Determination of Glyphosate and Glufosinate in Human Serum by MonoSpin TiO Extraction and Liquid Chromatography–Tandem Mass Spectrometry

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We developed and validated an assay for determination of glyphosate (GLYP) and glufosinate (GLUF) in human serum. Serum samples were extracted by using a MonoSpin® TiO column and analyzed by liquid chromatography–tandem mass spectrometry (LC–MS/MS). MonoSpin® TiO tends to specifically bind to phosphate groups. The assay was linear over a concentration range of 1–250 μg/mL. The recoveries for the 2 compounds were 1.0%–2.3%. The intra- and inter-day variations were <15%. Precision and accuracy were 5.6%–12.7% and 97.0%–103.9%, respectively. The validated method was applied to quantify the GLYP and GLUF content in the serum of GLYP and GLUF-poisoned patients. In conclusion, the method was successfully applied for accurate determination of GLYP and GLUF in serum obtained from patients with GLYP and GLUF poisoning.

Keywords: glyphosate, glufosinate, MonoSpin TiO column, serum

Introduction

Glyphosate (N-[phosphonomethyl]glycine) (GLYP) and glufosinate (GLUF) (DL-homoalanin-4-yl-[methyl]phosphinate ammonium salt) are phosphonic and amino acid group-containing chemicals that constitute an important category of pesticides and are extensively used as herbicides in many countries, including Japan. GLYP and GLUF-surfactant herbicides are the most frequently used nonselective herbicide. However, numerous cases of accidental and suicidal poisoning due to the ingestion of these herbicides in attempted suicides have been reported [1–6]. Although GLYP and GLUF herbicides poisoning was probably caused by surfactant [1], these surfactants are not disclosed by the manufacturers. Serum GLYP and GLUF concentrations were used for the alternative method of assessing poisoning states. Moreover, GLYP and GLUF products contained a surfactant—an isopropylamine salt or partial ammonium salt—before 2006. However, a part of GLYP products has been changed to contain 48% GLYP potassium salt in Japan since 2006. Thereafter, the incidence of mortality cases with hyperkalemia due to GLYP herbicide ingestion gradually increased. Although GLYP herbicide poisoning causes hyperkalemia, GLUF herbicide poisoning is still caused by a surfactant in the formulation. Although GLYP herbicide ingestion may be suspected by the presence of hyperkalemia, the early diagnosis and appropriate treatment of GLUF and GLYP may be lifesaving.

Recently, we developed a simple derivatization method for gas chromatography (GC)–mass spectrometry (MS) analyses of GLYP and GLUF [7–9], because the previously reported sample preparation method, especially the derivatization procedure, is time consuming [10–16]. However, GLYP and GLUF were frequently not detected after derivatization, especially after acylated derivatized sample analyses.

Recently, Yoshioka et al. reported an analytical method using liquid chromatography (LC)–MS/MS for bialophos (BIA), GLYP, and GLUF without derivatization [17]. Although their external standardization for extraction is usually more convenient and less expensive, an internal standard (IS) can be a useful tool for negative analytical results. Therefore, we developed rapid extraction method using an IS for identification of GLYP and GLUF. This method used a MonoSpin® TiO column for extraction and LC–MS/MS for analysis.

Materials and Methods

GLUF ammonium and GLYP were obtained from Wako (Osaka, Japan). DL-2-amino-3-phosphonopropionic acid was purchased from Sigma (St. Louis, MO, USA). MonoSpin® TiO cartridges (5010-21706) were used for extraction from serum samples (GL Sciences, Tokyo, Japan). Acetonitrile for LC–MS and formic acid were purchased from Sigma-Aldrich (St. Louis, MO, USA) and Nacalai Tesque (Kyoto, Japan), respectively. All other chemicals and reagents were of analytical grade (Wako, Japan). Serum samples for the validation study were collected from volunteers from hospital staff who were not using any medication. Frozen serum samples (−30 °C) were stored at room temperature until complete thawing and vortex-mixed 1 min before use.

Calibration Curve and Quality Control Samples

GLYP and GLUF (10 mg/mL) were prepared in 10% methanol. These solutions were further diluted with 10% methanol to concentrations of 1 mg/mL, 100 μg/mL, 10 μg/mL, and 1 μg/mL and stored for a maximum of 3 months at 4 °C. These solutions were used as calibrators. A stock solution of DL-2-amino-3-phosphonopropionic acid (1 mg/mL) was prepared in 10% methanol and used as an IS stock solution. The

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IS stock solution was further diluted with 10% methanol to a concentration of 100 μg/mL. Blank serum samples (final volume of 100 μL) were spiked with 10 μL of calibrator (10 μg/mL), 5 μL of calibrator (100 μg/mL), 2.5 μL of calibrator (1 mg/mL), 5 μL of calibrator (1 mg/mL), 10 μL of calibrator (1 mg/mL), and 2.5 μL of calibrator (10 mg/mL) resulting in calibration samples containing 0.1 (1 μg/mL), 0.5 (5 μg/mL), 2.5 (25 μg/mL), 5 (50 μg/mL), 10 (100 μg/mL), and 25 (250 μg/mL) μg of each compound, i.e., GLYP and GLUF.

Each quality control (QC) solution was prepared in the same manner as mentioned above. QC samples were prepared by using blank serum samples (final volume of 100 μL) containing 0.1 μg/mL of the response and the slope (S) of calibration curves at 15 psi.

Precise and accurate results were obtained by using blank serum samples spiked with 4 QC samples. For the intra- and inter-day assays, 6 replicates of each concentration were prepared and analyzed. Four QC samples were prepared every time and analyzed for 3 consecutive days. The accuracy was measured as the percentage deviation from the nominal concentration. The results for the intra- and inter-day percent relative standard deviations (%RSDs) were obtained by using blank serum samples spiked with 4 QC samples. The intra- and inter-day precision were determined as %RSDs.

**Extraction Procedure**

**Sample Preparation and Extraction.** A MonoSpin® TiO column was used for extraction from serum sample. The human serum standards for calibration were prepared by spiking the standard working solution and IS solution (20 μL of 100 μg/mL; 2 μg) into a pool of blank human serum (final volume of 100 μL). QC samples at 4 concentrations were prepared by spiking the appropriate standard working solutions and IS solution into blank human serum (100 μL). A 400-μL aliquot of 0.1% acetic acid was added to 100 μL of the serum sample and mixed on a vortexer for 15 s.

The MonoSpin® TiO column was conditioned with 200 μL of methanol at 3000 rpm for 1 min followed by 200 μL of 0.1% acetic acid at 3000 rpm for 1 min. The samples were applied to the conditioned MonoSpin® TiO column. The column was centrifuged for 1 min at 3000 rpm. The column was washed with 200 μL H2O at 3000 rpm for 1 min. Finally, the samples were eluted with 200 μL acetonitrile at 3000 rpm for 1 min and 200 μL acetonitrile-ammonia solution (9:1) at 3000 rpm for 1 min in a new tube. The collected samples were dried under stream nitrogen at 45 °C. The resulting samples were dissolved in 100 μL acetonitrile-0.1% formic acid (80:20) and injected into the LC–MS/MS system.

The calibration curves, QC, and clinical serum samples were extracted by using the same procedure.

**LC–MS/MS Analysis**

**LC–MS/MS Conditions.** Chromatographic separation was performed by using an Obelise N 5-μm (2.1 mm × 150 mm) column (SIELC Technologies Inc., Wheeling, IL, USA) at 40 °C in an Agilent 1200 LC system (Agilent, Santa Clara, CA, USA). Separation was performed with isocratic elution of 0.1% formic acid–acetonitrile (80:20, v/v) at a flow rate of 0.4 mL/min. The total data acquisition time was 7.0 min/sample. The injection volume was 10 μL.

Electrospray ionization–tandem mass spectrometry (ESI–MS/MS) detection was achieved on an Agilent 6410 triple quadrupole tandem mass spectrometer. The positive ionization mode was used, and the ions were monitored in the multiple reaction monitoring (MRM) mode. ESI source parameters were as follows: high-purity drying gas (N2) flow rate, 6 L/min; temperature, 300 °C; capillary voltage, 4000 V; and nebulizer, 15 psi.

**Limit of Detection (LOD) and Limit of Quantitation (LOQ).** LOD was defined on the basis of the standard deviation (SD) of the response and the slope (S) of calibration curves at levels approximating the LOD according to the following formula: LOD = 3.3(SD/S). Similarly, LOQ was defined on the basis of the SD of the response and S of calibration curves at levels approximating the LOQ according to the following formula: LOQ = 10(SD/S) [18].

**Precision and Accuracy.** The intra- and inter-day percent relative standard deviations (%RSDs) were obtained by using blank serum samples spiked with 4 QC samples. For the intra- and inter-day assays, 6 replicates of each concentration were prepared and analyzed. Four QC samples were prepared every time and analyzed for 3 consecutive days. The accuracy was measured as the percentage deviation from the nominal concentration. The criteria for the acceptability of data included accuracy within ±15% of the nominal value and a precision within ±15% of the RSD except for the lowest LOQ, which could not exceed 20% of RSD [19].

**Extraction Recovery and Matrix Effects.** We evaluated the extraction recovery of GLYP and GLUF from the blank serum by analyzing 6 replicates of 2 sets of QC samples. The first set A consisted of 100 μL of blank serum added to the 4 QC solutions and IS with 400 μL of 0.1% acetic acid and then extracted by using a MonoSpin® TiO column. The second set B consisted of 100 μL of blank serum added to 400 μL of 0.1% acetic acid, and the 4 QC solutions and IS were added after MonoSpin® TiO extraction of the sample. Then, the samples were prepared in the same manner and analyzed by the LC–MS/MS system. A comparison of the mean GLUF and GLYP integrated peak areas, (set A/set B) × 100%, provided the extraction recovery of GLUF and GLYP expressed in percentage.

The 4 QC and IS solutions were dissolved in acetonitrile just before LC–MS/MS analysis (C). The ratio (B/C × 100) defines the matrix effects. The reproducibility of the extraction procedure was determined as %RSD.

**Patient's Samples**

**Case 1.** A 55-year-old Japanese female attempted to commit suicide by ingesting an unknown volume of Basta® (18.5% glufosinate). She was admitted to the hospital by her husband and then transferred to a city hospital for further examination on the same day. On admission to City Hospital, she had a Glasgow Coma Scale of 1–1–4 (Japan Coma Scale 300), a systolic blood pressure of 170 mmHg, and a heart rate of 132 beats/min. She was transported to the intensive care unit (ICU) and intubated. However, she had marked blood pressure reduction and tachyrrhymia. Over a 7-day period, she was extubated in the ICU.

**Case 2.** A 65-year-old Japanese male attempted suicide by ingesting approximately 250 mL of Roundup® (48% glyphosate-potassium salt). Thereafter, he was in agony because of vomiting and diarrhea and was transferred to our emergency department, approximately 2 h after ingestion. On admission, he received cardiopulmonary resuscitation with intubation. However, he died despite intensive care 7 months later.

**Results and Discussion**

The precursors and product ions for each compound are presented in Table 1. These conditions were slightly different from those reported elsewhere [17]. The typical MRM chromatograms of the blank serum and spiked low QC and IS are shown in Figure 1. GLYP, GLUF, and IS were not detected in the blank serum. These chromatograms revealed that there was no interfering peak derived from the endogenous components at the elution times of GLYP, GLUF, and the IS. The peaks corresponding to GLUF, IS, and GLYP were clearly observed at 3.35, 4.45, and 5.18 min, respectively (Figures 1–3).
The LOD of GLYP and GLUF was 0.5 μg/mL for each. Based on the above criterion, the LOQ of GLYP and GLUF was determined to be 1 μg/mL for each, when a 100-μL serum sample was used.

The validation data for the analytical method in terms of accuracy and precision are summarized in Table 2. Data from the QC samples that were run in sextuplicate at each concentration on each of these 3 days are also presented in Table 2.

The calibration curves of GLYP and GLUF were linear over the concentration range from 1 to 250 μg/mL for both compounds. A linear regression analysis of the plot of the peak-area ratios (y) of the target to the IS versus the target concentration was performed. The equations of the calibration curves were calculated and are shown in Table 2.

Table 1. Precursor ions, product ions and their respective collision energy for each compound

<table>
<thead>
<tr>
<th>Compound</th>
<th>Precursor ion (m/z)</th>
<th>Product ion 1 (m/z)</th>
<th>Product ion 2 (m/z)</th>
<th>Collision energy (V)</th>
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</thead>
<tbody>
<tr>
<td>Glyphosate</td>
<td>170.0</td>
<td>88.1</td>
<td>50.1</td>
<td>5</td>
</tr>
<tr>
<td>Glufosinate</td>
<td>182.1</td>
<td>65.1</td>
<td>33.1</td>
<td>7</td>
</tr>
<tr>
<td>DL-2-amino-3-phosphonopropionic acid (IS)</td>
<td>170.0</td>
<td>124.0</td>
<td>106.0</td>
<td>13</td>
</tr>
</tbody>
</table>

aQuantitative ion.
bQualitative ion.

determination of Glyphosate and Glufosinate

Figure 1. Liquid chromatography–tandem mass spectrometry chromatograms obtained by MonoSpin® TiO column extraction of (A) blank human serum, (B) blank human serum with low QC level (glufosinate [GLUF], 2.5 μg/ml; glyphosate [GLYP], 2.5 μg/ml), and the internal standard (IS). 1: GLUF, 2: IS, and 3: GLYP
concentrations was performed. The regression equations for GLYP and GLUF were $y = 0.0529x + 0.1102$ and $y = 0.1495x + 0.2370$, respectively. The SDs of the slope and the intercept in GLYP regression equation were 0.0060 and 0.0370, respectively. Similarly, the SDs of the slope and the intercept in GLUF regression equation were 0.0142 and 0.0423, respectively. The average coefficient of determination ($r^2$) was always $>0.998$. The experimental peak-area ratios were interpolated on the calibration curve, and the concentrations were back-calculated. The mean back-calculated concentrations approached the spiked concentrations, as shown by an RSD of $<15\%$.

The LOD using our previous method for GLYP and GLUF was slightly better than that using the presented method [7–9, 12, 13]; previously, GLUF was analyzed in human serum and had an LOD of 0.1 μg/mL [7–9, 12, 13]. These values appeared to be in the same order of magnitude as the values estimated by the methodology presented herein.

The GLUF and GLYP concentrations in the serum after a few hours of GLUF and GLYP ingestion were $>24\mu g/mL$ and more than a few thousand μg/mL, respectively [4, 8, 9]. Therefore, the linearity range included the LOD and LOQ values for GLYP and GLUF in this method. Although the LC–MS/MS method is usually superior to the GC–MS method, the linearity range for GLYP and GLUF obtained by using our presented method was narrower than that obtained by using the previously reported GC–MS method [7–9, 12, 13]. This narrow linearity range can probably be explained by the extraction recovery method using MonoSpin® TiO. However, Yoshioka et al. reported [17] that GLYP and GLUF concentration in serum was analyzed using LC–MS/MS, and the higher linearity ranges for both compounds are up to 20 μg/mL according to their paper. The LODs for GLYP and GLUF in their method were 0.03 and 0.07 μg/mL, respectively. Similarly, the LOQs for GLYP and GLUF were 0.09 and 0.1 μg/mL, respectively. This higher linearity range was 10 times lesser than that in our method. Although our method is suitable for the analysis of higher blood concentrations of GLYP and GLUF in acute intoxication, their method is suitable for the analysis of even trace levels after intensive care.

Assay Precision and Accuracy. The %RSD of the intra-day assay (n = 6) for the serum GLYP and GLUF concentration was $<12.7\%$, and good inter-day (n = 6) RSD values ($<12.1\%$) were also obtained, as shown in Table 2.

Extraction Recovery. Table 2 presents the results of the extraction recovery for the 4 QC levels. The overall

![Figure 2. MRM chromatograms of standard of GLUF (upper). MRM chromatograms obtained by MonoSpin® TiO column extraction of blank human serum with GLUF (10 μg/mL) (bottom)](image-url)
recoveries in the serum samples ranged from 1.6% to 2.3% for GLYP and GLUF at the 4 QC levels. The overall extraction recoveries of GLYP and GLUF from the serum were very low.

Although TiO tends to bind to phosphate groups, the MonoSpin disk is thin. Therefore, the amount of phosphate groups held by TiO should be small, but this binding leads to low recovery. However, ion suppression was not detected and was approximately 100% in all QCs. We have previously reported that the greatest loss of GLUF and GLYP during sample processing occurred during deproteinizing [7]. Moreover, ion suppression was approximately 50%–60%. Although the recovery from MonoSpin TiO was very low, ion suppression did not occur because the selectivity for phosphorous is high.

Yoshioka et al. reported [17] that 500 μL of serum was ultra-filtered, and 300 μL of passage was used for further preparation. Their validation samples for recovery of GLYP and GLUF were also prepared in 500 μL of serum at 0.2 μg/mL and 5 μg/mL after ultrafiltration performed by using the same procedure. Although the recovery ranged between 94.5% and 108.2%, the details of their calculation method and ion suppression data were not described in their paper. Therefore, it is difficult to compare their method with the method presented here.

**Application.** The validated method was then applied to an analysis of samples from patients with GLUF and GLYP.

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**Table 2.** Method performance: recovery, accuracy, precision, and matrix effect

<table>
<thead>
<tr>
<th>QC conc. (ng/mL)</th>
<th>Recovery (%)</th>
<th>Accuracy (%)</th>
<th>Intra-day Precision (%)</th>
<th>Inter-day Precision (%)</th>
<th>Matrix effect</th>
</tr>
</thead>
<tbody>
<tr>
<td>Glufosinate</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>1</td>
<td>1.6</td>
<td>97.0</td>
<td>11.7</td>
<td>10.6</td>
<td>103.9</td>
</tr>
<tr>
<td>2.5</td>
<td>1.7</td>
<td>100.3</td>
<td>6.0</td>
<td>5.1</td>
<td>105.4</td>
</tr>
<tr>
<td>40</td>
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<td>101.2</td>
<td>9.7</td>
<td>8.8</td>
<td>104.7</td>
</tr>
<tr>
<td>200</td>
<td>2.0</td>
<td>103.9</td>
<td>8.1</td>
<td>7.4</td>
<td>104.7</td>
</tr>
<tr>
<td></td>
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<td>97.3</td>
<td>12.7</td>
<td>9.9</td>
<td>98.6</td>
</tr>
<tr>
<td>Glyosphate</td>
<td></td>
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<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>2.5</td>
<td>2.3</td>
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<td>99.4</td>
<td>10.0</td>
<td>7.4</td>
<td>100.9</td>
</tr>
</tbody>
</table>

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Figure 3. MRM chromatograms of standard of GLYP (upper). MRM chromatograms obtained by MonoSpin® TiO column extraction of blank human serum with GLYP (10 μg/mL) (bottom)
poisoning. Patients' serum concentrations of GLUF and GLYP were 487 and 6758 μg/mL.

Conclusion

In conclusion, we described a method for the determination of GLYP and GLUF in human serum. This method needs only a simple centrifuge for sample extraction and can be easily implemented into routine practice compared with previously reported methods. Compared with the previously published GC–MS method, the LC–MS/MS method reduces the analysis time from 20 min to 7 min, which makes the application of this method to clinical samples possible, in addition to rapid toxicological screening of GLYP and GLUF-poisoning samples. Particularly, this method is suitable for the rapid screening of acute intoxication by GLUF and GLYP, rather than after intensive care treatment.

Compliance with Ethical Standards

Conflict of Interest. All authors declare that they have no conflict of interest.

Ethical Approval. All procedures performed in studies involving human participants were in accordance with the ethical standards of Tokai University School of Medicine. Informed consent was obtained from all individual participants included in the study.

References