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- 1 Sensitive and selective quantification of glyphosate and
- 2 aminomethylphosphonic acid (AMPA) in urine of the general population by
- 3 gas chromatography-tandem mass spectrometry
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- 17 Glyphosate
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- 22 Environmental exposures
- 23

24 Abstract

Glyphosate is the highest volume herbicide used worldwide, and its main biodegradation 25 product is aminomethylphosphonic acid (AMPA), both are listed as priority substances in the 26 Human Biomonitoring for Europe initiative (HBM4EU) which aims at improving policy by 27 28 filling knowledge gaps by targeted research. The objective of the current study was to 29 advance the sensitivity of an existing gas-chromatography tandem mass spectrometry analytical method to measure environmental population exposures. A 50% lower limit of 30 quantification of 0.05 μ g/L was achieved for both analytes by slight modifications in sample 31 32 work-up, and use of another isotope labelled internal standard. In a pilot study on 41 urine samples from the general German population, glyphosate and AMPA could be quantified in 33 66% and 90% of the samples respectively, which is a sufficient number to reliably describe 34

35 distributions of urinary concentrations in the non-occupationally exposed population.

36 **1. Introduction**

Glyphosate (N-(phosphonomethyl)glycine) is a broad-spectrum herbicide, its main 37 environmental biodegradation product is aminomethylphosphonic acid (AMPA), and both are 38 39 often detected as residues on crops and plants, and in foodstuff [1]. There are over 750 herbicidal products containing glyphosate; it is the highest volume used herbicide in the 40 world, with over 6 million metric tons applied worldwide in the past decade [2, 3]. 41 Glyphosate is classified as 'Group 2A - probably carcinogenic to humans' by the 42 43 International Agency for Research on Cancer [4], a classification refuted by international regulatory agencies [5-7]. 44

45 Human biomonitoring (HBM) studies indicate a ubiquitous exposure to glyphosate [8-19]. However, existing analytical methodologies vary considerably in both their sensitivities and 46 specificities, depending on the underlying analytical principles (e.g. LC-MS/MS, GC-MS/MS, 47 ELISA), their limits of detection/quantification (LOD/LOQ), and their use of appropriate 48 49 isotope labelled internal standards, all of which make comparisons between study results difficult. Depending on their purpose, some methods have been developed to cover 50 specialised exposures (e.g. occupational), or are adapted for use on human matrices, e.g. from 51 water or food surveillance methods, or do not include AMPA and glyphosate simultaneously. 52 To date, only ten HBM exposure studies on glyphosate have also analysed for AMPA. When 53 quantified, however, AMPA was found in similar concentration ranges as glyphosate. Urinary 54 AMPA moderately correlates with urinary glyphosate [11], but it is not fully understood if (or 55 how much of) urinary AMPA stems from human metabolism of glyphosate or what 56 57 proportion is due to AMPA uptake concurrently (e.g. via food) with glyphosate [20]. AMPA has been reported to be of toxicological relevance as well [21] and measurement of both 58 compounds allows for a more comprehensive exposure assessment. Consequently, both 59 glyphosate and AMPA have been selected as priority substances in the European project 60 HBM4EU [22] (www.HBM4EU.eu) with the intention to establish an EU-wide HBM survey 61 62 for these substances, including thorough quality assessment of applied methods and 63 continuous quality assurance to increase data comparability [23].

After screening existing analytical methods and previously published HBM data the need for 64 more sensitive analytical methods to quantify both glyphosate and AMPA in urine below 0.1 65 66 μ g/L was identified, in order to capture the median and reliably describe distributions in the general population. Gas-chromatography tandem mass spectrometry in combination with 67 isotope dilution quantification is likely to be the most appropriate method, allowing sensitivity 68 and selectivity. Thus, we decided to advance a previously published analytical method [11] to 69 fulfil these aims. We demonstrate the applicability of this advanced method by the analysis of 70 urinary glyphosate and AMPA background levels in individuals from the general German 71 72 population.

74 2. Materials and methods

75 2.1. Chemicals and reagents

76 Water, methanol and acetonitrile (all LC-MS grade) were purchased from Honeywell Research Chemicals (Neu Wulmstorf, Germany). Trifluoroethanol (TFE) and trifluoroacetic 77 78 anhydride (TFAA) (both >99%) were purchased from Merck/Sigma-Aldrich (Darmstadt, Germany). Glyphosate (98% purity), (phosphonomethyl)glycine-¹³C₂-¹⁵N ('glyphosate-79 ¹³C₂, ¹⁵N'; 99% isotopic purity; product no. G765002), AMPA (95% purity), and P-80 (aminomethyl)phosphonic acid-¹³C,¹⁵N ('AMPA-¹³C,¹⁵N'; 98% isotopic purity; product no. 81 A617342) were purchased from TRC (North York, Ontario, Canada). N-Phosphonomethyl-d₂-82 glycine (glyphosate-d₂; 99% isotopic purity; product no. D-8030) was obtained from C/D/N 83 84 Isotopes (Point-Claire, Quebec, Canada) and helium (6.0), nitrogen (5.0), argon (6.0), and methane (5.5) were obtained from Air Products (Hattingen, Germany). 85

86 2.2. Standard solutions

All standard and internal standard solutions were prepared and stored in polypropylene 87 containers. Ten calibration solutions ranging from 0.01 to 5 μ g/L in water were prepared from 88 an aqueous stock solution containing 50 µg/L each of glyphosate and AMPA. An internal 89 90 standard working solution containing 8 µg/L AMPA-¹³C,¹⁵N, as well as 4 µg/L glyphosate-¹³C₂,¹⁵N or glyphosate-d₂ (to compare these two internal standards; Figure 1), was prepared 91 in water. Aliquots of 1 mL of each calibration solution were transferred into 2 mL 92 polypropylene N9 screw cap vials (Macherey-Nagel, Düren, Germany) immediately after 93 94 preparation. All solutions were stored at -18 °C until use. For the final validated method, we used glyphosate- d_2 instead of glyphosate- ${}^{13}C_2$, ${}^{15}N_1$, contrary to Conrad et al. (2017). 95

96 **2.3. Sample preparation**

97 Sample preparation was performed as described in Conrad et al. (2017)[11] with two 98 modifications: in the initial concentration step, 2.0 mL polypropylene (PP) HPLC vials were 99 used instead of 10 mL screw-capped glass tubes, and after derivatization, the mixture was not 100 evaporated to full dryness before adding methanol. 50 µL of urine (or calibration solution) 101 and 25 µL of internal standard working solution were added to 1 mL acetonitrile in a 2.0 mL PP vial. The solution was homogenised by vortex mixing and then evaporated to dryness at 102 0.1 mbar and 50 °C for 22 min using a vacuum centrifuge (Christ RVC 2-33 CDplus vacuum 103 centrifuge; Osterode, Germany). Afterwards, 0.5 mL TFE and 1 mL -18 °C cold TFAA were 104 added, the vial was capped, and the mixture was briefly homogenised by vortex mixing. 105 Derivatization was then performed for 1 h at 80-85 °C in a heating block thermostat (VWR, 106 107 Darmstadt, Germany). After cooling down to ambient temperature, the reaction mixture was concentrated to approx. 50 µL at 85 °C. After cooling down again, the solution was mixed 108 109 with 50 µL methanol and transferred into a GC vial with a glass micro insert.

110 **2.4. Instrumental analysis**

111 A 7890B GC coupled to a 7000A triple quadrupole-MS (both Agilent, Waldbronn, Germany) 112 was used for GC MS/MS analysis. Chromatographic separation was performed on a Zabron

112 was used for GC-MS/MS analysis. Chromatographic separation was performed on a Zebron

ZB-WAX column (30 m x 0.25 mm ID, 0.25 µm film thickness, Phenomenex, Offenbach, 113 Germany) with chromatographic conditions as reported by Conrad et al. (2017)[11]. A sample 114 volume of 1 µL was injected in pulsed splitless mode (0.5 min splitless time, 120 kPa; vent 115 116 flow 50 mL/min to split). A special deactivated 4 mm inner diameter liner with quartz wool 117 was used (Sky[®] liner; Restek, Bad Homburg, Germany). Injector and transfer line temperature were 255 and 240 °C, respectively. An initial GC oven temperature of 75 °C was kept for 118 119 0.5 min. Then, the temperature was increased to 170 °C at 20 °C/min and kept for 5 min. 120 Finally, the oven temperature was increased to 250 °C at 40 °C/min and kept for 3.5 min. The carrier gas (helium) was kept at 1.2 mL/min during the analysis and increased to 2.0 mL/min 121 after 10 min. Ionization of the analytes was performed in negative chemical ionization mode 122 123 (NCI) with methane as CI gas and an ionization energy of 240 eV. The ion source was kept at 150 °C. Argon was used as the collision gas. Analyte retention times and further MS 124 conditions are shown in Table 1. MassHunter GC/MS Acquisition software (B.05.02) was 125 126 used for instrument control, and MassHunter Workstation Software Quantitative Analysis 127 (B.07.00) was used for quantitative data analysis (Agilent, Waldbronn, Germany).

128 **2.6.** Calibration, validation and quality control

Calibration was performed by linear regression (ratio of analyte and internal standard peak 129 130 area vs concentration). Intraday and interday reproducibility were studied by repeated analysis of two different pooled urine samples ('Qlow' and 'Qhigh'), containing native concentrations of 131 glyphosate and AMPA. This material was also used as quality control material for routine 132 133 analysis. For the evaluation of the method accuracy, eight different urine samples (0.4 - 2.6)134 g/L creatinine) were analysed, spiked at two concentrations (0.2 and 1.0 μ g/L), as well as without spiking. Relative recoveries were calculated for glyphosate and AMPA after 135 subtraction of their respective native concentrations. 136

2.7. Study subjects

To demonstrate the applicability of the method, 41 convenience spot urine samples from the German general population (29 females, 12 males; aged 23-61 years), collected by IPA in 2016 and 2017, were analysed. Of the participants, 31 were non-smokers. Creatinine levels ranged from 0.12 to 2.31 g/L. No information on pesticide use was available on this sample population. The study protocol was registered with the Ethics Commission of the Faculty of Medicine of the Ruhr-University Bochum, Germany (Registry Numbers: 3867-10).

144 **3. Results and discussion**

145 **3.1. Method advancement**

A LOQ for both glyphosate and AMPA of 0.05 μ g/L was achieved, an improvement of a factor of 2 compared to the original method of Conrad et al. (2017)[11]. Demonstrative GC-MS/MS chromatograms for both glyphosate and AMPA are shown in *Figure S1*, *supplemental material*. The method's precision and accuracy data are presented in **Table 2** and **Table 3**. For the intra- and inter-day precision, relative standard deviations (RSD) of <6% and <11%, respectively, were achieved.

The LOQ was determined based on a signal-to-noise ratio of 10 in a matrix (as previously described in Bury et al. 2019 [24]), in accordance with the guidelines of the working group "Analyses in Biological Materials" of the Permanent Senate Commission for the Investigation of Health Hazards of Chemical Compounds in the Work Area of the German Research Foundation [25].

157 Method accuracy, calculated as relative recoveries from native, spiked urine samples at two 158 concentration levels (0.2 and 1.0 μ g/L) was on average 109 and 115% for glyphosate and 104 159 and 116% for AMPA, with single values in the range of 83-124% and 91-123%, respectively. 160 Recoveries were not matrix-dependent (creatinine content of urine samples).

During sample work-up, two modifications were essential to improve method sensitivity and 161 162 ruggedness while keeping reproducibility and accuracy high. Glyphosate has been shown to adsorb onto glassware [26, 27]. We also made similar observations (data not shown), when 163 comparing the original method with the current one. Exchanging non-silanised glass vials for 164 165 PP vials improved the LOQ by a factor of 2. This improvement was also observed for AMPA. Consequently, we replaced glassware with polypropylene (PP) material where 166 possible/appropriate and saw considerable improvements after replacement. Additional to 167 sample work-up, the use of glassware should also be avoided during preparation and storage 168 169 of the stock and working solutions.

- 170 Secondly, but of equal importance, was the choice of the standard. Glyphosate-d₂ was used as
- an internal standard, replacing glyphosate- ${}^{13}C_2$, ${}^{15}N$ used by Conrad et al. 2017 and others.
- 172 Initially, glyphosate- ${}^{13}C_2$, ${}^{15}N$ looks more appropriate than glyphosate- d_2 , considering the shift
- in molecular mass of +3 u instead of +2 u. However, the formation of the quantifier precursor ion via dissociative electron capture is accompanied by the elimination of a radical containing
- the two glycine carbons of glyphosate, and thus the ${}^{13}C_2$ label in case of glyphosate- ${}^{13}C_2$, ${}^{15}N$,
- resulting in a mass shift of only +1 u (**Figure 2**). Naturally occurring glyphosate- ${}^{13}C_1$ yields a
- precursor ion of the same nominal mass and both glyphosate- ${}^{13}C_2$, ${}^{15}N$ and glyphosate- ${}^{13}C_1$
- form the same quantifier fragment ion at m/z 245, consequently resulting in an interference on
- this mass transition for the internal standard caused by glyphosate itself. In the case of the glyphosate- d_2 used in the final method, the mass of the precursor ion is shifted by +2 u,
- 181 eliminating this interference (see Figure S2).

The German External Quality Assessment Scheme for analyses in Biological Materials (G-182 EQUAS) (http://www.g-equas.de) currently includes only glyphosate but not AMPA. With 183 the presented method we received certification of successful participation in the two most 184 185 recent G-EOUAS rounds for glyphosate (G-EOUAS 64 and 65). Additionally, with this method we successfully participated in all three rounds of the 186 HBM4EU (http://www.hbm4eu.eu) interlaboratory comparison investigations of selected pesticide 187 biomarkers in human urine for both glyphosate and AMPA. 188

189 **3.2. Biomonitoring results**

The applicability of this method was demonstrated through the analysis of 41 spot urine samples from the general German population, collected in 2016 and 2017. Glyphosate was

detected at concentrations at or above the LOQ in 66% (n=27) of the samples, with a median

of 0.09 µg/L and a maximum concentration of 0.33 µg/L. For AMPA, 90% (n=37) of the 193 samples were above the LOQ, with a median of 0.20 μ g/L and a maximum of 2.54 μ g/L. In 194 principle, these concentrations are comparable to levels found in a previous environmental 195 196 study of the German population (Conrad et al. 2017)[11]. However, due to the improved 197 sensitivity, the range of quantifiable data now is extended, allowing us to derive more 198 informative median concentrations, for glyphosate and AMPA (for glyphosate, using a 199 method with a LOQ of 0.1 µg/L would have resulted in only 24% (n=10) samples above the 200 LOQ, making this dataset unsuitable for calculating actual median concentrations). Having a higher fraction of measured values should assist with further elucidating sources of exposures. 201

202 4. Conclusions

Method modifications resulted in a 50% lower limit of quantification for both glyphosate and 203 AMPA (0.05 µg/L) [11] without impacting performance or quality. This increase in 204 205 sensitivity is essential for the analysis of environmental exposures of populations, to have 206 more values above the LOO and thus for deriving more informative median levels in human 207 biomonitoring studies in Europe. Glyphosate and AMPA have been selected as priority substances within the EU-funded European Joint Programme HBM4EU illustrating the need 208 for sensitive and reliable analytical methods for both substances. This method successfully 209 210 passed all three rounds of the HBM4EU interlaboratory comparison investigations for both glyphosate 211 and AMPA.

212 Declaration of Competing Interest

- 213 None.
- 214

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326 **Tables and Figures:**

327 Tables:

Table 1: Further MS parameters of the final method. Analyte retention times (t_R) and precursor and product ions with corresponding collision energies are given. Qualifier transitions in parentheses (no appropriate qualifier available for AMPA-¹³C-¹⁵N).

331

| Analyte | t _R [min] | Precursor ion [<i>m</i> /z] | Product ion [<i>m</i> /z] | Collision energy [eV] |
|--|----------------------|---------------------------------|-------------------------------|--------------------------|
| Glyphosate | 6.8 | 370 (351) | 245 (268) | 10 (5) |
| Glyphosate-d ₂ | 6.8 | 372 (353) | 245 (268) | 10 (5) |
| AMPA | 7.9 | 351 (271) | 268 (188) | 5 (5) |
| AMPA- ¹³ C- ¹⁵ N | 7.9 | 353 | 270 | 5 |

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333 **Table 2:** Method precision data for glyphosate and AMPA. Quality control materials (Q_{low}

and Q_{high}) were prepared from pooled urine samples containing native concentrations of glyphosate and AMPA.

| | Glyphosate | | | AMPA |
|-----------------------------|------------------|------------|------------------|------------|
| | Q _{low} | Q_{high} | Q _{low} | Q_{high} |
| Intra-day (n=10) | | | | |
| Measured conc. (μ g/L) | 0.19 | 0.85 | 0.38 | 0.90 |
| RSD (%) | 5.2 | 4.8 | 2.6 | 3.1 |
| Inter-day (n=10) | | | | |
| Measured conc. (μ g/L) | 0.21 | 0.83 | 0.42 | 0.90 |
| RSD (%) | 10.4 | 4.9 | 7.5 | 6.9 |

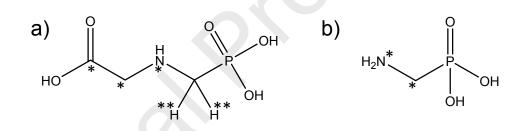
Table 3: Accuracy of the method for glyphosate and AMPA. Calculated as relative recoveries

from eight different urine samples varying in creatinine concentration (0.4 - 2.6 g/L),

analysed without spiking and spiked with 0.2 μ g/L and 1.0 μ g/L.

| | Glyp | hosate | AMPA | | |
|-------------------------------|--|----------------|--------------------|----------------|--|
| Spiked concentration (µg/L) | 0.2 | 1.0 | 0.2 | 1.0 | |
| Mean native concentration | | | | | |
| measured (range) (µg/L) | 0.11 (<loq -="" 0.40)<="" td=""><td colspan="3">1.14 (0.15 - 3.40)</td></loq> | | 1.14 (0.15 - 3.40) | | |
| Mean concentration | X | · · · | × | , | |
| measured (\sum native and | 0.32 | 1.26 | 1.34 | 2.3 | |
| spiked; range) (µg/L) | (0.22 - 0.64) | (1.13 - 1.56) | (0.35 - 3.61) | (1.34 - 4.63) | |
| Spiked conc. calculated (µg/L |) | | | | |
| Mean | 0.22 | 1.15 | 0.21 | 1.16 | |
| Range | 0.17 - 0.24 | 0.99 - 1.24 | 0.18 - 0.24 | 0.94 - 1.23 | |
| RSD (%) | 12.2 | 7.2 | 10.2 | 8.3 | |
| Accuracy (range) (%) | 109 (83 – 122) | 115 (99 – 124) | 104 (91 - 120) | 116 (94 – 123) | |

Figures:



- **Figure 1:** Chemical structures of (a) glyphosate, glyphosate- ${}^{13}C_2$, ${}^{15}N$ (isotopic labels marked with one asterisk), and glyphosate- d_2 (isotopic labels marked with two asterisks), as well as (b) AMPA and AMPA- ^{13}C , ^{15}N (isotopic labels marked with one asterisk).

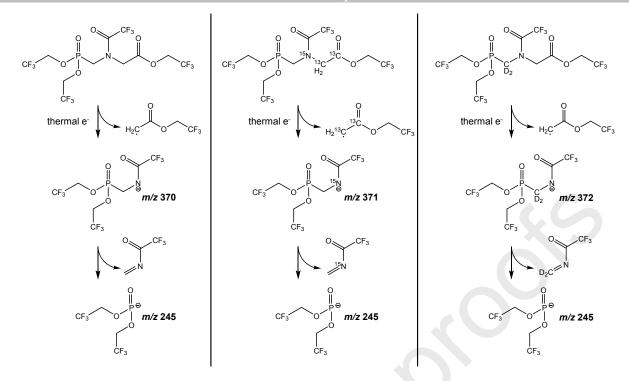


Figure 2: Quantifier mass transitions of glyphosate (left), glyphosate- ${}^{13}C_2$, ${}^{15}N$ (middle) and glyphosate- d_2 (right). Ionization of the internal standards by dissociative electron capture results in the formation of precursor ions differing from non-labelled glyphosate by +1 u (glyphosate- ${}^{13}C_2$ - ${}^{15}N$) and +2 u (glyphosate- d_2), respectively.

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358 Supplementary Data

- 359 Sensitive and selective quantification of glyphosate and
- aminomethylphosphonic acid (AMPA) in urine of the general population by
- 361 gas chromatography-tandem mass spectrometry
- 362 Alison Connolly^{a,b*}, Stephan Koslitz^a, Daniel Bury^a, Thomas Brüning^a, André
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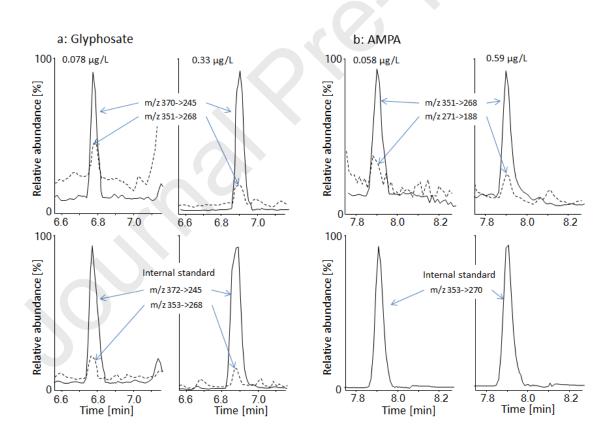
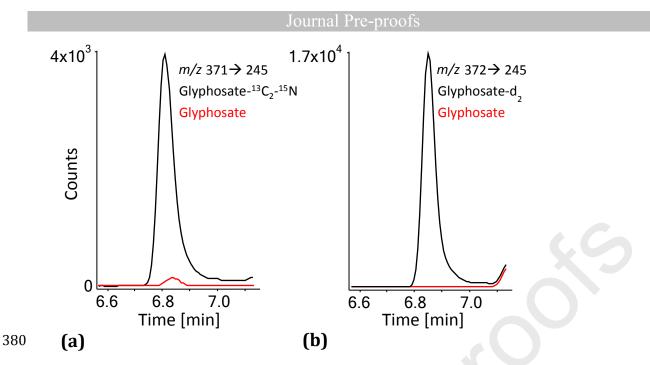


Figure S1: Chromatograms of native urine samples showing a) glyphosate and b) AMPA.
The upper row shows the target analytes with respective concentrations, the lower row shows
the respective internal standards; quantifier transitions as continuous lines and qualifier
transitions in dotted lines.



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Figure S2: Interference for glyphosate- ${}^{13}C_2$, ${}^{15}N$ caused by glyphosate. Extracted ion chromatograms for the quantifier mass transitions of (a) glyphosate- ${}^{13}C_2$ - ${}^{15}N$ and (b) glyphosate- d_2 in urine. Chromatograms of 2.5 µg/L glyphosate (red; without internal standard) showing the interference on the (a) glyphosate- ${}^{13}C_2$ - ${}^{15}N$ quantifier mass trace in comparison to 4 µg/L (i.e., the concentration used in the method) of glyphosate- ${}^{13}C_2$, ${}^{15}N$ (black). No significant interference was observed for (b) glyphosate- d_2 .