of the internal standards (10 µg/ml) were also prepared by the appropriate dilution of the individual standard stock solutions with hydrochloric acid and appropriately stored.

Eight mixed calibration solutions containing the SAs and trimethoprim analytes in the concentration range 5-150 ng/ml and dapsone in the range 2.5-25 ng/ml, as well as the internal standards at a fixed concentration of 20 ng/ml were prepared daily, by serial dilution of the mixed working solution of sulfonamides, the individual working solution of dapsone, the individual working solution of trimethoprim and of the individual working solutions of the internal standards in the aqueous mobile phase (0.1% *v/v* HCOOH in water).

### 2.3. Matrix Calibrations

Eight matrix-matched calibration solutions containing all the SAs and trimethoprim analytes in the range of 5-150 µg/kg and in the range of 2.5-25 µg/kg for dapsone and the internal standards at a fixed concentration of 20 µg/kg, were prepared by subjecting "blank" honey samples (5 g) to solid-phase extraction and further spiking of the extract with the appropriate volumes of the mixed sulfonamides working solution, the individual working solutions of dapsone and trimethoprim, as well as the individual working solutions of the internal standards.

## 2.4. Traceability

Honey samples were gathered from various botanical origins of north central and east Romania and analysed in a laboratory which is overseen by The National Veterinary Health and Food Safety Authority. The method presented in this work was developed, validated and accredited in order to offer a rigorous procedure for the determination and quantitative confirmation of sulfonamides, trimethoprim and dapsone residues in honey matrices at low levels of concentrations to meet the requirements.

The blank material was a blend of 20 different types of honey (varying in colour and texture), all of which were examined for residues before being used in the validation process, none of them contained detectable levels of SAs, trimethoprim and dapsone. 24 different honey samples were utilized to examine matrix effects.

## 2.5. Sample Preparation and Extraction

### *Testing Sample*

Approximately 50g of "blank" honey was heated at 45°C in a water bath for 1h. A quantity of 5 g of "blank" honey were accurately weighted in a 50 ml polytetrafluoroethylene (PTFE) centrifuge tube and then spiked with 100 µl of each individual working solution of internal standards-sulfadimidine-13C6 and sulfamethoxazole-13C6 of concentration 1 µg/ml. 10 ml of a 0.1 N hydrochloric acid (HCl) were added to the previously obtained mixture, which was further vortexed for 15 minutes and then sonicated for 15 minutes, until complete dissolution. Then, the testing sample was centrifuged at 3500 rpm for 10 minutes and stored over night at room temperature for a better static extraction. The next day it is applied to the solid phase extraction (SPE) cartridges.

### *Fortified (spiked) Sample*

The fortified (spiked) sample was prepared by accurately weighing 5g of honey in a 50 ml PTFE centrifuge tube. 250 µl of mixed sulfonamides working solution (1µg/ml), 25 µl of dapsone working solution (1 g/ml), 250 µl of trimethoprim working solution (1 µg/ml), as well as 100 µl of each working solution of internal standards (1 µg/ml) were added to the sample. 10 ml of HCl (0.1N) were further added in the centrifuge tube. The probe was vortexed for 15 minutes and then sonicated for 15 minutes. The sample was afterwards centrifuged at 3500 rpm for 10 minutes and stored over night at room temperature. The next day it is applied to the SPE cartridges.

### *Blank*

The blank was prepared of 10 ml HCl (0.1 N) and 100 µl of each working solution of internal standards (1 µg/ml), vortexed for 15 minutes and then sonicated for 15 minutes until complete dissolution and then centrifuged at 3500 rpm for 10 minutes and stored over night at room temperature. The next day it is applied to the SPE cartridges.

### *Samples Purification*

The solid-phase extraction (SPE) was performed on Strata™-X cartridges. The cartridges were pre-conditioned with 10 ml of methanol and 10 ml of ammonium acetate (0.2 mol/l, pH 5.3). The samples were percolated through the cartridges, these being afterwards rinsed with 10 ml of water and vacuum dried for 5 minutes. The retained analytes were further eluted with 10 ml of methanol. The eluate was concentrated on a water bath at 50°C to near dryness under a nitrogen stream. The solution was reconstituted with 1000 µl of HCl (0.1 N).

## 2.6. Instrumentation

The LC-tandem MS system included a Bruker UHPLC advance pump, a PAL HTC-xt autosampler, and a Bruker EVOQ Elite triple-quadrupole mass spectrometer with an electrospray ionization (ESI) interface. Data acquisition was done with Bruker MSWS software version 8.

A Pursuit XRs Ultra C18 column (100 mm × 2 mm/ 2.8 µm) for chromatographic separation (Agilent Technology, Netherlands). The temperature of the column was kept constant at 35°C. Two mobile phases were used in a multi-step binary elution gradient: phase A: water containing 0.1 % (v/v) HCOOH, and phase B: acetonitrile containing 0.1 % (v/v) HCOOH.

For both the standard and sample solutions, the flow rate was 200  $\mu\text{L}/\text{min}$  and a volume of 30  $\mu\text{L}$  was injected. The gradient is presented in Supplementary Material Table S1.

## 2.7. Method Validation

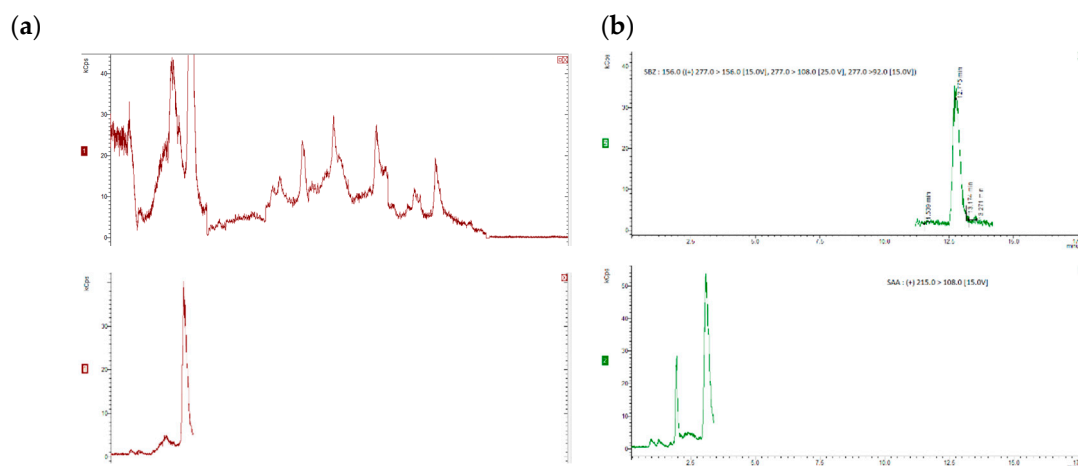
According to Commission Decision 2002/657/EC [31] and its amending guideline SANCO/2006/3228 [32], the method was validated as a quantitative confirmatory method. The retention times and relative ion intensities of matrix-matched standard calibration solutions were monitored to establish identification criteria for each analyte (Supplementary Material Table S1).

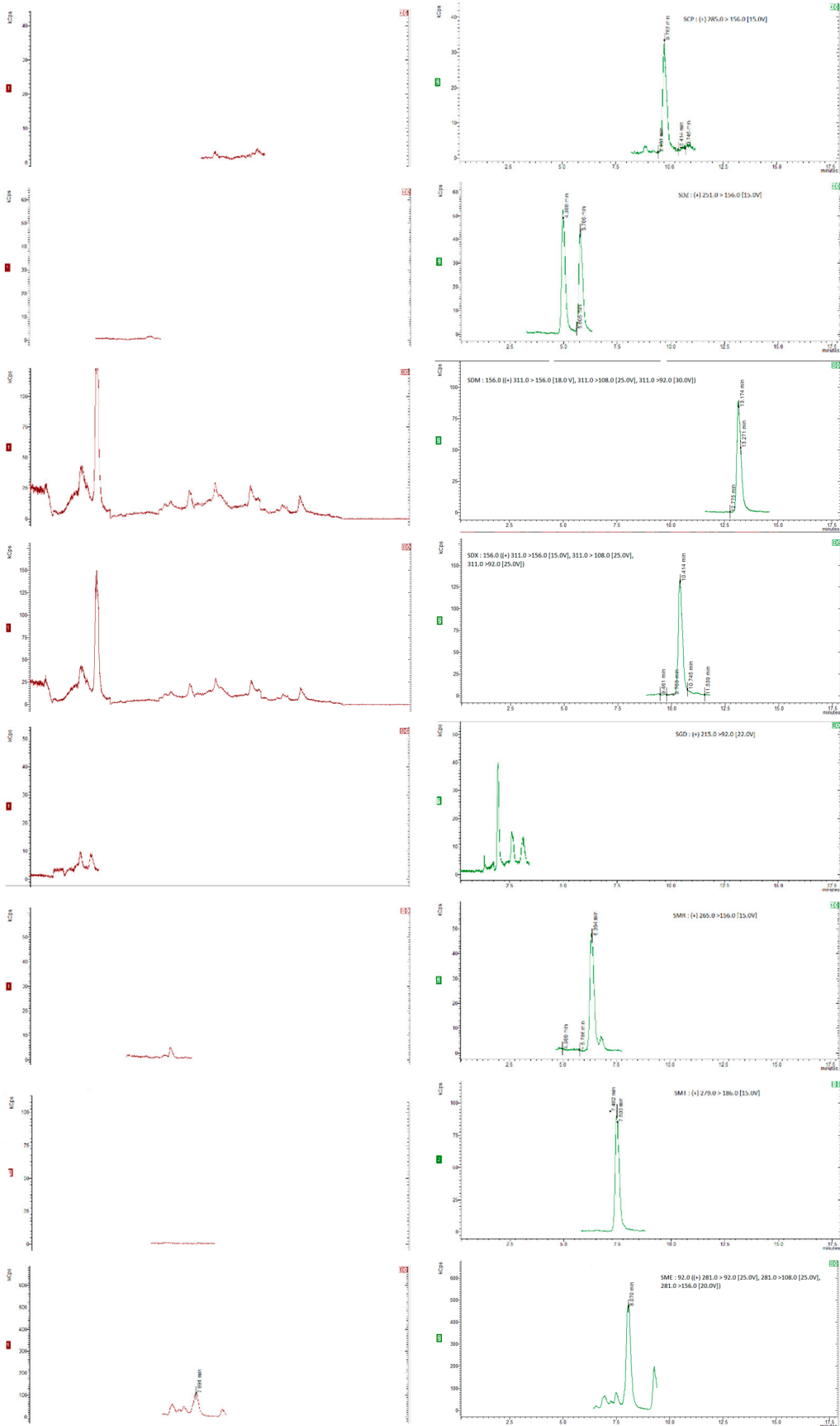
Specificity, sensitivity, linearity, precision (repeatability and within-laboratory reproducibility), trueness (in terms of recovery), decision limit ( $\text{CC}\alpha$ ), and detection capability ( $\text{CC}\beta$ ) were the performance criteria examined in the validation study [31–34] (Supplementary Material Table S2). Twenty organically cultivated honey samples of various origins were analyzed to determine the specificity. The linearity of the method was assessed using regression analysis of solvent calibration and matrix-matched calibration solutions, using the ratio of the standard area (derived from the most intense transition) to internal standard area against analytes concentrations.

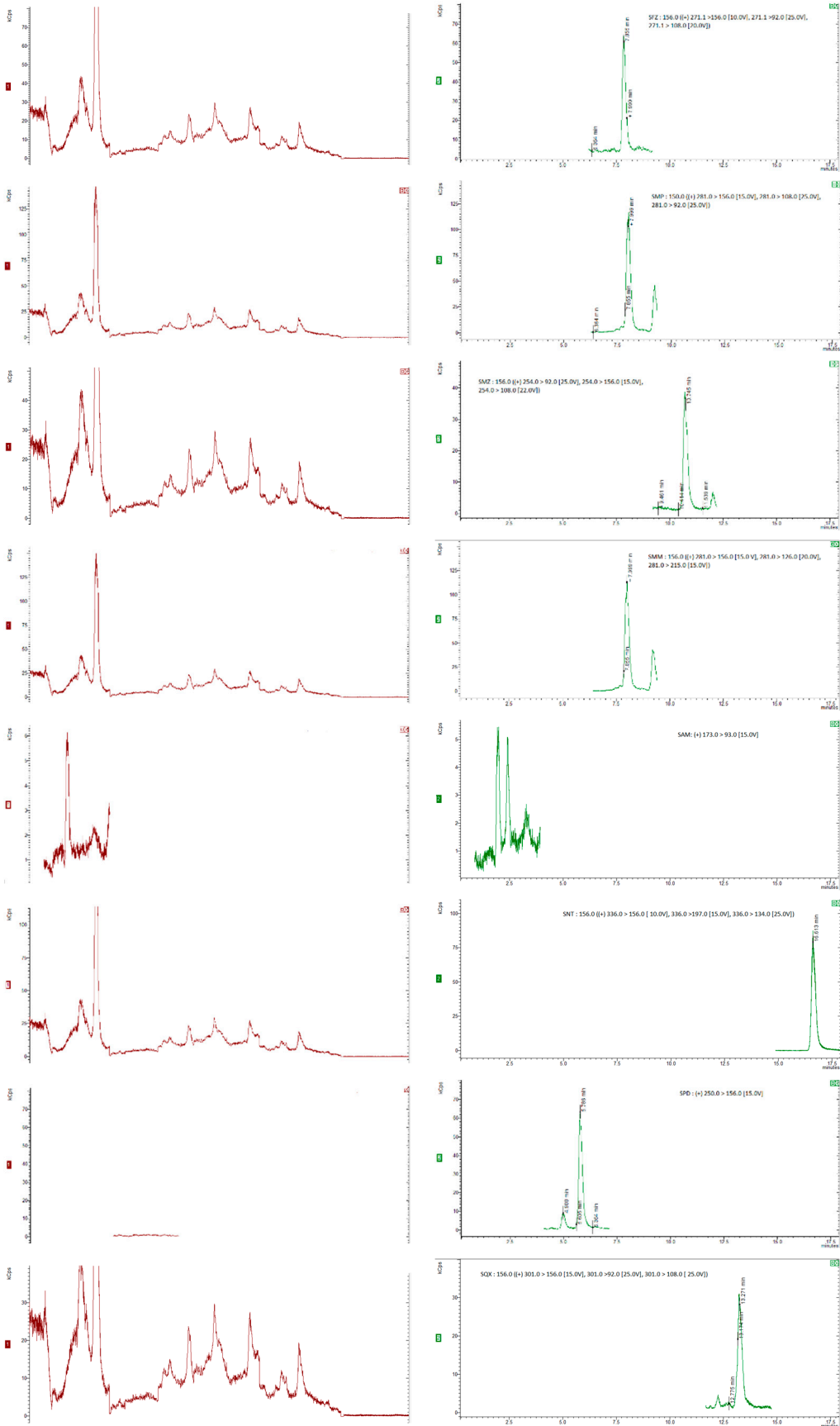
"Blank" honey samples (5 g) were spiked with appropriate amounts of the standard working mixture of analytes and the working mixture of deuterated standards for the recovery experiments, resulting in concentration levels of 25, 50, 75, 100  $\mu\text{g}/\text{kg}$  for SAs and trimethoprim, of 2.5, 5, 7.5, 10  $\mu\text{g}/\text{kg}$  for dapsone and 20  $\mu\text{g}/\text{kg}$  for the internal standards. To allow the sulfonamides to sufficiently bind to the sugars in the honey, the spiked samples were kept at room temperature for at least 1 hour [28,35]. Six replicates per spiking level were analysed and evaluated using a matrix-matched calibration curve prepared as previously described. The chromatograms were obtained for internal standards, for "blank" honey sample spiked with 20  $\mu\text{g}/\text{kg}$  of the internal standards and blank honey samples spiked with 20  $\mu\text{g}/\text{kg}$  of the internal standards and 50  $\mu\text{g}/\text{kg}$  of each compound (5  $\mu\text{g}/\text{kg}$  for dapsone), as shown in Figure 1 and Figure 2.

Each series, which included a matrix calibration curve and 24 spiked samples, was prepared on two distinct days for a total of 48 spiked samples with various time, operator, and LC–MS/MS calibration/operation status. These tests determined trueness (in terms of percent recoveries) and accuracy (in terms of repeatability and within-laboratory reproducibility in terms of respective percent relative standard deviations).

The decision limit ( $\text{CC}\alpha$ ) and detection capability ( $\text{CC}\beta$ ) were determined in accordance with the ISO Standard 11843, linear regression method, as proposed in European Decision No 2002/657/EC [31]. Both critical limits,  $\text{CC}\alpha$  and  $\text{CC}\beta$  were calculated using calibration curves obtained from "blank" honey samples spiked at four concentration levels of 25, 50, 75, 100  $\mu\text{g}/\text{kg}$  for SAs and trimethoprim and of 2.5, 5, 7.5, 10  $\mu\text{g}/\text{kg}$  for dapsone – six replicates per level – and subjected to SPE, using the ratio of the signal of the target analyte's less intense transition to that of the internal standard applicable in each case.

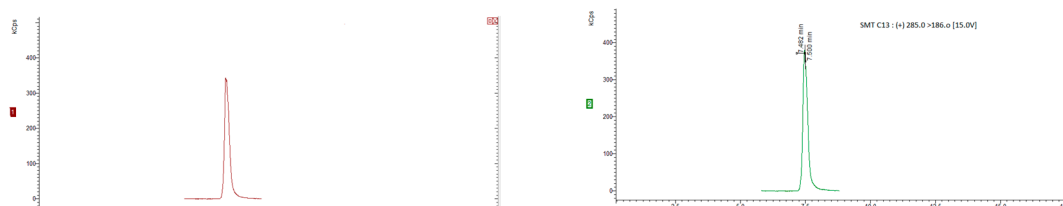


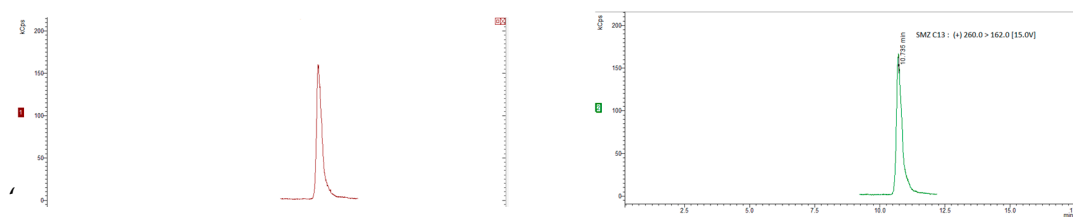






**Figure 1.** Chromatograms of “blank” honey samples with 20 µg/kg of the internal standards (a) and honey samples with 20 µg/kg of the internal standards and spiked with 50 µg/kg (5 µg/kg for dapsone) of each of the compounds of interest (b).





**Figure 2.** Chromatograms of “blank” honey samples with 20 µg/kg internal standards (a) and honey samples with 20 µg/kg internal standards and spiked with 50 µg/kg of each of the compounds of interest (b).

### 3. Results and Discussions

#### 3.1. LC-MS/MS Method Development

Sulfonamides are banned as antibiotics in honey in the European Union, as no maximum residue levels have been set for these foods. However, with SANCO/2006/3228 [36] a guideline has been established that sets recommended levels for substances for which there are no legal MRLs or minimum performance levels (MRPLs). The performance of analytical methods in relation to CCα must be below these values, even if these values are only analytical limits without regulatory significance. Non-authorized substances are monitored with two MRM (Multiple Reaction Monitoring) transitions. Since SAs contain a nitrogen atom, optimization in ESI mode was quite simple and the responses obtained in positive mode were high. It is known that the signals are weaker in negative ionization [21]. With the exception of some sulfonamides, the usual fragment ions with values of  $m/z$  156, 108 and 92 were used.

Optimization of the liquid chromatography settings with focus on the signals of the analytes and selection of the optimal separation conditions for the mobile phase, especially for the isomers SDX-SDM ( $m/z$  311) and SMM-SMP-SME ( $m/z$  281). A comparison of 0.1 % formic acid and 0.1 % PFPA in the aqueous mobile phase was investigated. PFPA had the advantage of successfully retaining the two most polar SAM and SGD analytes, but it interfered with the analytes' signals, so formic acid was finally chosen. Two columns were tested to observe the performance of the analytical columns: Pursuit XRs Ultra C18 column (100 mm × 2 mm/ 2.8 µm particle size, Agilent Technology) and Phenomenex Aqua C18 column (150 mm × 2.0 mm/ 3 µm particle size, Waters). The first column was chosen because it offers high reproducibility for the analytes investigated in this work.

Most published methods for measuring SAs in foods of various origins include at least one solid-phase extraction (SPE) step using polar [37], non-polar [38–40], or strongly cation-exchanging sorbents [41]. The advantage of an SPE step prior to LC–MS/MS analysis is that suppression by matrix components is reduced under certain circumstances and the detection limit of the method is lowered in most cases.

Simple extractions with HCl (0.1 N) in water versus trichloroacetic acid (TCA) (0.03 N) in water were both tested while improving the conditions for sample preparation. As a result, simple extraction procedures are required for the analysis of samples from national monitoring plans. TCAs show an inability to sufficiently break the bonds between the sulfonamides and the honey sugars. As the extraction yield and sensitivity were insufficient when using TCA, HCl was chosen to separate the sulfonamides from the honey components.

#### 3.2. Sample Extraction Procedure

The acid hydrolysis of the sulfonamides bound to the sugar was followed by an SPE purification phase using the sample extraction technique. The recovery of “blank” honey samples spiked with the target chemicals was used to evaluate the efficacy of the different sample extraction procedures. The influence of the pH of the honey sample after acid hydrolysis was investigated in the range of 2–7. The best sulfonamide recoveries occurred when the sample was adjusted to a pH of 4, where the

target species were in their isoelectric form (based on their respective pKa values). During our initial experiments, it was discovered that the cartridges were frequently clogged by undissolved material – most likely wax in the honey sample – resulting in low recovery values; as a result, the cartridges were conditioned and the sample solution was diluted to 250 mL with HCl (0.1 N) to reduce the likelihood of the sulfonamides re-binding to sugars and to avoid blocking of the SPE cartridges. The SPE cartridges were conditioned with methanol followed by acetate buffer 0.2 mol/L, pH 5.3, and extraction was carried out as rapidly as possible. When compared to acetonitrile, methanol was used to elute the target chemicals since it yielded greater recoveries. The eluate was concentrated on a water bath at 50°C to near dryness under a nitrogen stream. The solution was reconstituted with 1000 µl of HCl (0.1 N).

### 3.3. Method validation

#### 3.3.1. Specificity, Linearity, Sensitivity

The method's specificity was tested by examining a total of twenty "blank" honey samples. No interfering peaks overlapping with the analyte peaks were seen in the SRM chromatograms in these "blank" sample experiments, confirming appropriate specificity for the trace detection of the target compounds. The signal-to-noise ratios were considerably improved by using a more "narrow" resolution window on the first quadrupole, with the peak width set at 0.2 Da (FWHM) (Q1). Figure 1(a) and (b) show SRM chromatograms of a "blank" honey sample and a "blank" honey sample spiked with 2.5 mg kg<sup>-1</sup> of all the chemicals, respectively.

In the range of 0.5–100 mg L<sup>-1</sup>, calibration curves were created for both solvent and matrix matched calibration. The concentration range 0.5–100 mg L<sup>-1</sup> of matrix-matched calibration solutions corresponded to 0.1–20 mg kg<sup>-1</sup> of the target compounds in the honey sample. The slope (a), intercept (b), standard deviation of the slope (sa), standard deviation of the intercept (sb), and correlation coefficient ( $r^2 > 0.99$ ) of the linear regression equations were determined.  $R^2 > 0.99$  in all cases, suggesting good linearity in the concentration range examined. Supplementary Material Table S2 shows the linear regression parameters, as well as the instrumental limits of detection (LODs) determined from the matrix matched calibration solutions (where  $LOD = 3sb/a$ ). The LODs ranged from 0.3 to 0.9 mg L<sup>-1</sup> (corresponding to 0.06–0.18 mg kg<sup>-1</sup> in the honey sample), indicating strong detection sensitivity.

#### 3.3.2. Accuracy

Recovery studies utilizing fortified "blank" honey samples spiked with the target components were used to assess method accuracy – trueness and precision. For three different operating days ( $q=3$ ), six replicates were performed per day ( $m=6$ ) at each of the three concentration levels (1.5, 2.5, and 5.0 mg kg<sup>-1</sup>) ( $p=3$ ). The percent recovery, R percent, was computed using the following formula:  $R \% = (C_c/C_s) \times 100$ , where  $C_c$  is the analyte concentration in the spiked samples and  $C_s$  is the analyte concentration given to a "blank" honey extract after extraction, but before LC–MS/MS analysis.

The peak area ratio of each analyte versus the respective internal standard was used to derive  $C_c$  and  $C_s$  from the matrix matched calibration curves (except for trimethoprim (TMP)). The mean recovery values,  $R_m$  percent, ranged from 70 to 106 percent at the three concentration levels (Supplementary Material Table S2), meeting the requirements for trueness in the CD 2002/657/EC [42]. A total of 18 spiked samples per spiking level were subjected to one way analysis of variance (ANOVA) to estimate the mean recovery at each level; mean recovery values,  $R_m$  %, ranged from 70 to 106 % at the three concentration levels (Supplementary Material Table S2). ANOVA was also used to determine technique repeatability and within-laboratory reproducibility; experimental F values,  $F_{exp}$  (calculated as between-days variation over within-day variance), were consistently lower than the theoretical F value,  $F_{theor}$  (2, 15, 0.05) of 3.68. For all the compounds, the method's mean repeatability (expressed as percent relative standard deviation, %RSD percent) and within-laboratory reproducibility (expressed as percent relative standard deviations, RSDR %) both ranged from 6 to

18 % at the three concentration levels (data shown in Supplementary Material Table S2), meeting the CD 2002/657/EC criteria (i.e., RSD%<22 % ) [42].

The "modified" or "truncated" Horwitz equation (recommended by Thompson for concentrations less than 120 mg kg<sup>-1</sup> [43]) was used to calculate the "target" standard deviation sH (sH=140.22C, where C is the compound concentration in mg kg<sup>-1</sup>) and, as a result, the respective "target" relative standard deviations, RSDH. H, or the "Horrat" measure, is defined as follows:

$$H = RSDR/RSDH$$

where RSDR is the within laboratory reproducibility's experimental standard deviation. For all of the compounds, the "Horrat" values were r1 (Supplementary Table S2); a reasonable condition for intralaboratory validation is that the "Horrat" be in the range 0.2–1 [44].

3.3.3. Uncertainty

According to the LGC/VAM protocol [45], the experimental design used during method validation permitted estimate of measurement uncertainty from validation data. Given the minimal significance of type B contributions to uncertainty, the standard uncertainty, u(Y), can be determined using the following equation:

$$[u(Y)]^2 = [u(P)]^2 + [u(R_m)]^2$$

where u(P) represents the uncertainty associated with method precision and u(R<sub>m</sub>) represents the uncertainty associated with method recovery. The formula can be used to calculate relative uncertainty:

$$(u(Y))/Y = \sqrt{([u(P))/P]^2 + ([u(R_m))/R_m]^2}$$

The relative standard deviation of within-laboratory reproducibility, RSDR, obtained from the ANOVA test at each spiking level can be used to express the phrase u(P)/P [45].

To see if R<sub>m</sub> was substantially different from 1, a t-test was used with the formula  $t = 1 - uR_m/R_m$ , where uR<sub>m</sub> is the standard deviation of the recoveries per spiking level. The t values for SPD, STZ, SMR, SMZ, and SMTX were lower than the coverage factor k=2.33 [32], and hence R<sub>m</sub> values at the three spiking levels were not statistically different from 1.

As a result, there was no need for any additional bias correction, and the uncertainties associated with the recoveries were computed using relative standard deviations, uR<sub>m</sub>=R<sub>m</sub>.) The results of uncertainties are presented in Table 1.

**Table 1.** The values of uncertainties for the target compounds (µg/kg).

Compounds	Standard Uncertainty u <sub>c</sub> (µg/kg)	Expanded Uncertainty u <sub>c</sub> (k=2) (µg/kg)	Final results
Sulfaguanidine	4,386	8,772	50 ± 8,772
Sulfacetamide	4,309	8,618	50 ± 8,618
Sulfanilamide	6,155	12,310	50 ± 12,310
Sulfisomidine	4,157	8,314	50 ± 8,314
Sulfadiazine	3,184	6,368	50 ± 6,368
Sulfathiazole	4,030	8,060	50 ± 8,060
Sulfapyridine	3,468	6,936	50 ± 6,936
Sulfamerazine	3,330	6,660	50 ± 6,660

Sulfamethazine	2,773	5,546	50 ± 5,546
Sulfamethizole	3,433	6,866	50 ± 6,866
Sulfamethoxypyridazine	2,434	4,868	50 ± 4,868
Sulfamonomethoxine	3,498	6,996	50 ± 6,996
Sulfameter	3,012	6,024	50 ± 6,024
Sulfachloropyridazine	3,309	6,618	50 ± 6,618
Sulfadoxine	3,613	7,226	50 ± 7,226
Sulfamethoxazol	3,167	6,334	50 ± 6,334
Sulfisoxazole	3,195	6,390	50 ± 6,390
Sulfabenzamide	2,853	5,706	50 ± 5,706
Sulfadimethoxine	2,995	5,990	50 ± 5,990
Sulfaquinoxaline	3,053	6,106	50 ± 6,106
Sulfaphenazole	3,060	6,120	50 ± 6,120
Sulfanitran	4,484	8,968	50 ± 8,968
Trimethoprim	4,985	9,970	50 ± 9,970
Dapsone	0,421	0,842	5 ± 0,842

3.3.4. Decision Limits (CCalpha) and Detection Capabilities (CCbeta)

The decision limit (CCalpha), where α is 1% and the detection capability (CCbeta), where β is 5%, were calculated for this method using the following equations:

$$CC\alpha = cy + 2,33 \times sR$$

where: sR = standard deviation of the intra-laboratory reproducibility of the intercept

cy = the concentration corresponding to the y-intercept;

$$CC\beta = CC\alpha + 1,64 \times sR$$

The values of CCα and CCβ are shown in Table 2.

**Table 2.** Decision limits (CCα) and detection capabilities (CCβ) of the method (µg/kg).

Compounds	CCα (µg/kg)	CCβ (µg/kg)
Sulfaguanidine	3,50	4,33

Sulfacetamide	2,44	3,02
Sulfanilamide	22,12	27,36
Sulfisomidine	7,78	9,62
Sulfadiazine	2,30	2,85
Sulfathiazole	7,64	9,45
Sulfapyridine	5,58	6,90
Sulfamerazine	4,37	5,40
Sulfamethazine	2,16	2,67
Sulfamethizole	3,53	4,37
Sulfamethoxypyridazine	5,59	6,91
Sulfamonomethoxine	5,94	7,35
Sulfameter	9,36	11,57
Sulfachloropyridazine	7,20	8,91
Sulfadoxine	7,35	9,09
Sulfamethoxazol	7,89	9,76
Sulfisoxazole	7,87	9,74
Sulfabenzamide	7,66	9,48
Sulfadimethoxine	7,61	9,42
Sulfaquinoxaline	8,22	10,16
Sulfaphenazole	8,34	10,32
Sulfanitran	9,52	11,77
Trimethoprim	7,67	9,48
Dapsone	2,45	3,03

## 5. Conclusions

After finalizing the implementation and validation of the method in the laboratory, the method was verified by participating in a Proficiency Testing scheme organised by Fapas T02353 in which 59 international laboratories attended. Considering the z-scores obtained by the laboratory for this method, we can conclude that the method is adequate, rigorous and efficient for the correct identification and quantification of sulfonamides in honey. The method has low limits of determination which is very important considering the prohibition of sulfonamides in honey. Our method is used currently for the surveillance of antibiotic residues in honey, serving as an initial effective assessment tool.

Romania is an important honey producer, exporting honey on the European market, and therefore, this method is essential for investigating how safe this product is before reaching a large number of consumers from both national and European level.

**Supplementary Materials:** The following supporting information can be downloaded at the website of this paper posted on Preprints.org. Table S1: Fragmentation conditions for the target compounds, Table S2: The values of performance parameters obtained through the internal validation of the method.

**Author Contributions:** Conceptualization, M.M., G.V.V.; methodology, A.V.-L., G.V.V., M.-M.P.; formal analysis, G.V.V.; investigation, L.M.C., A.V.-L., G.V.V., A.I.G.-H., A.-A.C.; writing—original draft preparation, L.M.C., A.V.-L., G.V.V., A.I.G.-H., A.-A.C., M.-M.P.; writing—review and editing, G.V.V., A.I.G.-H., A.-A.C.; supervision, M.M.. All authors have read and agreed to the published version of the manuscript.

**Funding:** This research received no external funding.

**Data Availability Statement:** The data presented in this study are available on request from the corresponding author.

**Acknowledgments:** The authors would like to thank the D.S.V.S.A. Laboratory from Cluj Napoca for their support.

**Conflicts of Interest:** The authors declare no conflict of interest.

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