

Analytical Methods

Volume 15
Number 2
14 January 2023
Pages 135–248

rsc.li/methods



ISSN 1759-9679

 ROYAL SOCIETY
OF CHEMISTRY

PAPER

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**Indexed in
Medline!**

Cite this: *Anal. Methods*, 2023, 15, 142

Verification of exposure to chemical warfare agents through analysis of persistent biomarkers in plants†

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The continuing threats of military conflicts and terrorism may involve the misuse of chemical weapons. The present study aims to use environmental samples to find evidence of the release of such agents at an incident scene. A novel approach was developed for identifying protein adducts in plants. Basil (*Ocimum basilicum*), bay laurel leaf (*Laurus nobilis*) and stinging nettle (*Urtica dioica*) were exposed to 2.5 to 150 mg m⁻³ sulfur mustard, 2.5 to 250 mg m⁻³ sarin, and 0.5 to 25 g m⁻³ chlorine gas. The vapors of the selected chemicals were generated under controlled conditions in a dedicated set-up. After sample preparation and digestion, the samples were analyzed by liquid chromatography tandem mass spectrometry (LC-MS/MS) and liquid chromatography high resolution tandem mass spectrometry (LC-HRMS/MS), respectively. In the case of chlorine exposure, it was found that 3-chloro- and 3,5-dichlorotyrosine adducts were formed. As a result of sarin exposure, the *o*-isopropyl methylphosphonic acid adduct to tyrosine could be analyzed, and after sulfur mustard exposure the N1- and N3-HETE-histidine adducts were identified. The lowest vapor exposure levels for which these plant adducts could be detected, were 2.5 mg m⁻³ for sarin, 50 mg m⁻³ for chlorine and 12.5 mg m⁻³ for sulfur mustard. Additionally, protein adducts following a liquid exposure of only 2 nmol Novichock A-234, 0.4 nmol sarin and 0.2 nmol sulfur mustard could still be observed. For both vapor and liquid exposure, the amount of adduct formed increased with the level of exposure. In all cases synthetic reference standards were used for unambiguous identification. The window of opportunity for investigation of agent exposure through the analysis of plant material was found to be remarkably long. Even three months after the actual exposure, the biomarkers could still be detected in the living plants, as well as in dried leaves. An important benefit of the current method is that a relatively simple and generic sample work-up procedure can be applied for all agents studied. In conclusion, the presented work clearly demonstrates the possibility of analyzing chemical warfare agent biomarkers in plants, which is useful for forensic reconstructions, including the investigation into alleged use in conflict areas.

Received 11th October 2022
Accepted 27th November 2022

DOI: 10.1039/d2ay01650h

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1. Introduction

The importance of the Chemical Weapons Convention is highlighted by recent concerns of potential chemical attacks in Ukraine.¹ Over the last ten years, violations of the convention have taken place in the ongoing conflict in the Syrian Arab Republic, where the Organisation for the Prohibition of Chemical Weapons (OPCW) verified multiple attacks with sulfur mustard, sarin, and the toxic dual-use chemical

chlorine.²⁻⁶ More recently, Alexei Navalny, Sergei Skripal and four other individuals were poisoned by different nerve agents belonging to the group of Novichocks.^{7,8} These chemical warfare agents (CWAs) exert the same mechanism of action as other organophosphate nerve agents, although they are expected to be even more potent.^{9,10} These incidents emphasize that there is a continued risk of the use of CWAs in both large-scale attacks as well as small scale deployment. Because of the high impact of these events, often extensive international investigations are being conducted to verify the use of chemical weapons.

Analytical methods for the detection of chemical threat agents in matrices such as organic solvents, air samples,¹¹⁻¹³ soil,^{14,15} wipe^{16,17} and other solid materials have been broadly investigated.¹⁸ However, detection of intact agents is often not possible due to the relatively high volatility and reactivity of these chemicals.¹⁹ In such cases, biomedical samples may provide evidence for a longer period of time, where persistent

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† Electronic supplementary information (ESI) available. See DOI: <https://doi.org/10.1039/d2ay01650h>



biomarkers can be used for verification.^{20,21} Nonetheless, it may be problematic to get access to human biological samples because of ethical, cultural and safety reasons. In addition, it might be difficult to link the victims to the actual exposure area. In contrast, environmental samples such as plant leaves are abundant, easy to collect and transport, and not subject to informed consent or ethic regulations as is the case with human biomedical samples. The use of plant evidence has been scarcely examined for the presence of the intact CWAs or associated degradation products.^{22–26} In addition, a news article reported that researchers at Lawrence Livermore National Laboratory identified small chlorine biomarkers and protein adducts in grass.²⁷ However, to our knowledge, no research article has been published that specifically targets plant protein adducts to demonstrate CWA exposure. We hypothesize that combining the analysis of environmental samples with methods applied to biomedical samples could provide powerful new options to investigate CWA exposure at an incident site long after the alleged release of the agent.

Most CWAs and toxic industrial chemicals can form covalent, long-lasting adducts with proteins.²⁸ This principle has formed the basis for analyzing biomedical samples for verification of exposure to such chemicals. Previous studies have shown that in case of human exposure to sulfur mustard, histidine protein adducts are formed.²⁹ After exposure to nerve agents, such as sarin and Novichok A-234, phosphorylated serines or tyrosines are formed as specific biomarkers.^{30–32} Likewise, mono- and dichlorinated tyrosine adducts are the major protein markers for the presence of chlorine.^{33,34}

Also, plants also contain a lot of proteins, with ribulose-1,5-bisphosphate carboxylase-oxygenase (rubisco) being the most abundant protein on earth.³⁵ As a result, plant proteins could serve as potential scavengers for chemical threat agents and might form protein adducts in a similar way as unraveled for humans. Abundantly occurring plants, representing a wide range of vegetation, are bay laurel leaf (*Laurus nobilis*), basil (*Ocimum basilicum*) and stinging nettle (*Urtica dioica*), depicted in Fig. 1. Stinging nettle is a rapidly growing weed with a high protein content and is originally native to Europe.^{36,37} In contrast, basil thrives in tropical and subtropical climates and is an aromatic herb and medicinal plant.³⁸ Bay laurel leaf is used as a herb in many traditional practices and recipes, originating from the Mediterranean region, Asia, South and North America, and the Balkans.³⁹ It is expected that these species will be sufficiently representative to study CWA biomarker occurrence in plants in general. This will allow the use of the

presented methodology for any type of vegetation encountered on the incident scene.

Consequently, the aim of the current study was to explore whether persistent biomarkers of various CWAs could be detected in plants after vapor and liquid exposure. A novel approach was developed for analyzing modified protein adducts in vegetation, a strategy that to our knowledge has not been reported in literature before. The model plants used in this study were exposed to sulfur mustard, sarin, chlorine or Novichok nerve agent A-234. After exposure, leaves were collected, cut into small pieces, and washed to remove excess of intact CWA. The plant material was dried, then digested by pronase or trypsin followed by analysis with liquid chromatography tandem mass spectrometry (LC-MS/MS) and liquid chromatography-high resolution tandem mass spectrometry (LC-HRMS/MS). The present work demonstrates for the first time that plant biomarkers are promising indicators of CWA exposure that can be used for forensic reconstruction long after the alleged agent release.

2. Experimental

2.1. Safety

The organophosphate nerve agents Novichok A-234 and sarin (GB) and the blister agent sulfur mustard (HD) are highly toxic chemicals. All experiments were performed by trained personnel in the dedicated High-Tox facility at TNO Rijswijk, which is allowed under the Chemical Weapons Convention to synthesize and handle chemical warfare agents. The experiments, including chlorine gas exposure, were performed in a fume hood by trained personnel. The chemicals were handled in a leak-tight containment and personal protection measures, such as lab coats, gloves and safety glasses were worn. In case of an accidental exposure, an autoinjector with atropine sulphate and obidoxime as well as a diazepam preparation were readily available for administration by a specialized first aid team, to reduce the symptoms of intoxication.

2.2. Chemicals and materials

The chemicals A-234, GB, HD, *o*-isopropyl methylphosphonic acid adduct to tyrosine (GB-Tyr), N1-HETE-Histidine (N1-HETE-His), N3-HETE-Histidine (N3-HETE-His), A-234-Tyr and methylphosphonic acid (MPA)-Tyr were synthesized in the High-Tox facility at TNO, Rijswijk. The compounds were characterized by NMR, GC-MS and LC-MS. Purities of the synthesized agents, except for A-234-Tyr and MPA-Tyr which were only used for



Fig. 1 Photo of leaves of selected plants. (A) Bay laurel leaf (*Laurus nobilis*), (B) Basil (*Ocimum basilicum*), (C) Stinging nettle (*Urtica dioica*).



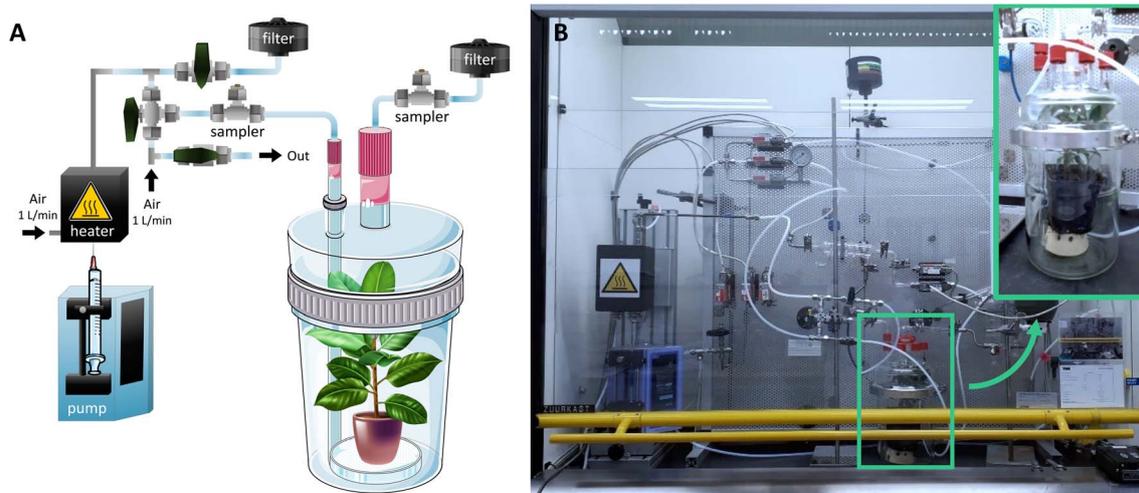


Fig. 2 Dedicated vapor generation set-up for controlled exposure of plants to chemical warfare agents. (A) Schematic view. (B) Photo of vapor generation set-up.

qualitative analysis, exceeded 95% (determined by NMR). 3,5-Di-chloro-L-tyrosine (di-Cl-Tyr) was purchased from BOC Sciences (London, UK) and $^{13}\text{C}_6$ -3-chloro-L-tyrosine was obtained from Cambridge Isotopic Laboratories (Andover MA, USA). Purities of the agents exceeded 97%. Acetic acid, ammonium bicarbonate (ABC), calcium hypochlorite, 3-chloro-L-tyrosine, dithiothreitol (DTT), formic acid, protease from *Streptomyces griseus* (pronase, ≥ 3.5 units per mg solid), sodium acetate trihydrate, sodium iodoacetate, trypsin from bovine pancreas ($\geq 10\,000$ BAEE units per mg protein) and urea were obtained from Sigma-Aldrich (Zwijndrecht, The Netherlands). Acetonitrile (ACN), acetone, ethanol and methanol (MeOH) were purchased from Biosolve (Valkenswaard, The Netherlands). Hydrochloric acid (HCl) was obtained from Thermo-Fisher Scientific (Landsmeer, The Netherlands). Additionally, MilliQ water (SimPak® 1) was used. Laurel (*Prunus Lusitanica Angustifolia*, Intratuin), *Ocimum Basilicum* (Aldi) and Nettle (local plant, Rijswijk) originated from the Netherlands.

2.3. Exposure of the plants

2.3.1. Liquid exposure. Laurel, basil, and stinging nettle were exposed to various concentrations of Novichok A-234, sarin or sulfur mustard. First, a single leaf was exposed in duplicate to a blank of ACN or 100 μL (in small drops of approximately 5 μL) of 0.02, 0.4 and 4 mmol L^{-1} of A-234, 0.004, 0.05 and 0.1 mmol L^{-1} of sarin or 0.02, 0.1, 1 and 10 mmol L^{-1} of sulfur mustard dissolved in ACN. The leaves had a total dry mass of about 30 mg and were left for 24 hours in a closed Petri dish. Afterwards, the samples were prepared for analysis as described in Section 2.4.

2.3.2. Vapor and gas exposure. Additionally, the plants were placed in a 2 L glass vessel and exposed to vapors of sarin or sulfur mustard in a dedicated set-up (Fig. 2). First, vapors at a concentration of 2, 5, 25 or 250 mg m^{-3} sarin or 2.5, 12.5, 50 or 150 mg m^{-3} sulfur mustard were generated. The temperature was maintained at 20–22 $^{\circ}\text{C}$. A syringe pump (Hamilton 1801N,

10–100 μL) delivered the agent at constant flow rate (10–250 nL min^{-1}) in a dry air stream (1 L min^{-1}) in a heated evaporation chamber at 140 $^{\circ}\text{C}$. The incoming vapor flow entered the vessel at the bottom, while the outgoing flow exited at the top to promote a homogenous distribution. The exiting flow was led through a charcoal filter to trap and convert the threat agents. After a 30 minute exposure, the valves could be switched without opening the setup to clean the vessel by air flow for approximately 30 minutes. This was verified at the sampling spots with a handheld CWA detector (LCD 3.3, Smiths Detection).

For chlorine exposure, the plants were placed in a closed 2 or 20 L vessel and chlorine was generated by the reaction of calcium hypochlorite and HCl, similar to the method applied by de Bruin-Hoegée *et al.*²¹ A 1.5 mL Eppendorf tube with 1, 10 and 50 mg calcium hypochlorite was placed in the vessel above the plant. A syringe was put through a septum and slowly, over a period of one minute, 1 mL of a 12 M HCl solution was added to the calcium hypochlorite, after which 50 mg m^{-3} , 500 mg m^{-3} , 5 g m^{-3} or 25 g m^{-3} chlorine gas was generated. The plants were left for 30 minutes in the closed vessel.

2.4. Protein precipitation and digestion

Fig. 3 summarizes the sample preparation steps. After exposure, two leaves of each plant species were collected and washed with methanol to remove potential intact CWAs. Also, two unexposed leaves were included. Subsequently, the leaf was placed in a 2 mL vial and cut into minuscule homogenous pieces using a small nail scissors. The sample was then transferred into a 15 mL centrifuge tube and washed with 10 mL acetone by centrifuging at 4000 rpm for 10 minutes (Megafuge 1.0 R centrifuge). The washing step was included to dehydrate and clean the sample, because the aim of the study was to find protein adducts. It should be noted that potential other, small molecule biomarkers could have been removed during this procedure. This was repeated three times for HD and chlorine.





Fig. 3 Schematic overview of the sample preparation steps. Figures adapted from <https://www.smart.servier.com>, CC BY 3.0.

Subsequently, the plant material was dried overnight at ambient temperature. It should be noted that drying is not required if only qualitative analysis is necessary, since digestion can be performed with wet leaves as well.

Afterwards, the samples were digested. The applied method was adapted from protocols originally described for the detection of CWAs in biomedical samples.^{4,21,40} For pronase digestion, 400 μL ABC buffer (50 mM) and 100 μL pronase (10 mg mL^{-1} in 50 mM ABC) were added to 10 mg of dried leaf. The digests after HD and chlorine exposure were incubated overnight at 37 $^{\circ}\text{C}$ and a rotational speed of 800 rpm. Subsequently, the samples were filtered through a 10 kDa Amicon ultra-centrifugal filter at 14 000 rpm for 10 minutes in an Eppendorf centrifuge. The samples after sarin exposure were incubated with pronase for 90 minutes in a Thermoshaker (grant-bio PHMT) at 50 $^{\circ}\text{C}$ and 500 rpm. It turned out that both digestion methods, for either sulfur mustard and chlorine and the procedure for sarin, provided similar results and could be used interchangeably.

The filtrates were purified with reversed phase solid phase extraction (SPE) using a C18 column (Bakerbond SPETM). To wet the sorbent bed and activate the nonpolar sorbents, 1 mL MeOH was percolated through the column and 1 mL water was used to equilibrate the column. Afterwards, the sample was loaded on the column and washed with 10% MeOH in water. The retained analytes were eluted with 1.5 mL MeOH and dried under nitrogen. The dried sample was then dissolved in 50 μL water with 0.2% v/v formic acid and analyzed as described in Section 2.5.

For trypsin digestion, 1 mL urea (8 M solution in 50 mM ABC) containing 5 μL of DTT (800 mM in water) was added to 10 mg of plant material. Subsequently, the sample was incubated for 45 minutes at 37 $^{\circ}\text{C}$ and 800 rpm. After the incubation, 100 μL sodium iodoacetate (150 mM in water) was added. Next, the sample was incubated for 30 minutes in the dark at 37 $^{\circ}\text{C}$ and 800 rpm. Thereafter, the sample was filtered through a 10 kDa Amicon ultra-centrifugal filter at 14 000 rpm for 10 minutes in an Eppendorf centrifuge (5417 R). The residue was washed on the filter four times with 400 μL ABC buffer (50 mM) and afterwards 400 μL water and 30 μL trypsin solution (10 mg mL^{-1} in 50 mmol L^{-1} acetate buffer, pH = 3.55) were added. The sample was incubated overnight at 37 $^{\circ}\text{C}$ and 800 rpm. After digestion, the samples were filtered through a 10 kDa filter at 14 000 rpm for 10 minutes and analyzed as described in Section 2.5.

2.5. Chemical analysis

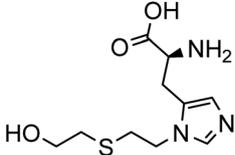
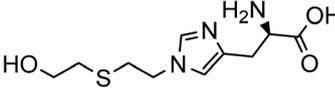
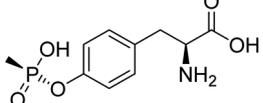
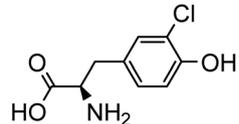
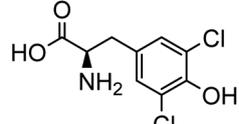
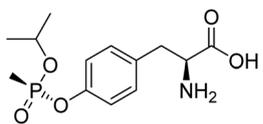
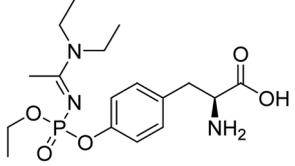
2.5.1. Targeted analysis by LC-MS/MS. Prior to analysis, the pronase digests were diluted up to 100-fold by water with 0.2% v/v formic acid depending on the exposure concentration. The samples were analyzed by a Waters Acquity ultra-high-pressure liquid chromatographic (UPLC) system equipped with a Waters Acquity HSS T3 C18 column (100 \times 2.1 mm I.D., 1.8 μm). Samples were kept in the autosampler at 8 $^{\circ}\text{C}$ and a volume of 5 μL was injected, after which the analysis was performed at room temperature with a gradient flow of 100 $\mu\text{L min}^{-1}$. All compounds could be analyzed with a single chromatographic method. The optimized gradient elution settings can be found in Table A1 of the ESI.[†] The UPLC system was coupled to a Waters (Milford, MA, USA) Xevo TQ-S triple-quadrupole mass spectrometer, equipped with electrospray ionization. Analytes were quantified in positive ionization mode with a capillary voltage of 3.5 kV. The nitrogen cone gas flow was set to 150 L h^{-1} and the argon collision gas flow was set to 0.19 mL min^{-1} . Cone voltages of 40 V for A-234 and GB, 10 V for chlorine and 14 V for HD exposure were used. Data was acquired in selected reaction monitoring (SRM) mode.

The collision energy was optimized for each compound. Table 1 shows the chromatographic and mass spectrometric parameters for the protein adducts and metabolites which were analyzed by LC-MS/MS. The identification of the adducts was confirmed by comparing their retention times, precursor ion and characteristic fragment ion m/z values with synthetic reference standards. GB-Tyr, Cl-Tyr, di-Cl-Tyr, N1-HETE-His, N3-HETE-His and ¹³C₆-Cl-Tyr were quantitatively analyzed, while A-234-Tyr and MPA-Tyr were only utilized for qualitative analysis. For the limit of detection (LOD) a signal-to-noise (S/N) ratio of at least 5 was required. Further results of the optimization and validation of the corresponding targeted LC-MS/MS method can be found in Section S1 of the ESI.[†]

2.5.2. Data-dependent acquisition by LC-HRMS/MS. The trypsin digests were analyzed on an LC-HRMS/MS instrument based on a previously published method.²¹ An Ultimate 3000 RSLCnano system (Thermo Scientific Dionex Softron GmbH, Germany) coupled to an Orbitrap mass spectrometer (Q Exactive plus, Thermo Scientific, Bremen, Germany) was used. A volume of 10 μL was injected onto an Acclaim PepMap 100 C18 μ -pre-column (5 mm \times 300 μm I.D., 5 μm , 100 \AA , Thermo Fisher Scientific) and subsequently separated on an Acclaim PepMap



Table 1 Chromatographic and mass spectrometric parameters for analytes and internal standards analyzed by LC-MS/MS

Analyte	Precursor ion (m/z)	Retention time (min)	Production (m/z)	Collision energy (eV)	Linearity range (ng mL^{-1})	Chemical structure
N1-HETE-His	260.0	3.7	105.0	11	1–100	
			81.8	25		
			61.0	25		
N3-HETE-His	260.0	4.2	105.0	11	1–100	
			81.8	25		
			61.0	25		
MPA-Tyr	260.1	5.1	214.1	13	n.a.	
			197.1	13		
$^{13}\text{C}_6\text{-Cl-Tyr}$	222.0	6.3	176.3	15	1–100	
Cl-Tyr	216.2	6.3	141.3	25	1–100	
			170.3	15		
			135.3	25		
di-Cl-Tyr	250.1	6.7	204.0	15	1–100	
			169.0	30		
GB-Tyr	302.1	8.1	260.1	13	0.05–20	
			214.1	13		
A-234-Tyr	386.2	9.3	313.0	15	n.a.	
			285.0	15		
			244.1	15		
			74.1	15		
				15		

C18 Analytical column (250 mm \times 75 μm I.D., 2 μm , 100 \AA , Thermo Fisher Scientific). Afterwards, the raw data was analyzed using Peaks® X+ software (2019, Bioinformatics Solutions Inc., Waterloo, Canada) to identify the peptides that were present in the sample. A peaks database search was employed which compared all theoretically possible peptides with the FASTA database. Also, a *de novo* search was applied. The error tolerance for the precursor mass and the fragment ion were set at 10 Da and 0.5 Da monoisotopic mass, respectively.

3. Results & discussion

3.1. Visual examination of vegetation

Any discoloration or other physical changes of the leaves were monitored to assess potential visual cues for plant and leaf collection in the field. After chlorine exposure, typical yellow staining was immediately observed (Fig. 4B). Alternatively, leaves exposed to sulfur mustard (Fig. 4D), A-234 or sarin

showed discoloration, turning the leaf from green to a brown color. However, this was only visible for relatively high vapor concentrations, whereas lower concentrations ($<25 \text{ mg m}^{-3}$) did not result in immediate physical changes. More pictures are included in Section S2 of the ESI.† Further knowledge of the physical state of the plant might be obtained by applying fluorescence detection before sample selection.⁴¹

3.2. Liquid exposure

Protein adducts were formed in plants after exposure to various chemical threat agents. To the best of our knowledge, this is the first time that these biomarkers are detected in plants. After liquid exposure to A-234, phosphorylated tyrosine adducts were found. Fig. 5 shows representative extracted ion chromatograms of the sample preparation blank, a standard of tyrosine exposed to A-234, and nettle exposed to A-234. The retention time (t_R) and fragmentation pattern of the unknown sample correspond to the standard. No peaks were visible in the blank



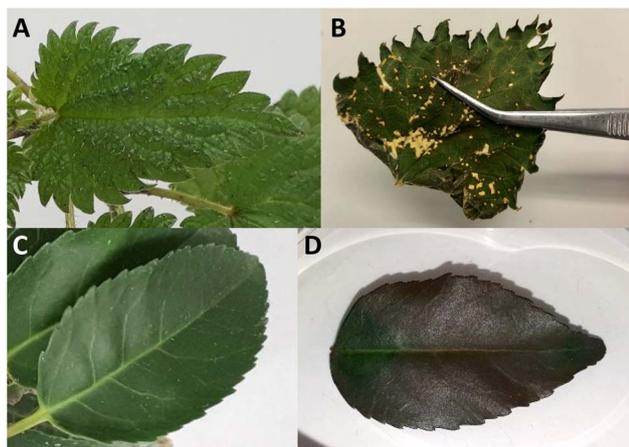


Fig. 4 Physical state of nettle leaf upon chlorine gas exposure and laurel upon sulfur mustard exposure. (A) Nettle leaf before exposure, (B) Nettle leaf after 25 g m^{-3} chlorine exposure, (C) Laurel before exposure, (D) Laurel leaf after $20 \mu\text{mol}$ liquid sulfur mustard exposure.

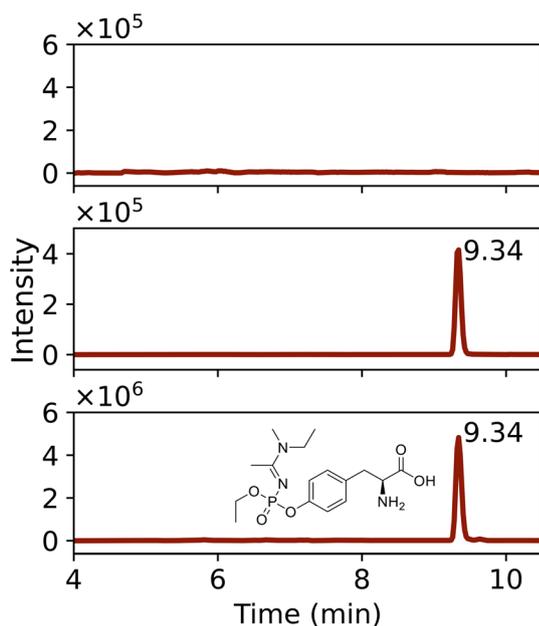


Fig. 5 Extracted ion chromatograms (m/z 386.2 \rightarrow 74.1) of phosphorylated tyrosine adduct with $t_R = 9.34$ min, after liquid exposure to A-234. (A) Sample preparation blank. (B) Standard A-234-Tyr. (C) A-234-Tyr adducts measured in nettle plant after exposure to 40 nmol A-234.

runs even after a 10-fold expansion of the base line signal. In Fig. 6 the HRMS/MS spectrum of A-234-Tyr is presented with a mass error of only 0.22 ppm compared to the theoretical m/z value. The cleavage products with m/z 313.0949, 285.0635, 244.0734 and 74.0964 were clearly visible, which were only reported by one other study for the A-234 tyrosine adduct in biological fluids.⁴² This is the first study to identify the A-234 tyrosine adduct by comparing it to a synthetic reference standard. The lowest m/z of the adduct moiety diethylamine was observed in other studies for the nonapeptide fragmentation

pattern as well.³² An increasing exposure concentration resulted in higher adduct formation in leaves exposed to liquid drops (Fig. 7A), with a limit of detection of 2 nmol (448 ng).

Furthermore, the *o*-isopropyl methylphosphonic acid adduct to tyrosine was also detected in vegetation after liquid exposure to sarin, confirmed by a reference standard. The MPA-Tyr adduct was not detected, indicating no aging of the adduct. A liquid exposure of 0.4 nmol sarin (56 ng) could still be observed in nettle leaves. When the exposure level was increased, a higher concentration of biomarkers was measured (Fig. 7B).

Third, the N1- and N3-HETE-His adducts were detected after liquid exposure of leaves to sulfur mustard. An exposure of a single nettle leaf to 0.2 nmol HD (32 ng) could still be detected. Both adducts were formed to a similar extent at each sulfur mustard concentration (Fig. 7C). The next section will describe the results of the exposure of plants to GB and HD vapor including the chromatograms.

3.3. Vapor and gas exposure

3.3.1. Analysis of phosphorylated tyrosine adducts after sarin exposure. Similar to the liquid exposure results, the *o*-isopropyl methylphosphonic acid adduct to tyrosine was also detected in living plants after vapor exposure to sarin as confirmed by a reference standard (Fig. 8). The same product was observed for all investigated plant species. The lowest observable exposure limit in basil was 2.5 mg m^{-3} and in laurel and nettle 25 mg m^{-3} after full plant exposure. This concentration is higher than the lethal dose for humans of 0.19 mg m^{-3} for a 30 minute exposure.⁴³ It can be expected that in real cases, the exposure pattern of plants is completely different compared to humans, because exposed people will leave the incident scene as soon as possible, while the vegetation can be exposed for a longer duration. This will then result in a higher adduct concentration. Besides, vegetation might also be found very close to the center of the release and thus be exposed to much higher concentrations. The persistence of the protein adducts was remarkable, GB-Tyr could still be detected in living plants, as well as in dried leaves, up to three months after the actual exposure. The extracted ion chromatograms analysed three months after exposure, are presented in Section S3 of the ESI.† Dried leaves did not require storage at reduced temperatures. This would facilitate the transport and storage of plant-based evidence during investigations of alleged use of chemical weapons. It is expected that the detection window could be extended even further, because the concentration of the adduct in the leaves did not significantly drop after three months, and for a high exposure experiment the MS response still exceeded the limit of quantification by a factor of 10 000.

3.3.2. Analysis of chlorotyrosine adducts. In case of exposure to the toxic industrial chemical chlorine, the 3-chloro- and 3,5-dichlorotyrosine adducts were identified in vegetation in line with studies involving biomedical samples (Fig. 9). The lowest observable exposure limit was 500 mg m^{-3} after full plant exposure of laurel and nettle, whereas for the experiments with basil, chlorine contact could be established from a concentration of 5 g m^{-3} . The LOD was lowered when a single



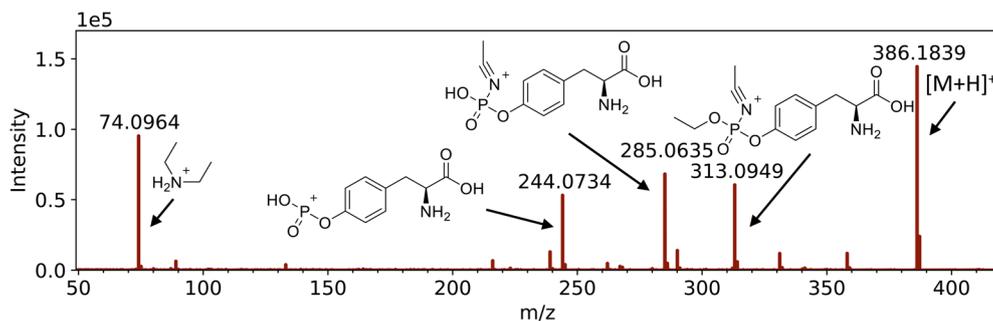


Fig. 6 HRMS/MS spectrum with fragmentation pattern of the tyrosine adduct of Novichock A-234 (m/z 386.1839).

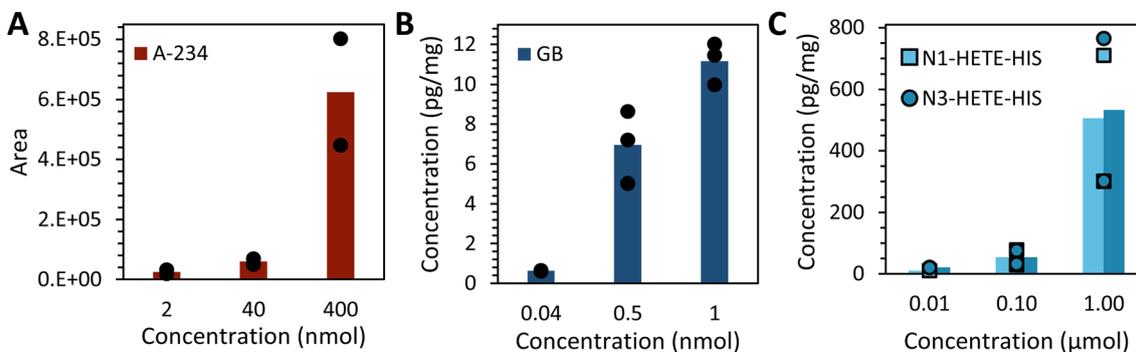


Fig. 7 Effect of concentration of added droplets on the average adduct formation in nettle plant per mg dried leaf (A) A-234, (B) GB and (C) HD exposure.

leaf was exposed to the same chlorine concentration, resulting in a LOD of 50 mg m^{-3} after exposure of nettle. This is slightly below the lethal dose for humans, since a severe 30 minute

exposure at 81 mg m^{-3} can result in cell death, pulmonary edema, or a sudden death due to narrowing of the airways.⁴⁴ Additionally, an increasing chlorine concentration resulted in

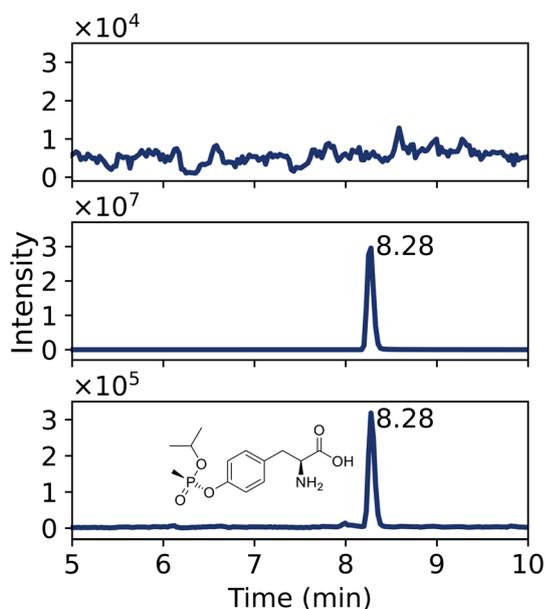


Fig. 8 Extracted ion chromatograms of phosphorylated tyrosine adduct with $t_R = 8.28$ after sarin vapor exposure. (A) Sample preparation blank. (B) Standard GB-Tyr. (C) GB-Tyr measured after plant exposure to 250 mg m^{-3} sarin.

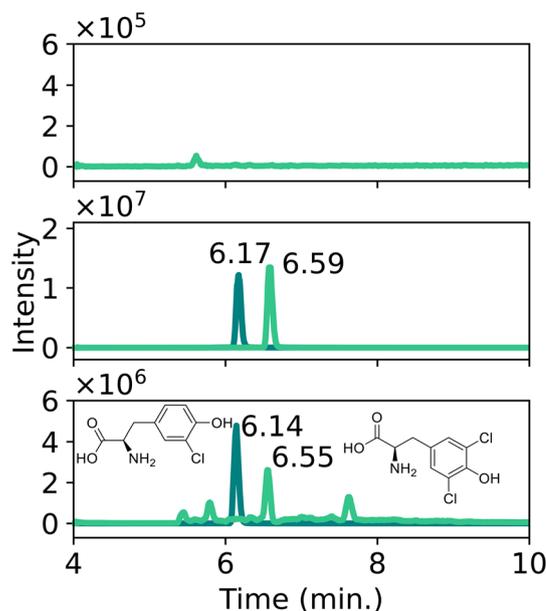


Fig. 9 Extracted ion chromatograms of chlorinated tyrosine adducts after chlorine gas exposure. (A) Sample preparation blank. (B) Standard Cl-Tyr with $t_R = 6.17$ min and di-Cl-Tyr with $t_R = 6.59$ min. (C) Cl-Tyr and di-Cl-Tyr, diluted 100 times, measured after 5 g m^{-3} plant exposure.



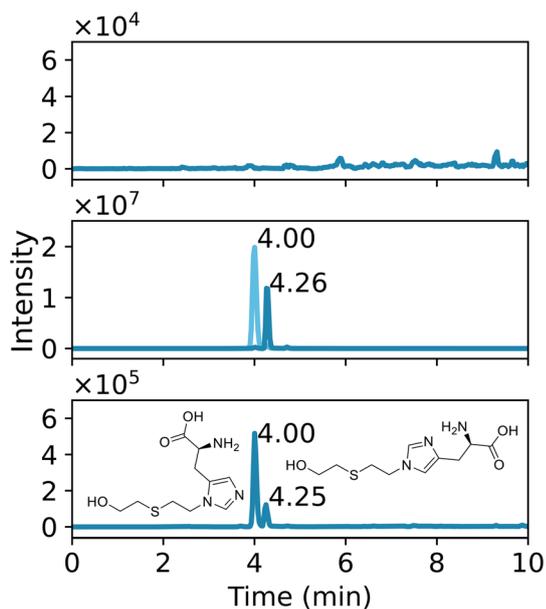


Fig. 10 Extracted ion chromatograms of N1-HETE-His and N3-HETE-His adducts after sulfur mustard vapor exposure. (A) Sample preparation blank. (B) Standard N1-HETE-His with $t_R = 4.00$ min and N3-HETE-His with $t_R = 4.26$ min. (C) N1-HETE-His and N3-HETE-His measured after 150 mg m^{-3} plant exposure.

an increased formation of di-Cl-Tyr adduct compared to Cl-Tyr. Interestingly, the observed persistence of the chlorine biomarkers was similar to the findings for the A-234 and sarin experiments. Each protein adduct could still be detected three months after the exposure experiment at a concentration significantly higher than the LLOQ (Section S3 of the ESI†).

It should be noted that these tyrosine adducts do not immediately relate to a targeted attack with chlorine gas. Similar reactions can also occur when plants are exposed to accidental chlorine exposure for instance after contact with household bleach. Further research should be conducted to investigate whether it would be possible to differentiate between various exposure scenarios and the chemicals involved through detailed analysis of plant biomarkers. Possibly this will require the analysis of chlorinated peptides as was recently demonstrated for human biological samples.³⁰ However, the context of a given case and the conditions at the incident scene might also provide insight into the nature of the chlorine source.

3.3.3. Analysis of histidine adducts after sulfur mustard exposure. Fig. 10 shows the N1- and N3-HETE-His adducts that were detected after exposure of plants to sulfur mustard. LC gradient separation was satisfactory to differentiate between these two isomers ($\Delta t_R > 0.4$ min). Both adducts could be detected in laurel at a vapor concentration as low as 12.5 mg m^{-3} . In nettle the N1-HETE-His adduct was detected at the same exposure level, while the N3-HETE-His adduct was only visible after a higher exposure of 50 mg m^{-3} . The biomarkers in basil could only be detected after an exposure of 150 mg m^{-3} . In line with the other chemical warfare agents, the biomarkers of sulfur mustard could still be observed three months after

exposure at a concentration significantly exceeding the LLOQ (Section S3 of the ESI†). Life-threatening effects in humans already occur after a 30 minute exposure to 2.7 mg m^{-3} .⁴⁵

Further improvements in limit of detection might be obtained by processing more plant material. In this study only 10 mg of dried plant material was prepared for analysis, which was often less than the mass of a single leaf. Especially after large scale attacks, it is expected that much more vegetation has come into contact with the chemical agents and could hence be collected and processed for forensic investigations. Another potential advancement in sensitivity, which has remarkably decreased the detection limit for the analysis of nerve agents in biomedical samples, might be realized by isolating specific proteins using antibodies.⁴⁶

Over time a slight increase in adduct concentration in the living plant was observed after sarin, chlorine, and sulfur mustard exposure, which might be due to accumulation in the soil or plant as is observed for free metabolites as well.²⁵ However, given the small sample size, the results must be interpreted with caution as the variation in concentration observed in individual leaves was substantial, which might be due to the position relative to the vapor flow or the growth phase of the leaf.

3.4. Influence of plant exposure conditions

The conditions of the leaves were varied to mimic different weather conditions. Freshly picked leaves were compared with wetted, frozen, and dried leaves. After chlorine exposure, the concentration of Cl-Tyr and di-Cl-Tyr in the frozen and dried leaves was significantly lower than the adduct concentration detected in fresh leaves as shown in Fig. 11. Nevertheless, as described before, it was still possible to identify protein adducts in dried leaves after three months. The di-Cl-Tyr concentration of the wetted leaves was also slightly reduced, while the Cl-Tyr concentration did not show a significant difference. Additionally, during sarin and sulfur mustard exposure, the humidity in the vapor generation setup was increased from dry air to 60% relative humidity. However, the adduct concentration was not affected by these conditions. Finally, a doubled exposure time with GB and HD resulted in more adduct formation as expected. At even higher exposure levels and durations, the biomarker formation is expected to level out when all available amino acid sites have reacted with the CWA. However, such a high exposure concentration was not applied due to safety reasons.

The observed biomarker concentration was considerably affected by the exposure area of the plant. An increase in biomarker formation with a factor of four was observed for a single leaf compared to a whole plant exposed to 5 or 25 g m^{-3} chlorine gas. This suggests that all the available chlorine fully reacts with an excess of biomaterial and that hence the biomarker formation is distributed over the various leaves of a full plant. As a result, the size of the plant and its surface area vs. mass indirectly affects the formed adducts. To improve the exposure design in future studies, it is recommended to precisely select the plant size and number of leaves to limit experimental variation. However, in real-life incidents CWA



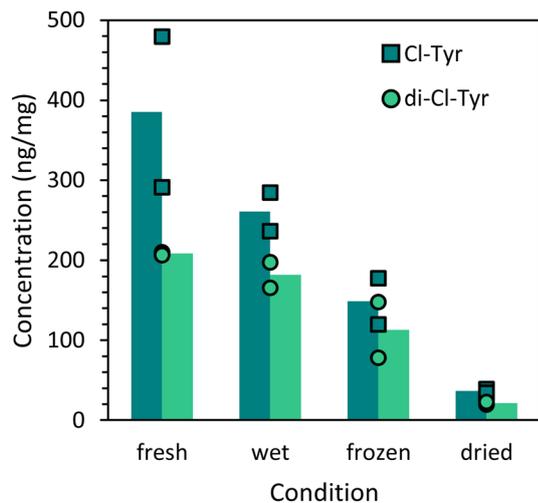


Fig. 11 Influence of nettle leaf conditions on the average adduct concentration.

exposure will be uncontrolled and will vary greatly depending on the exposure conditions and the location of the vegetation at the scene.

In this study, no consistent differences in adduct concentration were observed for the various plant species. The variation within the individual experiments of one plant species was similar to the variation between different plant species. Nevertheless, all chemical warfare agent adducts were visible in all investigated plant species. A plausible explanation for this might be that green plants in general contain a lot of similar proteins consisting of amino acids that are accessible to chemical threat agents, as will be discussed in more detail in Section 3.5. It is therefore expected that these findings can be extrapolated to other vegetation as well.

3.5. Protein identification with high resolution mass spectrometry

To provide more insight in which plant proteins react with the threat agents, peptide identification was performed by LC-HRMS/MS after trypsin digestion. It was anticipated that a majority of the adducts would originate from the abundant plant protein rubisco. This protein was also mentioned in the news article about the research of Lawrence Livermore National Laboratory.²⁷ However, the applied pronase digestion method is non-specific, and the monitored biomarkers can result from any plant protein containing tyrosine. Because of the lack of specific databases of nettle, laurel, and basil, only peptides that occur generically in plants could be identified. Consequently, this approach is generally applicable to a wider range of plant species.

Table 2 shows the identified peptides that were present in two nettle samples exposed to chlorine gas. No chlorinated peptides were found in the blank. As expected, mono and dichlorinated rubisco peptides were found. Besides this protein, various peptides from other proteins were detected, such as chlorinated chloroplastic ATP synthase, which are involved in the CO₂ signalling pathway and control gas-exchange between plants and the atmosphere,⁴⁷ chlorophyll binding protein and serine-glyoxylate aminotransferase. In one sample the peptide VGLTALTIAEY*FR was both mono chlorinated and dichlorinated.

After duplicate sulfur mustard exposure of laurel and basil, post-translational modifications were found as well (Table 2). Slightly different proteins were found in both samples of laurel and basil. The proteins calmodulin, cytochrome, photosystem II D2 and chloroplastic chlorophyll binding protein were modified. Of the latter protein, even two peptides were affected, which might indicate that this protein is an accessible target.

Additionally, Table 2 shows the phosphorylated peptides that were formed after sarin exposure. Also, in this case

Table 2 Overview of peptides containing at least one chlorinated or phosphorylated tyrosine residue or histidine adduct of sulfur mustard, detected in duplicate samples of nettle, laurel or basil after exposure

No.	Peptide	Protein	Accession	Exposure	Species
1	GYPGY(Cl)LYTDLATIYER	ATP synthase	B0K8E7 VATB_THEP3	Cl	Nettle
2	EAQAVADDVFSLFISEEVDKVELLY(Cl)TK	ATP synthase, chloroplastic	Q01908 ATPG1_ARATH	Cl	Nettle
3	VGLTALTIAEY(Cl)FR	ATP synthase, chloroplastic	O03081 ATPB_PSINU	Cl	Nettle
4	H*NLTDQQLSEFK	Calmodulin	Q39752 CALM_FAGSY	HD	Laurel
5	Y(Cl)AGVGAIEYAVLHLK	Carbonic anhydrase, chloroplastic	P42737 BCA2_ARATH	Cl	Nettle
6	GPLENLADHLADPVNNNAWAY(Cl)ATKLCPGK	Chlorophyll binding protein, chloroplastic	P09755 CB22_SOYBN	Cl	Nettle
7	Y*GELLHGR	Chlorophyll binding protein, chloroplastic	Q07473 CB4A_ARATH	GB	Basil
8	EVPYLH*LQFDLSDQNLAK	Chlorophyll binding protein, chloroplastic	Q9XF88 CB4B_ARATH	HD	Basil
9	EFLVLH*LQFELDSLQDGLAK	Chlorophyll binding protein, chloroplastic	Q9XF88 CB4B_ARATH	HD	Basil
10	H*HLTFPLSPDPTTK	Cytochrome	A6MMV7 CYF_ILLOL	HD	Basil
11	WSFAAEH*HPENLLFPEEVLPR	Photosystem II D2	Q2MIA5 PSBD_SOLLC	HD	Basil
12	Y*TLPGHKKAFLPR	Putative beta-D-xylosidase	P83344 XYNB_PRUPE	GB	Laurel
13	GHYLNATAATC(Cl ₂)EEMLR	RuBisCO	Q1KVV0 RBL_TETOB	Cl	Nettle
14	LYTYTPEY(Cl)ETK	RuBisCO	P48706 RBL_LACSA	Cl	Nettle
15	VAY(Cl)PIDLFEEGSVTNLFTSIVGNVFGFK	RuBisCO	P48711 RBL_PICAB	Cl	Nettle
16	WY*MVHSGTVVGQLEWR	RuBisCO	P19163 RBL_NEUMU	GB	Basil
17	Y(Cl)NLSLGLGLNK	Serine-glyoxylate aminotransferase	P84187 SGAT_MAIZE	Cl	Nettle
18	KNQY*LEAEWNAK	Transketolase, chloroplastic	O20250 TKTC_SPIOI	GB	Laurel



different proteins were found in laurel compared to basil. Modifications in chloroplastic chlorophyll binding protein, chloroplastic transketolase, putative beta-D-xylosidase and rubisco were identified. The finding of rubisco was consistent with data obtained after chlorine exposure of nettle. In addition, similar chloroplastic chlorophyll binding proteins were targeted by all three chemical threat agents. An overview of all the detected plant-based peptides, including those that were only detected once, can be found in Table of the ESI.† The comprehensive list of peptides showed that multiple proteins were modified by all studied chemicals, such as chloroplastic ATP synthase, ferredoxin-dependent glutamate synthase, oxygen-evolving enhancer protein 1, photosystem II CP47 reaction center protein, RuBisCO activase and RuBisCO. From these results, it can be concluded that many proteins are targeted by the various chemicals. Interestingly, some proteins showed modifications in all three plant species by all studied chemical threat agents. This indicates that a specific peptide biomarker approach as recently presented for human biological samples might also be feasible for plant-based evidence.

4. Conclusions

In this study a novel approach was developed for analyzing persistent biomarkers in vegetation for retrospective investigation of chemical warfare agent exposure. We demonstrate that these plant biomarkers are identical to the analytes that have been accepted as unequivocal biomarkers of exposure in biomedical samples of victims and that have been used by the designated laboratories of the OPCW to analyze samples associated with chemical weapons attacks. This finding was confirmed through the use of reference standards. The experience with complex matrices, protein isolation protocols and enzymatic digestion steps is a distinct advantage when implementing the method developed in this study in forensic investigations, guaranteeing rapid implementation of the technology. In addition, the focus on plant-based evidence can offer a number of distinct advantages. This work shows that CWA biomarkers formed in plants are very persistent and can be detected up to three months after exposure. It is also expected that these biomarkers will be formed generically in plants irrespective of the species and exposure conditions. Furthermore, plant-based evidence is relatively easy to sample, ship, store and process and typically is abundantly available at an incident scene. Finally, for perpetrators it will be difficult to remove or manipulate plant-based evidence after a CWA attack. Future research could focus on processing more plant material to further reduce the detection limit and thereby the minimum exposure concentration. The analysis of case-work samples is required to establish the applicability of the method for concentrations and conditions encountered after a CWA attack. Both in terms of verifying the presence of plant biomarkers after a confirmed CWA exposure and of showing the absence of these biomarkers in situations where no threat agent was released. In addition, the combination of various analytical techniques can provide complementary information or additional proof through the presence of specific biomarkers. In conclusion, the

analysis of plant protein adducts can become a powerful tool to verify (or refute) the alleged use of chemical weapons.

Author contributions

Mirjam de Bruin-Hoegée: methodology, validation, formal analysis, writing – original draft, visualization; Latifa Lamriti: methodology, validation, formal analysis; Jan Langenberg: conceptualization, funding acquisition; René C. M. Olivier: methodology, formal analysis; Lai Fun Chau: methodology, formal analysis; Marcel van der Schans: conceptualization, methodology, writing – review & editing, supervision, project administration; Daan Noort: conceptualization, methodology, writing – review & editing, supervision; Arian C. van Asten: writing – review & editing, supervision.

Conflicts of interest

The authors declare that there is no conflict of interest.

Acknowledgements

This research is partially funded by the European Union, through a research contract awarded by the Organisation for the Prohibition of Chemical Weapons (OPCW), in support of the Plant Biomarker Challenge. This work is part of the Forensic Attribution for CWA INtelligence (FACING) project, a collaboration between the Van 't Hoff Institute for Molecular Sciences (HIMS) of the University of Amsterdam and TNO Defence, Safety & Security. The FACING project is financed by the DO-AIO fund of the Dutch Ministry of Defence.

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