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Strategic Environmental Research and Development Program (SERDP)

## Preparative, Extraction, and Analytical Methods for Simultaneous Determination of Legacy and Insensitive Munition (IM) Constituents in Aqueous, Soil or Sediment, and Tissue Matrices

Rebecca A. Crouch, Jared C. Smith, Bobbi S. Stromer,August 2021Christian T. Hubley, Samuel Beal, Guilherme R. Lotufo,Afrachanna D. Butler, Michelle T. Wynter, Dale A. Rosado,Amber L. Russell, Jessica G. Coleman, Jay L. Clausen,and Anthony J. Bednar



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### Abstract

No standard method exists for determining levels of insensitive munition (IM) compounds in environmental matrices. This project resulted in new methods of extraction, analytical separation and quantitation of 17 legacy and 7 IM compounds, daughter products of IM, and other munition compounds absent from USEPA Method 8330B. Extraction methods were developed for aqueous (direct-injection and solid-phase extraction [SPE]), soil, sediment, and tissue samples using laboratory-spiked samples. Aqueous methods were tested on 5 water sources, with 23 of 24 compounds recovered within DoD QSM Ver5.2 limits. New solvent extraction (SE) methods enabled recovery of all 24 compounds from 6 soils within QSM limits, and a majority of the 24 compounds were recovered at acceptable levels from 4 tissues types. A modified chromatographic treatment method removed analytical interferences from tissue extracts. Two orthogonal high-performance liquid chromatographyultraviolet (HPLC-UV) separation methods, along with an HPLC-mass spectrometric (HPLC-MS) method, were developed. Implementing these new methods should reduce labor and supply costs by approximately 50%, requiring a single extraction and sample preparation, and 2 analyses rather than 4. These new methods will support environmental monitoring of IM and facilitate execution of risk-related studies to determine longterm effects of IM compounds.

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### Preface

This study was conducted for the Strategic Environmental Research and Development Program (SERDP) under MIPR W74RDV70604798 under Project ER-2722, "Development and Optimization of Analytical Methods for Simultaneous Determination of IM and Legacy Explosive Compounds". The technical monitor was Dr. Elizabeth A. Ferguson.

The work was performed by the Environmental Chemistry Branch of the Environmental Processes Division, US Army Engineer Research and Development Center, Environmental Laboratory (ERDC-EL). At the time of publication, Amber L. Russell was Chief, Warren P. Lorentz was Chief, and Elizabeth Ferguson was the Technical Director for Military Environmental Engineering and Sciences. The Deputy Director of ERDC-EL was Dr. Jack E. Davis and the Director was Dr. Edmond J. Russo Jr.

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COL Teresa A. Schlosser was the Commander of ERDC, and Dr. David W. Pittman was the Director.

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### **1** Introduction

### 1.1 Background

No standard method currently exists for analyzing insensitive munition (IM) compounds in environmental matrices (with or without concurrent legacy compounds). Lacking established methods, contract and research laboratories either do not measure IM compounds at all; quantify some, but not all, using US Environmental Protection Agency (USEPA) Method 8330B (USEPA 2006) developed for legacy compounds (leading to inaccuracies for certain IM compounds and degradation products [Felt et al. 2016]); or use methods originally developed for neat materials with no proven performance metrics.

IM compounds are high-energy compounds that, while designed to match the performance of legacy materials, are less likely to detonate when exposed to unintended stimuli such as shock, heat, and adjacent detonating munitions (Barrie 2013). The development of IM began in the 1980s upon review of several unintended detonations at US Army facilities as a means to improve soldier safety and reduce resource loss (Newman 2010). The primary goal of the IM Program is to oversee the development and production of safer alternatives to replace 2,4,6-trinitrotoluene (TNT) and Composition-B (Comp-B) in US Military ordnance (Kiebler and Manz 2010). Comp-B contains TNT and hexahydro-1,3,5-trinitro-s-triazine (RDX). The compounds and formulations developed under the IM Program are required to meet certain criteria. They must be as effective as legacy munitions; they must show reduced sensitivity compared to legacy munitions; they must be inexpensive; and they must be scalable for production (Roos 2007). Of the numerous compounds and mixtures evaluated for use in IM formulations, three compounds-3-nitro-1,2,4triazol-5-one (NTO); nitroguanidine (NQ); and 2,4-dinitroanisole (DNAN)-hold the most promise as legacy replacements. IMX-101 (NTO, NQ, and DNAN) is the current frontrunner for replacement of TNT in US munitions packing (Picatinny Public Affairs 2010), and IMX-104 (NTO, DNAN, and RDX) is set to replace Comp-B (Mainiero 2015). Other compositions, such as PAX-21 (RDX, DNAN, and ammonium perchlorate) and PAX-41 (DNAN and RDX), have already seen use in theater (Fung et al. 2012).

Though care may be taken to follow strict munition handling guidelines, unintentional release of munition compounds into the environment could occur at any point during the manufacturing, testing, operation, or demilitarization processes. Contamination could occur due to leaky plumbing at manufacturing facilities, or at wastewater treatment plants, or from unintentional spills during transport. On the battlefield and on testing ranges, unconsumed munition compounds are dispersed at the target and at the point from which the shot was fired (from propellant) as small particles (Jenkins et al. 2001; Jenkins et al. 2005; Jenkins et al. 2006; Pennington et al. 2001, Pennington et al. 2002, Pennington et al. 2003, Pennington et al. 2004, Pennington et al. 2005, Pennington et al. 2006; Taylor et al. 2004). Corrosion and leakage of unexploded ordnance (UXO) can be a major long-term point source on impact areas (Brannon et al. 2000; Chendorain, Stewart, and Packer 2005).

Contamination found on munitions ranges is complex. The various chemical and physical properties of each munition compound, their transformation products, and the interferences arising from the environmental matrix make extraction and quantitation a challenge. Two factors complicate the assessment of environmental impacts from munitions contamination. First, new IM formulations will be used on training ranges where legacy munition contamination may be present. Second, daughter products of IM components, such as 2-/4-nitrophenol (2-/4-NP) or 2,4-dinitrophenol (2,4-DNP) (Hawari et al. 2015) from the degradation of DNAN, appear as IM compounds degrade. Aminotoluenes and nitrotoluenes from the degradation of TNT are examples of degradation products that might also be present if legacy munitions had previously been used at the site (Walsh and Jenkins 1990). Other legacy munitions, such as picric acid (PA) may also be present on training ranges but are not included in USEPA Method 8330 (Goerlitz 1979).

Current high-performance liquid chromatography (HPLC)–and gas chromatography (GC)–based analytical methods require preconcentration by solid-phase extraction (SPE) prior to analysis in order to achieve sub- $\mu$ g/L<sup>1</sup> detection limits in water samples. While methods of extracting legacy munitions from water by SPE have been standardized, and several

<sup>1.</sup> For a full list of the spelled-out forms of the units of measure used in this document, please refer to US Government Publishing Office Style Manual, 31st ed. (Washington, DC: US Government Publishing Office, 2016), 248–52, https://www.govinfo.gov/content/pkg/GPO-STYLEMANUAL-2016/pdf/GPO-STYLEMANUAL-2016.pdf.

SPE methods are available for the extraction of nitrophenols, only one has been developed for NTO (Walsh 2016). To date, no unified SPE method has been developed that allows for coextraction of NTO, NQ, DNAN, PA, 2,4-DNP, and 2-/4-NP. Lack of such a method means that detection limits are significantly elevated, because IM-containing waters must undergo direct-injection analysis by HPLC.

The traditional acetonitrile (ACN) extraction method for explosives in soils described in USEPA Method 8330B results in poor extraction efficiency of NTO, NQ, PA, 2,4-DNP, and 2-/4-NP. The compounds are only slightly soluble in acetonitrile but have relatively high solubility in water at 13 g/L for NTO, 2.6 g/L for NQ, 17.6 g/L for PA, 5.6 g/L for 2,4-DNP, 2.1 g/L for 2-NP, and 16 g/L for 4-NP. Furthermore, PA, NTO, and 2,4-DNP exist as anions at neutral pH due to their acidic nature, with pKa = 0.38, 3.76, and 4.11, respectively. Alternatively, NQ (pKa 12.2) can exist as either a nitroimine (predominant at environmental pH) or nitroamine, and at very low pH could become protonated to form a cation (De Vries and St. Clair-Gantz 1954; Kaplan, Cornell, and Kaplan 1982). DNAN is the exception of the IM compounds, with water solubility comparable to TNT (approximately 200 mg/L) (Lide 2005). Indeed, DNAN has been extracted via the standard SPE procedure outlined in USEPA Method 3535 (USEPA 2007).

HPLC separations, coupled with ultraviolet (UV) absorbance or mass spectrometric (MS) detection and quantitation, are the gold standard for analysis of munition compounds (USEPA 2006). Many munition compounds are nitroaromatics (for example, nitrotoluenes and nitrobenzenes) or nitramines (for example, RDX) and are analyzed using USEPA Method 8330. Other energetic compounds, including NTO, DNAN, NQ, and PA, can also be determined using HPLC separations, though the mobile and stationary phases may be modified from the conditions reported in USEPA Method 8330B (USEPA 1996; Goerlitz 1979; Russell et al. 2014). The chemical properties of some of these compounds preclude direct application of the extraction, preconcentration, and analysis methods described in USEPA Methods 3535 and 8330B protocols without modification. Therefore, when analysis of samples containing IM and legacy compounds is required, multiple analytical procedures would have to be employed, leading to increased costs and delays in data availability.

Much research is underway to determine the effects of NTO, DNAN, and NQ in the environment. The lack of uniform methods for extraction of munition compounds in water, soil, and tissue is a barrier to determining the long-term environmental impact of IM compounds. The wide-ranging solubilities and ionic properties of NTO, NQ, DNAN, 2,4-DNP, 2-/4-NP, and legacy munition compounds necessitated the development and optimization of methods to effectively extract or extract and preconcentrate-denoted as *extract/(preconcentrate)* throughout this technical report-all of the IM and legacy compounds studied. This work fills current gaps in energetics determination methodology by developing a direct-injection and SPE method for waters, solvent extraction (SE) methods for soils and tissues, and two HPLC-UV and one HPLC-MS method for analysis of IM, IM transformation products, and legacy munition compounds. The implementation of these methods for simultaneous determination of IM and legacy munition compounds should lead to resource and labor cost savings, with a minimal increase in environmental monitoring efforts and no modification to existing samplecollection procedures.

### 1.2 Objective

The primary objective of this project was to fulfill the requirements of SERDP SON ERSON-17-02 by developing new methods of extraction, preconcentration, and analytical separation and quantitation of 17 legacy and 7 IM compounds, daughter products of IM, and other munition compounds absent from USEPA Method 8330B. The goal was to produce a single standardized method for simultaneous analysis of legacy and IM compounds in environmental matrices. Two HPLC-UV and one HPLC-MS methods were developed, as well as extraction methods for waters (direct-injection and SPE), soils (two-stage SE), and tissues (one-stage SE). An analytical interference reduction procedure was also developed for tissues.

#### **1.3 Technical approach**

HPLC methods were developed for simultaneous separation, detection, and quantitation of 24 legacy and IM compounds of interest and 2 surrogates. Primary- and secondary-column HPLC-UV methods, as well as a confirmatory HPLC-MS method, were developed by the Environmental Laboratory (EL) at the US Army Corps of Engineers (USACE) Engineer Research and Development Center (ERDC), adapted from existing methods (USEPA 2006; Russell et al. 2014). The compounds of interest were the 17 legacy compounds included in USEPA Method 8330B (USEPA 2006) and 7 IM compounds, daughter products, and others absent from USEPA Method 8330B, specifically, NQ, NTO, PA, DNAN, 2,4-DNP, 2-NP, and 4-NP. The linear dynamic ranges (LDRs), method detection limits (MDLs), and matrix effects for IM compounds, IM daughter products, and legacy munition compounds were studied using the developed extraction/(preconcentration) and analytical methods in aqueous and solid matrices.

SPE is a method for extraction and preconcentration of energetic compounds from environmental water samples in both USEPA Method 8330B and internal ERDC-EL IM methods. An SPE method was developed to extract and preconcentrate all 24 compounds of interest (and 2 surrogates) by evaluating a series of cation-exchange, anion-exchange, reverse-phase (polymeric and traditional C-18), and activated carbonbased SPE cartridges. Methods developed using reagent water samples were applied to four additional water matrices: (1) tap water–Vicksburg Municipal Water; (2) ground water—well water from Rayville, Louisiana; (3) river water—Yazoo River near Vicksburg, Mississippi; (4) sea water— Houston Shipping Channel. Water characterization, including pH, TOC, and TDS appear in Table 8 (section 3.4) The developed SPE method efficiently extracted all 24 compounds of interest and 2 surrogates within acceptable Department of Defense (DoD) Quality Systems Manual (QSM) Ver5.2 limits, except NQ (approximately 50% recovery), from each water type, with slight pH-related variability, particularly with NTO.

Current USEPA Method 8330B solvent extraction methods were modified to enable extraction of all 24 legacy and IM compounds of interest (and 2 surrogates) from soil samples by adjusting the solvent system and using a two-stage extraction procedure. The developed SE method efficiently extracted all 24 compounds of interest and 2 surrogates within acceptable DoD QSM Ver5.2 limits, using six different soil sources: (1) ASTM fat clay—CH-1; (2) Aberdeen clay—Aberdeen, Maryland; (3) Jefferson clay— Madison, Indiana; (4) Memphis silt—Memphis, Tennessee; (5) Ft. Riley clay—Riley, Kansas; (6) Yuma sand—Yuma, Arizona. All six soils showed a wide range of organic matter, pH, and other properties (listed in Table 14, section 3.6).

A tissue extraction method was also developed by slight modification to existing USEPA Method 8330B processes, with the majority of the 24

compounds and 2 surrogates being extracted within acceptable DoD QSM Ver5.2 limits via a single 18 h MeOH<sup>2</sup> (rather than ACN) extraction. Four different tissue matrices were tested, including (1) earthworms (*Eisenia fetida*, soil-dwelling invertebrate); (2) fathead minnows (*Pimephales promelas*, freshwater vertebrate); (3) polychaete worms (*Alitta virens*, marine invertebrate); and (4) perennial ryegrass (*Lolium perenne*, perennial plant). Varying degrees of analytical interferences arising from coextracted nontarget matrix components were observed for each tissue type, with the most interference occurring in earthworm tissue. A chromatographic interference reduction method, using common organic laboratory chromatography materials, was developed to remove these matrix-associated analytical interferences.

Stability studies were carried out using laboratory-spiked reagent water and three laboratory-spiked soils (Riley, Yuma, and ASTM fat clay) to verify whether current standard pre- and postextraction hold times of 7/14 days and 40 days, respectively, for semivolatile organics (USEPA 2014) would remain valid for mixtures of the 24 legacy and IM compounds and 2 surrogates. Samples were stored in triplicate under three different conditions: (1) cold/dark (4°C/amber vials); (2) room temperature/dark (25°C/amber vials); and (3) room temperature/light (25°C/clear vials) in order to distinguish photo- versus thermal-related degradation of each compound. In addition, the developed extraction and analysis (primary HPLC-UV) methods were applied to laboratory-spiked water (five samples) and soil (six samples) for verification by inter-laboratory testing at ERDC-EL and ERDC Cold Regions Research and Engineering Laboratory (CRREL). For these tests, samples were extracted and analyzed within current standard hold times for semivolatile organics.

# **1.4** Benefits and implications for future research and implementation

The development of these extraction and analysis methods to simultaneously analyze environmentally copresent legacy and IM compounds was necessary to enable future monitoring of co-occurring components at firing ranges, demilitarization facilities, manufacturing facilities, and environmental sites, where munitions are tested, produced,

<sup>2.</sup> For a full list of the spelled-out forms of the chemical elements used in this document, please refer to US Government Publishing Office Style Manual, 31st ed. (Washington, DC: US Government Publishing Office, 2016), 265, https://www.govinfo.gov/content/pkg/GPO-STYLEMANUAL-2016/pdf/GPO-STYLEMANUAL-2016.pdf.

and detected (Felt et al. 2013; Walsh et al. 2014). Currently, separate extraction procedures must be carried out to extract legacy and IM compounds. Furthermore, separate primary and secondary analytical methods must be used to quantify legacy and IM compounds. It is estimated that the simultaneous analysis of IM and legacy compounds using the developed primary and secondary HPLC-UV methods may reduce total analysis time by approximately 50% versus using four different analytical methods, as the previous process required. In addition, the labor and supply cost savings due to coextraction and sample preparation is at least 50%, as only a single extraction procedure and analytical sample preparation are required, versus up to four (including confirmation analyses and associated quality control [QC] samples), since the sample preparation for the primary and secondary methods are the same. Overall, these savings will lead to reduced laboratory costs and more rapid results acquisition. The current work will continue under ESTCP ER19-D1-5078 (Validation of Sample Extraction and Analysis Techniques for Simultaneous Determination of Legacy and Insensitive Munitions [IM] Constituents), to validate the methods described herein. Furthermore, the products of this work will contribute to the ability to conduct fate and transport studies for IM compounds by providing a standardized method for quantitation of IM and legacy materials needed to determine long-term effects.

### **2** Materials and Methods

### 2.1 Primary HPLC-UV method

The HPLC-UV and -MS methods developed under SERDP ER-2722 were modified from existing methods and established laboratory techniques. Previously developed (Russell et al. 2014) and currently used explosives analysis methods (USEPA 8330B, in-house IM method; see Figure 4, section 3.1) were used as a starting point and reference in the development of the primary and secondary HPLC-UV methods for SERDP ER-2722.

The primary HPLC-UV method for simultaneous separation and quantitation of 17 legacy and 7 IM compounds, daughter products, and other munition compounds not included in USEPA Method 8330B was developed by adapting previous methods. Method parameters and solvent gradient scheme are shown in Table 1. Method development was carried out using Phenomenex Synergi 4 $\mu$ m Hydro-RP, 80Å, 250 × 4.6 mm HPLC columns. An optional Phenomenex SecurityGuard AQ C18 precolumn guard cartridge was included, which could extend column lifetime—especially when analyzing samples containing complex environmental matrices. As indicated in Table 1, either a 0.1% TFA or 0.25% FA (v/v) solution in water can be used. (This was investigated during HPLC-MS method development [section 2.3], due to ion-suppression effects caused by TFA.) An example chromatogram is shown in Figure 1 (acquired using TFA). All chromatographic peaks were identified using individual reference standards.

The HPLC methods developed under Strategic Environmental Research and Development Program (SERDP) Environmental Restoration (ER)-2722 were adapted from a method originally developed for internal ERDC efforts funded by the Environmental Quality and Installations (EQI) program that investigated IM biogeochemistry. The SERDP ER-2722 project and previous EQI-funded efforts built on previously completed SERDP and Environmental Security Technology Certification Program (ESTCP) projects focused on legacy compounds (SERDP ER-1155, SERDP ER-1481, and ESTCP ER-2000628). The current work will continue under ESTCP ER19-D1-5078 (Validation of Sample Extraction and Analysis Techniques for Simultaneous Determination of Legacy and Insensitive Munitions (IM) Constituents), to validate the methods described herein. The efforts undertaken for SERDP ER-2722 were carried out in coordination with those for SERDP ER-2724 (PI: Dr. Roman Kuperman).

Total run time: 48.0 min; Detection wavelengths: 210, 254, 315 nm Column temperature: 25°C; Flow rate: 1.0 mL/min; Injection volume: 50 µL					
Time (min)	Reagent water (%)	MeOH (%)	0.1% TFA/water (%)*	ACN (%)	
0.00	89	3	3	5	
2.00	89	3	3	5	
2.20	52	40	3	5	
12.5	52	40	3	5	
19.0	57	35	3	5	
28.0	48	44	3	5	
32.0	48	44	3	5	
44.0	32	60	3	5	
44.1	89	3	3	5	
48.0	89	3	3	5	

# Table 1. Primary high-performance liquid chromatography–ultraviolet (HPLC-UV)separation solvent gradient scheme and other method conditions.(\*or 0.25% FA/water)

Figure 1. HPLC chromatograms of a mixed standard containing 5 mg/L of 24 legacy and IM compounds and 2 surrogates analyzed by the developed primary HPLC-UV method (210 nm: bottom line; 254 nm: middle line; 315 nm: top line). (Retention times are listed in Table 6.)





# 2.2 Secondary high-performance liquid chromatography–ultraviolet (HPLC-UV) method

A secondary HPLC-UV method was developed for confirmatory analysis using Restek Pinnacle II Biphenyl, 5  $\mu$ m, 150  $\times$  4.6 mm HPLC columns. Again, an optional Phenomenex SecurityGuard AQ C18 precolumn guard cartridge was included. The method parameters are shown in Table 2. As indicated, again either a 0.1% TFA or 0.25% FA (v/v) solution in water can be used to acidify the mobile phase for optimal chromatographic resolution. An example chromatogram is shown in Figure 2 (acquired using FA; retention times listed in Table 3). All chromatographic peaks were identified using individual reference standards.

Total run time: 35.0 min; Detection wavelengths: 210, 254, 315 nm Column temperature: 25°C; Flow rate: 0.9 mL/min; Injection volume: 50 μL					
Time (min)	Reagent water (%)	MeOH (%)	0.1% TFA/water (%)*	ACN (%)	
0.00	75	10	10	5	
2.50	75	10	10	5	
2.60	39	46	10	5	
9.00	39	46	10	5	
9.10	33.5	51.5	10	5	
15.0	44	41	10	5	
29.0	25	60	10	5	
29.1	75	10	10	5	
35.0	75	10	10	5	

Table 2. Secondary HPLC-UV separation solvent gradient scheme and other method conditions. (\*or 0.25% FA/water)



Figure 2. HPLC chromatograms of a mixed standard containing 10 mg/L of 24 legacy and IM compounds and 2 surrogates analyzed by the developed secondary HPLC-UV method (210 nm: bottom line; 254 nm: middle line; 315 nm: top line). (Retention times are listed in Table 3.)

Bold, italicized are IM compounds and degradation products.

Table 3. Retention	times (min) for the 24	compounds of	of interest and	2 surrogates,
	analyzed by the secor	ndary HPLC-UV	/ method.	

Peak #	Compound	Retention time (min)
1	NQ	2.33
2	NTO (210nm)	2.67
2	NTO (315nm)	2.67
3	PA (210nm)	5.86
4	PA (315nm)	5.86
5	o-NBA (surrogate)	6.87
6	НМХ	8.82
6	4-NP	8.98
7	RDX	9.22

Peak #	Compound	Retention time (min)	
8	2-NP	11.54	
9	NG	12.94	
10	NB	12.68	
11	3,5-DNA (210nm)	12.90	
12	3,5-DNA (254nm)	12.90	
13	2,4-DNP	<i>13.55</i>	
14	1,2-DNB (surrogate)	13.89	
14	4-Am-2,6-DNT	15.08	
15	2-Am-4,6-DNT	15.62	
16	1,3-DNB	17.13	
17	2-NT	18.56	
18	4-NT	20.47	
19	3-NT	20.92	
20	DNAN	22.53	
21	2,6-DNT	23.37	
22	PETN	24.51	
23	1,3,5-TNB	25.62	
24	2,4-DNT	26.63	
25	Tetryl	28.67	
26	2,4,6-TNT	30.59	

# 2.3 High-performance liquid chromatography-mass spectrometric (HPLC-MS) method

An HPLC-MS scan mode method was developed using an Agilent 6120B single quadropole (SQ) system equipped with an atmospheric-pressure chemical ionization (APCI) source for confirmatory analysis of the 24 compounds of interest and 2 surrogates. Spray-chamber and MS-detector parameters for the scan mode method are provided in Table 4. Chromatographic separation was achieved via the primary HPLC-UV method described in section 2.1. An example MS total ion chromatogram (TIC) is shown in Figure 3. Several ion masses were observed for each of the 26 compounds analyzed (shown in Table 5), with the exception of nitroglycerine (NG) and 2-, 3-, or 4-nitrotoluene (2-NT, 3-NT, 4-NT), which did not produce any detectable MS signal under the developed spray chamber and detector conditions. In addition, 1,3-dinitrobenzene (1,3-DNB) yielded only a single observed ion. Ions were identified where possible. Observed ions are listed in order of abundance for each compound.

Parameter	Value
Source	APCI
Ionization mode	Negative
Drying gas (°C)	350
Vaporizer (°C)	325
Drying gas (L/min)	4.0
Nebulizer (psig)	40
Corona (µA)	10
Capillary (V)	1,500
Mass range	40-400
Fragmentor	100
Gain	1.00
Threshold	0
Step size	0.20
Speed (µ/s)	743
Peak width (min)	0.060
Cycle time (s/cycle)	0.57

## Table 4. Spray chamber and detector parameters for the developed HPLC-MS scan mode method.



Figure 3. Total ion chromatogram (TIC) for a mixed standard containing 10 mg/L of 24 legacy and IM compounds and 2 surrogates analyzed by the developed

Bold, italicized are IM compounds and degradation products.

Table 5. Retention times (min), ion masses (m/z), and identities obtained for mixed
standard containing 24 legacy and IM compounds and 2 surrogates using the
developed HPLC-MS scan mode method.

Retention time (min)	Compound	lon mass (m/z)	lon identity
3.78		203.6	
	NQ	103.0	[M]-H
4.50	4/70	129.0	[M]-H
	ΝΤΟ	55.0	
7.64	o-NBA (surrogate)	122.0	
		167.0	[M]-H
9.21	НМХ	341.0	[M]-H+FA
		147.0	[M]/2-H
13.40		267.0	[M]-H+FA
	RDX	335.0	[M]-H+TFA
16.00	( )/2	108.0	
	4-NP	138.0	[M]-H

Retention time (min)	Compound	lon mass (m/z)	lon identity	
10.00		213.0	[M]-H	
16.80	1,3,5-INB	183.0		
18.20	24	228.0	[M]-H	
	PA	211.9		
10.00		183.0	[M]-H	
18.80	2,4-DNP	153.0		
00.70		138.0		
22.70	1,2-DNB (surrogate)	168.0	[M]-H	
23.55	1.3-DNB	168.0	[M]-H	
04.00		138.0	[M]-H	
24.60	2-1112	122.0	[M]-H2O-H	
07.40		182.0	[M]-H	
27.10	3,5-DNA	138.0		
		182.0	[M]-H+HOAc	
27.80	NB	153.0		
		183.0	[M]-CH3	
29.00	DNAN	152.9		
	<b>-</b>	241.0	[M]-H+NO <sub>2</sub> -*	
30.00	Tetryl	288.0	[M]-H	
	NG	226.1	[M]-H (not obsv.)	
24.20		226.0	[M]-H	
31.30	2,4,6-1N1	210.0	[M]-H <sub>2</sub> O-H	
		196.9	[M]-H	
35.10	4-AM-2,6-DN1	167.0		
		196.0	[M]-H	
35.95	2-AM-4,6-DN1	179.9	[M]-H <sub>2</sub> O-H	
07.54		152.0	[M]-30*	
37.54	2,6-DNT	122.1		
27.07	0.4 51/7	165.0	[M]-H <sub>2</sub> O-H	
37.87	2,4-DN1	181.0	[M]-H	
42.00 - 44.50	2-NT/3-NT/4-NT	136.1	[M]-H (not obsv.)	
		62.0	NO3-*	
45.10	PEIN	270.9		

Bold, italicized are IM compounds and degradation products.

\*Avci, Anilanmert, and Cengiz 2017; Gapeev, Signman, and Yinon 2003; Jiang 2010; Kinghorn, Milner, and Zweigenbaum 2005; Thurman and Ferrar 2012; and Xu et al. 2004.

### 2.4 Method for preparation of waters for direct-injection analysis

Water samples known or suspected to contain the compounds of interest at levels detectable without extraction and preconcentration can be analyzed by direct injection using the developed analytical HPLC-UV or -MS methods. In order to ensure the accurate analysis of ionic species (including NTO), an acidification step was added for direct-injection water samples. A solution of 1% HCl/MeOH was used to acidify and dilute water samples for final acid concentration of 0.5% (v/v) and final solvent ratio of 50/50 MeOH/water.

### 2.5 Solid-phase extraction (SPE) method for waters

The SPE method developed under SERDP ER-2722 involved the sequential stacking of three different commercially-available SPE cartridges: (1) Strata X (polymeric reverse phase, 500 mg/6 mL, Phenomenex); (2) Strata X-A (polymeric strong anion exchange, 500 mg/6 mL, Phenomenex); and (3) Supelclean ENVI-Carb (granular activated carbon, GAC, 500 mg/6 mL, Supelco). The SPE cartridge conditioning, loading, and elution procedures were developed using common SPE methods for environmental water extraction and preconcentration. Specifically, SPE cartridges were conditioned with two 5 mL aliquots of MeOH and then equilibrated with two 5 mL aliquots of reagent water. Approximately 2 mL of reagent water was then added to each conditioned cartridge (to prevent drying) before the cartridges were stacked in the following order for loading using SPE tube adapters (Supelco): (1) Strata X at the top, (2) Strata X-A in the middle, and (3) ENVI-Carb on the bottom. Each triple-stacked SPE unit was loaded with 100 mL laboratory-spiked water sample containing all 26 compounds of interest (including two surrogates) or blank (unspiked) water sample. In order to allow for mass-balance analysis, a small amount of pass-through water was collected (for example, breakthrough determinations).

After allowing loaded SPE cartridges to air-dry under vacuum for approximately 10 min to remove all remaining water, the stacking order of the cartridges was reversed so that ENVI-Carb was at the top, Strata X-A was in the middle, and Strata X was on the bottom. Each sample-loaded SPE unit was eluted first with 5 mL MeOH, followed by 5 mL 2% HCl/MeOH. Extracts were collected separately, stored at  $\leq$ 4°C, and later prepared for analysis. MeOH and acidified MeOH extracts were analyzed individually, or by combining known ratios of each, and diluting with reagent water, for a final solvent ratio of 50/50 MeOH/water.

### 2.6 Solvent extraction (SE) method for soils

The SE method developed under SERDP ER-2722 for soils was a two-stage extraction procedure, modified from the current USEPA Method 8330B. Sample collection, storage, drying, and grinding procedures were not modified. Specifically, soils were ground to <0.85 mm (US Standard sieve #20) particle diameter using a ceramic mortar and pestle, dried at 25°C in a forced-air convection oven in the absence of light. Soil samples were spiked (varying concentrations) with a mixed reference standard containing all 26 compounds of interest (including two surrogates) in a mixture of MeOH/ACN or with unspiked MeOH/ACN, and vortexed for 1 min to mix. Samples were allowed to dry again in the oven at 25°C and then mixed using a horizontal sample roller for up to 18 h at 4°C in the absence of light.

Samples were then extracted using a two-stage ultrasonication procedure. For each sample, 5 mL of MeOH per gram of soil was added, and soilsolvent mixtures were placed in a cooled (by constant water replacement) ultrasonic bath for 6 h in the absence of light. Soil-solvent mixtures were centrifuged, and the supernatant (MeOH extract) was collected by syringe filtering through a 0.45 µm PTFE hydrophobic disk filter. Next, 5 mL of 50/50 MeOH/water per gram of soil was added, and the soil-solvent mixtures were placed in the cooled ultrasonic bath for another 14 h in the absence of light. Again, soil-solvent mixtures were centrifuged, and the supernatant (MeOH-water extract) was collected by syringe filtering through a new 0.45 µm PTFE hydrophobic disk filter. Extracts were collected separately, stored at  $\leq 4^{\circ}$ C, and later prepared for analysis. Extracts were analyzed individually, or by combining known ratios of each, and diluting for a final solvent ratio of 50/50 MeOH/water.

### 2.7 SE method for tissues

The SE method developed under SERDP ER-2722 for tissues was a single 18 h ultrasonic extraction procedure, modified from the current USEPA Method 8330B. Briefly, whole tissues were ground without lyophilization (wet) under liquid nitrogen, using either a mill or stainless steel mortar and pestle, to as fine a particle size or paste as possible. Aliquotted samples were spiked with a mixed reference standard containing all 26 compounds of interest (including two surrogates) in a mixture of MeOH/ACN or with unspiked MeOH/ACN, vortexed for 1 min, and were homogenized well prior to extraction. No post-spike drying step was performed for tissue samples. Homogenized samples were extracted using a single 18 h ultrasonication with MeOH. For each sample, 5 mL of MeOH per gram of wet sample was added, and samples were placed in a cooled (by constant water replacement) ultrasonic bath for 18 h in the absence of light. Tissue-solvent mixtures were centrifuged, and the supernatant (MeOH extract) was collected by syringe filtering through a 0.45  $\mu$ m PTFE hydrophobic disk filter. Extracts were stored at  $\leq 4^{\circ}$ C and later prepared for analysis. MeOH extracts were analyzed after diluting with water for a final solvent ratio of 50/50 MeOH/water. (At cold temperatures, some tissue extracts were observed to develop a precipitate. These samples were filtered further using either 0.45  $\mu$ m PTFE hydrophobic or 0.20  $\mu$ m Anotop 10 inorganic membrane disk filters.)

# 2.8 Chromatographic interference reduction method for tissue extracts

The tissue interference reduction method developed under SERDP ER-2722 was a modified chromatographic method (Larson et al. 1999). Other common organic laboratory procedures were tested, including storage at low temperature or acidification to induce complete precipitation, followed by filtration. However, these methods were not as effective as chromatographic methods. Initial tests were performed using either neutral alumina or florisil, and later tests included combinations of these, basic alumina, or silica. In later tests, MeOH extracts of unspiked tissues were spiked for a final concentration of 6 mg/L of the 24 compounds of interest and 2 surrogates. Small-scale chromatography columns were prepared using 5 <sup>3</sup>/<sub>4</sub> in (14.6 cm) borosilicate pipettes, loaded with a total of 0.2 g packing material. Five different column packing schemes were used: (1) 0.1 g neutral alumina layered on top of 0.1 g florisil, (2) 0.1 g florisil layered on top of 0.1 g neutral alumina, (3) 1:1 mixed neutral alumina and florisil, (4), basic alumina, and (5) silica gel. Chromatography columns were wetted with MeOH just before loading 1 mL tissue MeOH extract. After the 1 mL MeOH extract had completely passed through the column, 1 ml of MeOH was used to elute, followed by 1 mL 2% HCl/MeOH. The resulting 3 mL of column-treated MeOH extract was vortexed to mix. Treated extracts were again stored at  $\leq 4^{\circ}$ C and later prepared for analysis by diluting with water for a final solvent ratio of 50/50 MeOH/water. Several of the treatment options were effective at

reducing the analytical interferences observed for the tissues. However, the silica gel option proved most effective for the majority of compounds in the majority of tissue types.

### 2.9 Stability study methods

Photo and thermal stability was investigated for each of the 24 compounds of interest and 2 surrogates by testing the current pre- and postextraction hold times for water and soils, which are 7/14 days for extraction, respectively, and 40 days for analysis for munition compound determinations performed using USEPA Method 8330B. Extraction and analysis hold times were not previously reported for tissues in Method 8330B.

Reagent water was batch-spiked, aliquotted, extracted (using the developed SPE method), and analyzed (using the primary HPLC-UV method) on Day o. Portions of laboratory-spiked water were stored under three different conditions until Day 7. The three storage conditions were (1) cold (4°C), dark (amber containers); (2) room temperature (25°C), dark; and (3) room temperature, light (clear containers). On Day 7, the stored water samples were extracted and analyzed. In addition to storing portions of laboratory-spiked water on Day 0, aliquots of Day 0 MeOH/acidified MeOH extracts were also stored under the same three conditions until Day 40, the stored MeOH and acidified MeOH extracts were analyzed using the primary HPLC-UV method. Included as part of the stability studies in water were direct-injection (no SPE) water samples, which were spiked at a higher level to enable detection without preconcentration.

Three soils were batch-spiked and extracted (using the methods described in section 2.6) and analyzed (using the primary HPLC-UV method) on Day 0/1 (due to the two-day extraction procedure). Portions of the laboratoryspiked soils were stored under the same three conditions as for water samples until Day 14. On Day 14, the stored soil samples were extracted (Day 14/15) and analyzed (Day 15). Again, in addition to storing portions of laboratory-spiked soils on Day 0, aliquots of Day 0/1 MeOH and MeOH/water extracts (combined) were also stored under the same three conditions until Day 40. On Day 40, the stored extracts were analyzed using the primary HPLC-UV method.

### 2.10 Interlaboratory batch analysis methods

An interlaboratory batch study was carried out at ERDC-EL and ERDC-CRREL. Briefly, five water sources (reagent, tap, river, sea, and ground) and six soil sources (Aberdeen, Jefferson, Memphis, Riley, Yuma, and ASTM fat clay) were batch-spiked, homogenized, aliquotted, and distributed to each laboratory for extraction using the developed SPE and SE methods, followed by analysis using the developed primary HPLC-UV method. Samples were spiked with the 24 compounds of interest and 2 surrogates. High-level, direct-injection water samples were spiked at 2 mg/L, and low-level water samples for SPE extraction and preconcentration were spiked at 0.10 mg/L. Soils were spiked at 20 mg/kg. Each laboratory received identical protocols for sample extraction, analytical sample preparation, and analysis.

During cross-laboratory studies using the direct injection method for waters, much lower recoveries for NTO, and in some cases *o*-NBA surrogate, were observed for field water samples than for reagent water. Therefore, further experiments were carried out in which the pH of samples to be analyzed by direct injection was adjusted using 2% HCl/MeOH (the same used for stage two of SPE elution), so that the final acid concentration was the same as in SPE extracted and preconcentrated samples, or double. Briefly, reagent water, Vicksburg municipal water (tap), and three field water samples were spiked with the 24 compounds of interest and 2 surrogates for a final concentration of 1 mg/L and either 0.5% or 1.0% HCl (v/v). The acidified, laboratory-spiked water samples were diluted 1:1 with MeOH and analyzed using the developed primary HPLC-UV method.

### **3** Results and Discussion

### 3.1 Primary HPLC-UV method

Previously developed (Russell et al. 2014) and currently employed munition compound analysis methods (USEPA 8330B, in-house IM method) were used as references in the development of the primary HPLC-UV method. Figure 4 shows the isocratic or gradient methods referenced. The separation solvent gradient scheme and other method conditions for the primary HPLC-UV method itself are shown in Table 1 in section 2.1. Peak identities (obtained by analysis of individual reference standards), method detection limits (MDLs), and linear dynamic ranges (LDRs) are provided in Table 6.

Figure 4. Previous HPLC methods used for munition compound determinations, including (a) current USEPA Method 8330B employed at the Engineer Research and Development Center–Environmental Laboratory (ERDC-EL) for HPLC-UV analysis of legacy compounds, (b) method developed at ERDC-EL for HPLC-MS analysis of insensitive munition (IM) compounds (Russell et al. 2014), (c) current method employed at ERDC-EL for HPLC-UV analysis of IM compounds, and (d) primary column HPLC-UV method developed for simultaneous analysis of legacy and IM compounds for SERDP ER-2722.



Table 6. Retention times (min), method detection limits (MDLs) and linear dynamic range (LDR) upper limits (ULs) for the 24 compounds of interest and 2 surrogates, determined in direct-injection water (Est. MDL), reagent water that underwent SPE preconcentration, and Ottawa sand that underwent SE, analyzed by the primary HPLC-UV method.

		Retention	Est. MDL ( <i>n</i> = 10 at 40	MDL in reagent water ( <i>n</i> = 10 at	MDL in Ottawa sand ( <i>n</i> = 10 at 4	LDR UL
Peak #	Compound	time (min)	µg/L) (µg/L)	40 µg/L) (µg/L)	mg/kg) (mg/kg)	(mg/L)
1	NQ	3.51	16	12	2.64	≥80
2	NTO (210nm)	4.29	6	20	1.26	≥50
2	NTO (315nm)	4.29	9	15	0.82	≥60
3	o-NBA (surrogate)	7.45	12	49	0.32	≥80
4	НМХ	8.96	6	11	1.18	≥70
5	RDX	12.81	6	9	0.66	≥70ª
6	PA (210nm)	14.64	10	23	3.88	<i>≥70</i> ª
6	PA (315nm)	14.58	12	11	1.44	≥80¢
7	4-NP	15.31	10	7	0.94	≥80
8	1,3,5-TNB	16.17	11	13	1.14	≥80
9	2,4-DNP	17.86	9	22	4.42	≥80
10	1,2-DNB (surrogate)	20.35	15	16	2.38	≥70
11	1,3-DNB	21.80	7	12	0.94	≥80
12	2-NP	22.91	12	14	1.26	≥80
13	NB	25.42	9	10	1.74	≥80
14	3,5-DNA (210nm)	26.16	13	16	3.02	≥70
14	3,5-DNA (254nm)	25.93	13	13	0.38	≥70
15	DNAN	26.89	11	16	1.58	≥70
16	Tetryl	28.43	12	8	2.08	≥70
17	NG	29.30	17	27	5.06	≥70
18	2,4,6-TNT	30.06	15	11	2.14	≥70
19	4-Am-2,6-DNT	32.60	11	17	2.56	≥80
20	2-Am-4,6-DNT	33.43	11	22	2.50	≥80
21	2,6-DNT	35.25	14	37	2.82	≥70
22	2,4-DNT	35.57	7	12	0.70	≥80
23	2-NT	40.18	13	14	1.08	≥80
24	4-NT	41.42	12	16	2.00	≥80
25	3-NT	42.84	14	21	0.84	≥80
26	PETN	44.38	16	22	3.46	≥80

Bold, italicized are IM compounds and degradation products.

a. coelute at concentrations >30 mg/L; b. RDX not detected appreciably at 315 nm

In the table, LDR values represent the highest concentration of spiked standard analyzed within ±10% of the known spike concentration, using a complete mixed reference standard of the 24 compounds of interest and 2 surrogates. Because most of the commercially available individual reference standards and mixes were acquired at 1,000 mg/L in MeOH or ACN, solutions at higher concentrations (>50 mg/L) contained greater than 1:1 MeOH/water ratios, which was determined to be the optimal analytical sample solvent mixture (see Figure 9). Because previous studies indicated that MeOH/water ratios of 9:1 caused chromatographic deterioration (see Figure 9), 80 mg/L with 8:1 MeOH/water content was the highest concentration tested. The upper limit of the linear range for some compounds may be higher than 80 mg/L. Results for multiple wavelengths were included for NTO, PA, and 3,5-DNA, to allow for quantitation using either wavelength. In some cases, detection of NTO and PA at 315 nm may be preferred in order to minimize the effects of analytical interferences-for example, with tissue samples. The PA chromatographic peak at 315 nm is inverted (versus at 210 nm) and lies below the baseline. For 3,5-DNA, the detection limits obtained for 210 nm and 254 nm were typically very similar, and either wavelength could be used for quantitation. MDLs were determined for direct-injection water samples (*Est. MDL*), SPE water samples (*MDL in reagent water*), and SE solid samples with Ottawa sand (MDL in Ottawa sand).

For simultaneous legacy and IM compound analysis, it was clear that a more complex gradient of the major aqueous (water) and organic phases (MeOH) would be required, with the final method incorporating elements of all three of the methods referenced (Figure 4a-c). Specifically, the primary HPLC-UV method gradient started with very low percentage of MeOH that increased sharply after 2 min, initiating a period of approximately 30 min during which the percentage of water was to varying degrees slightly greater than the percentage of MeOH, followed by a period of approximately 10 min during which the percentage of MeOH surpassed the percentage of water and continued to increase up to 60%. A brief re-equilibration period was placed at the end of the gradient method in order to allow the system time between sample injections to reequilibrate to initial conditions. The small decrease in MeOH from 40% at 12.5 min to 35% at 19 min (Figure 4d) was introduced in order to improve separation of the hydrophobic compounds that elute between 20–30 min, as neither isocratic conditions nor a linear increase in percentage of MeOH from 12.5 to 28 min resulted in well-resolved, stable chromatography. The

developed primary HPLC-UV method incorporated conditions favorable for separation of the 24 chemically disparate compounds of interest (and 2 surrogates). Further details on the optimization of the initial percentage MeOH, as well as the isocratic percentage of 0.1% TFA (or 0.25% FA) and percentage of ACN throughout the method are discussed below.

Throughout primary column HPLC-UV method development, NQ and 2,6-/2,4-DNT resolution remained the most challenging. Resolution of 2,6-DNT and 2,4-DNT was observed to vary greatly across manufacturer production lots and individual parts; however, resolution tended to be improved for well-conditioned, newer columns. Therefore, a decrease in the resolution of these two compounds could signal that a column has degraded, and a new column could improve analyses. Meanwhile, NQ, as anticipated, seemed to suffer the most from solvent effects, specifically changes in MeOH concentration, resulting in peak splitting and migration, so that the optimal initial percentage of MeOH determined for the primary HPLC-UV separation served to minimize these negative effects but did not completely eliminate them. In fact, NQ resolution (retention time and peak shape) depended greatly on the aqueous-to-organic ratio in both sample solvent matrix and elution solvent gradient, with comprehensive studies discussed below. Therefore, secondary confirmation of NQ by the secondary HPLC-UV or -MS method may be necessary to confirm results obtained using the primary column.

As previously mentioned, solvent effects on the gradient method, as well as on sample preparation, were studied. Preliminary method development results indicated that a complex, multistage gradient would be required for separation of all compounds of interest and that the most effective initial percentage of MeOH concentration was likely to be <10%. It was, thus, necessary to determine the initial percentage of MeOH that would yield the most stable retention times and peak shapes for NQ and NTO, the most polar and earliest-eluting compounds of interest, to ensure reliable quantitation. The effect of varying the solvent gradient initial percentage MeOH from 2% to 15% was studied, while holding the isocratic 0.1% TFA and ACN constant at 3% and 5%, respectively, using a 5 mg/L mixed standard containing all 24 compounds of interest and 1,2-DNB (surrogate) prepared in 50/50 MeOH/water. Little effect was observed for any compounds other than NQ and NTO (data not shown). Both NQ and NTO underwent large retention time shifts (>0.5 min) and changes in peak shape, including reduced intensity and splitting of NTO at >10% MeOH

(Figure 5). Meanwhile, NQ shifted almost entirely into the void peak but did become sharper as the MeOH concentration increased. Overall, it was determined that 3% MeOH yielded the best NTO peak shape and intensity, as well as NQ distance from both the void peak and NTO, with sufficient baseline resolution for reliable quantitation.

The effect of varying the isocratic mobile phase component 0.1% TFA/water from 2% to 15% was studied, while holding the initial MeOH and isocratic ACN constant at 3% and 5%, respectively, using a 5 mg/L mixed standard containing all 24 compounds of interest and 1,2-DNB (surrogate) prepared in 50/50 MeOH/water. Little effect was observed for compounds with retention times greater than 20 min (data not shown), which were generally nonionic. NQ was for the most part unaffected by changes in % TFA (Figure 6a). The most significant effects were observed for PA, NTO, and 2,4-DNP, which were the most acidic compounds studied (pKa = 0.38, 3.76, and 4.11, respectively). Meanwhile, NTO underwent a great deal of peak broadening (from a peak width of 0.1 min to 0.3 min) and retention time shift of approximately 0.5 min (Figure 6a) as 0.1% TFA/water increased from 2% to 15%.



In addition, both PA and 2,4-DNP underwent extreme peak shifting (ranging 12–15 min and 16.5–18 min, respectively), and shifted inversely with respect to one another, on the basis of the TFA concentration. At lower pH, PA shifted to earlier retention times, while 2,4-DNP shifted to later retention times. No analytical (coelution) challenges arose because of 2,4-DNP peak shifting.

Conversely, PA coeluted with 4-NP at 2% 0.1% TFA/water, shifted left to coelute with RDX with the 0.1% TFA/water set to 8% of the mobile phase composition, and eluted before RDX by 15% 0.1% TFA/water. In another part of these studies (data not shown), PA was also observed to undergo retention time shifts over a range of approximately 1 min as a result of increased or decreased sample pH arising from changes in the concentration of PA itself. Higher concentrations of PA resulted in earlier elution (~13 min), and lower concentrations of PA resulted in later elution (~14 min), similar to the TFA-dependent shifting observed in Figure 6b. This wide variability of PA retention time was due to its very low pKa, which
was not buffered well by the TFA in the mobile phase. Over the course of the current effort, PA concentration–dependent PA peak shifting was observed over a range of concentrations because of pH variability in the sample matrix or in the prepared 0.1% TFA/water mobile phase, or both. Considering all effects of TFA concentration on the primary HPLC-UV method chromatography, a 3% isocratic mobile phase composition of 0.1% TFA/water was chosen, to enable efficient and consistent quantitation of NTO and PA. As a technical note, the extreme pH sensitivity observed for PA highlighted the importance of careful preparation of the 0.1% TFA/water solvent to ensure chromatographic stability.

The effect of varying the solvent gradient isocratic percentage of ACN from 5% to 8% was studied, while holding the initial MeOH and isocratic 0.1% TFA/water constant at 3%, using a 5 mg/L mixed standard containing all 24 compounds of interest and 1,2-DNB (surrogate) prepared in 50/50 MeOH/water (Figure 7). While retention time shifts for NQ and NTO from 3.6 min to 3.5 min and 4.3 min to 3.9 min, respectively, as percentage of ACN increased from 5% to 8%, peak shape, resolution, and intensity were not affected by changes in percentage of ACN (Figure 7a). However, ACN concentration was shown to impact the resolution of middle- and lateeluting compounds, including most of the legacy munition compounds (present in the current USEPA Method 8330B), resulting in an overall decrease of 2-3 min method run time, and one or more coelutions at >6% ACN (Figure 7b). Because there were no coelutions at either 5% or 6% ACN, and because some compounds were better resolved either at 5% (1,3-DNB/2-NP) or at 6% ACN (Tetryl/NG), the use of either 5% or 6% isocratic ACN would be appropriate.

Cursory experiments were carried out to investigate whether the gradient method could be shortened by ramping up the percentage of ACN during the final 20 min of the method (Figure 8). For this test, 6% ACN was used from 0–28 min then was increased from 6% to 20% from 28–44 min. Total run time was shortened by 5 min; however, the introduction of an ACN gradient would only be recommended in cases where the additional complication to the method resulted in significant benefit to the user.

Further experiments showed that NQ underwent substantial changes in retention time (from 3.3 min to 3.9 min) and peak shape, including splitting and broadening, resulting in several small peaks that appeared near (and possibly included within) the void peak (2.7 min) and one main

peak that shifted closer to the NTO peak as percentage of MeOH composition increased (Figure 9a). Overall, a 50/50 MeOH/water sample solvent composition was determined to be optimal. Neither the NTO nor any of the later-eluting analytes were negatively affected by a 50% MeOH composition, and adequate baseline resolution enabled analysis using the main NQ peak (3.65 min), which yielded a linear calibration curve from 0.020 mg/L to 20 mg/L ( $R^2 \ge 0.97$ ) with MDL of 16 µg/L (Table 6). For the majority of included compounds, no peak broadening or shifting was observed up to 75% MeOH, with the first signs of negative chromatographic effects appearing at 90% MeOH between 8 and 28 min (Figure 9b). Overall, NQ posed the greatest analytical challenges, because of its highly polar character. The solvent gradient and sample solvent composition conditions developed for the primary HPLC-UV method served to minimize negative effects on NQ retention time and peak shape but did not eliminate them, so that a confirmation method (such as the secondary HPLC-UV or -MS method described) should be used to confirm NQ determinations.

Figure 6. Effect of varying the solvent gradient isocratic 0.1% TFA/water from 2% to 15%, while setting the initial MeOH at 3% and holding ACN constant at 5%, using 5 mg/L mixed standards containing all 24 compounds of interest and 1,2-DNB surrogate prepared in 50/50 MeOH/water. (a) Effect on NQ and NTO, (b) Effect on PA and 2,4-DNP retention times (min) and peak shapes.



(a)



(b)

Figure 7. Effect of varying the solvent gradient isocratic % ACN from 5% to 8%, while setting the initial % MeOH at 3% and holding % 0.1% TFA/water constant at 3%, using 5 mg/L mixed standards containing all 24 compounds of interest and 1,2-DNB surrogate prepared in 50/50 MeOH/water. (a) Effect on NQ and NTO, (b) Effect on middle- and late-eluting compounds.







Figure 9. Effect of varying the sample solvent composition % MeOH from 0% to 100%, using the developed primary HPLC-UV method to analyze 5 mg/L mixed standards containing all 24 compounds of interest and 1,2-DNB surrogate prepared with indicated MeOH/water solvent ratios. (a) Effect on NQ and NTO, (b) effect on middle-and late-eluting compounds.



(a)



## 3.2 Secondary HPLC-UV method

A secondary HPLC-UV method was developed to enable confirmatory analysis of the 24 legacy and IM compounds of interest and 2 surrogates. The solvent gradient scheme and other method parameters are shown in Table 3, and an example chromatogram is shown in Figure 2 (with peak assignments determined using individual reference standards), in section 2.2.

A graphical representation of the solvent gradient scheme for the secondary HPLC-UV method is shown in Figure 10. The initial portion of the gradient (0-2.5 min) was designed to elute the highly polar NQ and NTO. The next stage consists of a brief isocratic period (2.5-9 min), followed by a spike and then a gradual decrease of the percentage of MeOH (9-15 min), was designed to elute and resolve the more hydrophobic compounds. The third stage of the gradient (15-29 min), with an increase to 60% MeOH composition, was designed to elute the most hydrophobic compounds, decrease their retention times, and reduce overall method run time. A brief period at the end of the method returned the system to initial conditions to re-equilibrate before the next sample injection.

Three difficulties were addressed during the development of this method. Initially, (1) PA eluted within a void peak (6.5 min) resulting from a sharp shift in aqueous-to-organic ratio at 4.1 min (rather than the final 2.6 min), visible at 210 nm; (2) HMX, 4-NP, and RDX co-eluted with each other; and, (3) NQ was poorly resolved at 210 nm. To mitigate these issues, the isocratic percentage of 0.1% TFA/water and percentage of ACN were optimized, and small adjustments to solvent gradient timing and the percentage of MeOH shifts were tested. By making slight changes in the percentage of MeOH gradient, the HMX, 4-NP, and RDX became well resolved. While NQ and PA became better resolved through solvent gradient method optimization, the issues with their chromatography were not eliminated. Therefore, an alternative option for improved chromatography could be to use alternate detection wavelengths, such as 254 nm for NQ and 315 nm for PA, rather than using 210 nm, where void peaks did not appear, as these wavelengths were distant enough from the MeOH UV cutoff (205 nm) and should result in acceptable detection limits.





## 3.3 HPLC-MS method

An HPLC-MS scan mode method was developed to enable confirmatory analysis of the 24 legacy and IM compounds of interest and 2 surrogates. The solvent gradient scheme and other HPLC method parameters were unchanged from the primary HPLC-UV method described in section 2.3.

Efforts were made to transfer the LC-MS scan method to a selective-ion monitoring (SIM) method to improve detection limits by selectively detecting either one or two of the ion masses observed for each compound in scan mode experiments. The experimentally acquired ion masses obtained using the developed scan mode method, described in section 2.3 (Table 5 and Figure 3), were used to create ion-time windows for a SIM method. Several iterations with varying numbers of windows and times were attempted, with the most successful shown in Table 7 and Figure 11. The only other parameter that was changed from the scan mode method for the chromatogram shown in Figure 11 was the gain (2.0), which was increased to boost the signal. The peak width setting of 0.06 min resulted in total dwell times of between 330 and 345 msec per SIM group, which was equally divided among the SIM ions in each group, as a fixed function of the system's software. Retention times were approximately the same as those for the scan mode method (Table 5). The compounds in SIM Group 1 (NQ, NTO, o-NBA, and HMX) were not observed. Most compounds in SIM Groups 2 and 4 were detected well, except for PA, 2,4-DNP, NB, and NG, which were not observed. Meanwhile, all the compounds in SIM Group 3 were observed. For SIM Groups 5 and 6, just one compound each was observed, 4-Am-2,6-DNT and PETN, respectively. Because of the issues encountered during SIM mode method development, confirmatory analysis should be performed using the scan mode method.

SIM window start time (min)	SIM window group #	Compound	SIM ion (m/z)	Dwell time (ms)
		NQ (1)	203.6	42
		NQ (2)	103.0	42
		NTO (1)	129.0	42
0.00		NTO (2)	55.0	42
0.00	1	o-NBA (surr.) (1)	122.0	42
		o-NBA (surr.) (2)	167.0	42
		HMX (1)	341.0	42
		HMX (2)	147.0	42
		RDX (1)	267.0	37
		RDX (2)	335.0	37
		4-NP (1)	138.0	37
		4-NP (2)	108.0	37
11.50	2	1,3,5-TNB (1)	213.0	37
		1,3,5-TNB (2)/ <i>2,4-DNP (1)</i>	183.0	37
		2,4-DNP (2)	153.0	37
		PA (1)	228.0	37
		PA (2)	211.9	37
		1,2-DNB (surr.) (1)/ <i>2-NP (1)</i>	138.0	115
20.50	3	1,2-DNB (surr.) (2)/1,3-DNB (1)	168.0	115
		2-NP (2)	122.1	115
		NB (1)/3,5-DNA (1)	182.0	37
25.50	4	NB (2)	153.0	37
		3,5-DNA (2)	138.0	37

Table 7. Selective-ion monitoring (SIM) windows developed from UV peak retention times (min) and ion masses (m/z) observed using the developed scan mode method.

SIM window start time	SIM window		SIM ion	Dwell time
(min)	group #	Compound	(m/z)	(ms)
		DNAN (1)	183.0	37
		DNAN (2)	152.9	37
		Tetryl (1)	241.0	37
		Tetryl (2)	288.0	37
		2,4,6-TNT (1)/NG (not obsv.)	226.0	37
		2,4,6-TNT (2)	210.0	37
		4-Am-2,6-DNT (1)	196.9	42
		4-Am-2,6-DNT (2)	167.0	42
		2-Am-4,6-DNT (1)	196.0	42
20 50		2-Am-4,6-DNT (2)	179.9	42
32.50	5	2,6-DNT (1)	152.0	42
		2,6-DNT (2)	122.1	42
		2,4-DNT (1)	165.0	42
		2,4-DNT (2)	181.0	42
		2-NT/3-NT/4-NT (not obsv.)	136.1	115
40.00	6	PETN (1)	62.0	115
		PETN (2)	270.9	115

**Bold, italicized** are IM compounds and degradation products. \*SIM window start time = LC-MS method run time (no MS delay)



Figure 11. HPLC-MS TIC acquired by SIM mode method. SIM window start times are indicated by dotted lines, and SIM group numbers are noted above the plot area. (Retention times are listed in Table 7.)

Bold, italicized are IM compounds and degradation products.

## 3.4 Method for preparation of waters for direct-injection analysis

During the direct-injection water portion of the interlaboratory study (section 3.10), much lower recoveries for NTO, and in some cases o-NBA surrogate, were observed for field water samples than for reagent water. As this trend seemed related to pH, with NTO recovery being lower for higher-pH water samples (Table 8), further experiments were carried out in which the pH of samples to be analyzed by direct injection was adjusted. A solution of 2% HCl/MeOH (the same used for stage two of SPE elution) was used, for final acid concentrations equal (0.5%, v/v) or double (1%, v/v)v/v) that of corresponding SPE extracted and preconcentrated samples. Recovery results for the acidified direct-injection water samples are shown in Table 9. For each field water source, acidification with HCl resulted in near 100% recovery for NTO and o-NBA. Moreover, an acid concentration of 0.5% was sufficient, and 1% acid was not necessary. Therefore, for water samples with  $pH \ge 6.5$  (approximately) to be analyzed by direct injection, acidification is recommended. Furthermore, for higher-pH samples extracted and preconcentrated by SPE that are known or suspected to

contain NTO, additional acidification may also be necessary (as for the pH 9.2 tap water used in the current study).

			Total orga (TOC) (	nic carbon (mg/L)	Total dissolved solids (TDS) (mg/L)		
Water source	рН	Conductivity (µS/cm)	AVG ( <i>n</i> = 3)	Std Dev	AVG ( <i>n</i> = 3)	SD	
Reagent (ERDC-EL)	6.30	3.53	0.00	0.00	-11	1.2	
Tap (Vicksburg, MS)	9.16	208.1	1.11	0.12	125	1.2	
Ground (Rayville, LA)	8.10	456.6	1.31	0.33	264	3.3	
Yazoo River (Vicksburg, MS)	8.22	254.2	5.62	0.03	198	2.0	
Sea (Houston Shipping Channel)	7.84	13,350	5.14	0.05	8,910	7.2	

Table 8. Characteristic properties of five water sources used in interlaboratory batch studies. (AVG = average; SD = standard deviation)

	Recover	ri <b>es</b> (%)																		
	0.5% H0	CI (vol/vo	ol)								1% HCI	(vol/vol)	)							
	Reagen	t water	Tap wat	er	Yazoo R water	iver	Houstor Shippin Channe	n g I water	Well wat	ter	Reagen	t water	Tap wat	er	Yazoo R water	iver	Houstor Shippin Channe	ı g I water	Well wa	ter
	AVG		AVG		AVG		AVG		AVG		AVG		AVG		AVG		AVG		AVG	
Compound	( <i>n</i> = 2)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD
NQ	99	5.4	110	2.2	101	2.2	107	7.4	101	6.4	109	6.1	108	12.2	120	14.4	112	17.6	106	19.0
NTO (210nm)	111	10.4	114	3.9	117	7.0	119	2.4	111	8.0	118	9.9	108	6.5	132*	15.6	125*	21.4	112	11.0
NTO (315nm)	110	14.7	112	6.2	120	3.9	121	3.8	114	4.6	121	10.2	113	6.2	131*	15.6	121	20.4	117	11.2
o-NBA (surrogate)	94	9.0	98	5.8	98	5.8	99	1.9	96	7.1	104	4.1	104	8.5	112	11.9	102	17.2	102	11.6
HMX	94	9.1	97	5.4	95	5.9	98	4.2	94	0.9	101	10.1	98	8.1	106	16.0	97	17.3	94	10.2
RDX	96	12.3	100	4.0	95	6.7	100	1.1	96	6.2	101	6.7	102	10.4	109	17.7	102	17.5	99	9.6
PA (210nm)	83	5.4	<i>93</i>	4.5	87	6.3	87	8.6	89	6.3	95	10.8	<u>98</u>	7.2	104	<i>23.7</i>	105	23.6	97	8.7
PA (315nm)	91	8.0	96	5.2	<i>93</i>	7.0	<i>93</i>	5.1	90	7.8	97	6.0	97	8.6	106	14.0	<u>98</u>	18.8	95	10.0
4-NP	95	12.0	<u>98</u>	5.1	<u>99</u>	6.1	<u>99</u>	4.8	94	6.3	<u>99</u>	6.7	95	8.9	108	14.1	100	17.2	96	10.5
1,3,5-TNB	92	10.9	98	3.0	96	4.8	96	3.3	92	4.5	97	6.1	98	5.3	105	15.4	98	17.9	94	8.9
2,4-DNP	86	9.6	<i>93</i>	7.7	92	5.9	94	7.3	87	3.8	100	3.2	<i>92</i>	11.2	107	13.6	94	18.2	92	10.6
1,2-DNB							2.6				. –						100	10.6		
(surrogate)	91	11.0	98	6.5	92	5.0	96	3.6	93	2.9	97	9.0	98	10.1	110	13.8	100	18.6	94	14.0
1,3-DNB	92	11.2	100	7.4	95	4.5	96	3.4	91	4.6	103	12.1	99	7.2	107	13.5	99	17.5	96	10.8
2-NP	88	13.5	<u>92</u>	6.8	<u>92</u>	5.9	<u>92</u>	4.1	88	2.4	<u>90</u>	4.5	84	9.6	99	17.7	<u>95</u>	13.1	91	<u>9.7</u>
NB	88	107.0	85	6.8	90	7.4	92	2.0	88	5.1	94	5.2	82	6.0	103	14.7	96	19.4	93	9.9
3,5-DNA (210nm)	106	12.4	109	6.6	101	4.9	106	3.1	98	3.7	102	1.1	103	11.4	115	17.1	106	18.1	102	10.8
3,5-DNA (254nm)	107	11.8	111	8.5	106	5.7	109	1.9	104	2.1	108	4.6	107	9.1	119	18.0	109	17.5	105	10.1
DNAN	87	12.3	92	5.0	86	3.9	88	2.1	83	3.8	89	3.6	89	6.8	<u>96</u>	10.0	<u>95</u>	15.4	88	11.2
letryl	94	7.5	99	3.5	96	5.1	97	5.6	95	7.1	98	4.4	104	15.7	107*	13.9	101	18.5	9/	9.4
	106	5.0	103	8.3	04	2.8	99	8.3	109	4.6	05	11.5	113	18.9	12/*	9./	118	15.4	108	9.6
2,4,6-1N1	92	0./	99	0.8	94	8.0	93	3.9	94	0.4	95	6.0	100	15.5	110	10.5	100	20.3	93	10.1
4-Am-2,6-DN1	93	11.0	80	4.4	95	7.0	98	6.4	94	/.6	96	8.3	/6	9.2	108	14.5	100	18.6	95	11.8
2-Am-4,6-DN1	93	11.4	83	2.3	95	5.6	102	4.6	99	1.8	99	8./	85	9.5	108	10.1	101	20.0	94	11.1
2,0-DN1	92	8.4	93	5.8	95	2.1	102	/.1	89	4.4	95	5.1	96	/.0	109	17.0	102	10.2	95	10.2
2,4-DNI	<u> </u>	9.5	9/	4.5	90	0.0 6.2	98	0.3	94	4.8	103	0.5	98	ð.ð	107	13.1	100	19.2	90	12.0
	98	0.9 0.9	07	1.Z	98	0.3	101	2.4 5.7	93	<i>L.L</i>	101	4.5	98	0.3	103	$\frac{2}{10}$	100	10.2	93	11.4
4-IN I 2 NIT	07	0.2	101	4.0	91	5.0	95	6.2	92	0.4 1 2	90	1.5	91	0./ //7	114	12./	95	14.1	94	0.5
	97	13.0	101	5.5 77	101	0.1	99	0.5	92	4.5	90	1.1	90	4./	107	13.0	100	14.0	93	9.3
PEIN	99	13.3	100	/./	101	0.0	98	1.3	94	1.2	97	12./	110	4.0	110	11.ð	100	19.0	94	/ y.l

Table 9. Recoveries (%) for five water samples acidified using 2% HCI/MeOH and analyzed by direct injection without undergoing SPE.(AVG = average; SD = standard deviation)

 Bold, italicized are IM compounds and degradation products.

 Green: 75%-125%; light green (\*): 57%-74% or 126%-135%; red (†): <57% or >135% (DoD QSM Ver5.2 for Waters—lowest LCL: MNX, 57%; highest UCL: HMX, 135%).

#### 3.5 SPE method for waters

Current USEPA Method 8330B and ERDC-EL SPE methods were modified to enable extraction and preconcentration of all 24 compounds of interest (and 2 surrogates) from environmental water samples. A series of cation-exchange, anion-exchange, reverse phase, and GAC-based SPE cartridges were evaluated using slightly modified SPE methods for environmental water extraction and preconcentration. The ability of each cartridge, individually or sequentially, to recover the 24 compounds of interest and 2 surrogates from laboratory-spiked reagent water samples was compared.

The results in Table 10 show the varying abilities of seven individual SPE cartridges to extract and preconcentrate 25 compounds (prior to addition of the second surrogate, *o*-NBA). The individual cartridges compared in this experiment were (from left to right in the table) (1) Strata X (polymeric reverse phase, 500 mg/6 mL, Phenomenex, PRP500); (2) ENVI-18 (traditional C-18, 1 g/6 mL, Supelclean, C181G); (3) Strata X-A (polymeric strong anion exchange, 200 mg/6 mL, Phenomenex, PSA200); (4) Strata X-A (polymeric strong anion exchange, 500 mg/6 mL, Phenomenex, PSA500); (5) ENVI-Carb (GAC, 250 mg/3 mL, Supelclean, GAC250); (6) Strata ABW (mixed mode strong cation exchange (4-ethylbenzene sulfonate) and weak anion exchange (propylamine), 200 mg/3 mL, Phenomenex, ABW200); and (7) a new Phenomenex GAC cartridge undergoing product testing (500 mg/6 mL, ACP500).

Overall, the reverse phase (PRP500 and C181G) and anion exchange (PSA200 and PSA500) tested were able to extract most compounds within DoD QSM Ver5.2 limits. The GAC250 was also able to extract many compounds, but fewer than the reverse phase and anion exchange. In some cases, particularly for the acidic compounds, when recoveries were low, mass balance analysis indicated that the compounds were effectively adsorbed onto cartridges but failed to elute. On the other hand, NQ tended to be the only compound that was poorly adsorbed onto any individual cartridge. Of the seven individual cartridges, ABW200 performed best for NQ with 61% recovery; however, this result proved to be irreproducible.

The ABW200 and ACP500 cartridges did not recover many analytes within accepted ranges and so were not included in further studies. Aside from ABW200, GAC250 performed best for NQ, with 30% recovery. Meanwhile, PSA500 was twice as effective as PSA200 for NQ recovery (11% versus 6%), and PRP500 was better than C181G for NQ (10% versus 5%), NTO (3% versus 1%), and PA (102% versus 11%) recovery. Furthermore, both of the PSA cartridges were able to recover all of the remaining compounds within DoD QSM Ver5.2 limits (apart from the anomalously high NG recoveries for this experiment). The PRP500 and C181G failed only to recover NTO or NTO and PA, respectively, indicating that inclusion of an anion exchange cartridge would be important for extraction of the highly acidic compounds.

On the basis of the results with individual cartridges, several double- and triple-stacked options were tested, to identify a sequential option that yielded acceptable recoveries for all compounds. The stacking order for sample loading and elution was reversed to prevent irreversible binding of analytes to incompatible SPE cartridge matrices, which was observed during evaluation of individual cartridges. In addition, two sizes of ENVI-Carb cartridges were tested, the 250 mg/3 mL (GAC250) and a larger 500 mg/6 mL (GAC500). Results for six stacked options are shown in Table 9, where loading-stage stacking order from top to bottom is indicated, including (A) PRP500-PSA500-GAC250, (B) PRP500-PSA500-GAC500, (C) C181G-PSA500-GAC250, (D) C181G-PSA500-GAC500, (E) PSA500-C181G-GAC250, and (F) PSA500-GAC250.

		Recoveries (%)											
Compound	PRP500	C181G	PSA200	PSA500	GAC250	ABW200	ACP500						
NQ	<i>10</i> †	<b>5</b> †	<b>6</b> †	<b>11</b> †	<i>30</i> †	<i>61</i> *	<b>2</b> †						
NTO (210 nm)	<b>3</b> †	<b>1</b> †	120	118	123	<b>2</b> †	<i>235</i> †						
НМХ	113	104	118	118	118	88	27						
RDX	112	115	113	113	116	102	1						
PA (210 nm)	102	<b>11</b> †	119	113	<b>1</b> †	ND†	102						
4-NP	108	112	112	110	121	<i>16</i> †	<i>65</i> *						
1,3,5-TNB	114	114	112	112	88	3†	15†						
2,4-DNP	108	106	126*	117	<b>4</b> †	<b>ND</b> †	102						
1,2-DNB (surr.)	100	102	100	101	100	99	4†						
1,3-DNB	111	114	112	112	112	21†	52†						

Table 10. Compound recoveries from laboratory-spiked reagent water samples using individual SPE cartridges for extraction/preconcentration of the 24 compounds of interest and 1,2-DNB surrogate.

			R	ecoveries	(%)		
Compound	PRP500	C181G	PSA200	PSA500	GAC250	ABW200	ACP500
2-NP	95	97	95	96	95	56†	<b>49</b> †
NB	112	114	112	113	19†	9†	ND†
3,5-DNA (254 nm)	111	114	111	114	113	110	ND†
DNAN	106	108	108	107	108	<b>1</b> †	<i>10</i> †
Tetryl	112	115	81	81	105	86	6†
NG	114	115	148†	175†	105	111	ND†
2,4,6-TNT	110	113	104	105	79	8†	28†
4-Am-2,6-DNT	117	114	112	112	118	53†	52†
2-Am-4,6-DNT	115	115	112	113	6†	13†	ND†
2,6-DNT	113	118	115	108	115	100	24†
2,4-DNT	105	98	100	119	113	2†	21†
2-NT	109	111	108	109	109	85	3†
4-NT	109	113	110	109	109	59*	31†
3-NT	109	112	110	110	109	75*	14†
PETN	126*	110	109	111	110	102	2†

Bold, italicized are IM compounds and degradation products. ND: not detected.

Green: 75%–125%; light green (\*): 57%–74% or 126%–35%; red (†): <57% or >135% (DoD QSM Ver5.2 for Waters–lowest LCL: MNX, 57%; highest UCL: HMX, 135%)

Table 11. Compound recoveries (%) from laboratory-spiked reagent water samples
using stacked SPE cartridges for extraction/preconcentration of the 24 compounds
of interest and 1,2-DNB surrogate.

			Recove	ries (%)		
	А	A B C D		Е		
	PRP500	PRP500	C181G	C181G	PSA500	F
	I	I	I	I	1	PSA500
	PSA500	PSA500	PSA500	PSA500	C181G	I
	I	I	I	I	I	GAC250
Compound	GAC250	GAC500	GAC250	GAC500	GAC250	
NQ	<b>2</b> †	<b>19</b> †	<b>48</b> †	<b>4</b> †	<b>43</b> †	<i>38</i> †
NTO (210 nm)	107	98	109	103	107	111
HMX	97	95	117	102	113	110
RDX	91	92	114	99	107	109
PA (210 nm)	94	86	96	67*	102	101

		Recoveries (%)											
	A PRP500   PSA500	B PRP500 I PSA500	C C181G   PSA500	D C181G   PSA500	E PSA500   C181G	F PSA500 I							
						GAC250							
Compound	GAC250	GAC500	GAC250	GAC500	GAC250								
4-NP	87	90	110	97	103	104							
1,3,5-TNB	93	91	113	100	106	109							
2,4-DNP	93	94	110	93	112	115							
1,2-DNB (surr.)	90	88	101	97	95	98							
1,3-DNB	90	87	113	97	106	109							
2-NP	76	76	95	85	91	91							
NB	92	90	114	100	107	109							
3,5-DNA (254 nm)	79	84	111	80	105	107							
DNAN	77	84	106	93	102	105							
Tetryl	94	88	109	96	71*	81							
NG	94	103	114	101	184†	108							
2,4,6-TNT	93	89	113	100	96	99							
4-Am-2,6-DNT	93	92	112	100	105	108							
2-Am-4,6-DNT	95	92	113	99	106	108							
2,6-DNT	85	86	115	99	109	113							
2,4-DNT	97	89	105	96	99	95							
2-NT	74*	75*	108	71*	101	105							
4-NT	78	77	109	77	102	104							
3-NT	77	76	112	74*	101	105							
PETN	89	89	110	98	101	105							

*Bold, italicized* are IM compounds and degradation products. Green: 75%–125%; light green (\*): 57%–74% or 126%–135%; red (†): <57% or >135%

(DoD QSM Ver5.2 for Waters–lowest LCL: MNX, 57%; highest UCL: HMX, 135%)

NQ recovery for each option was expected to be  $\geq$ 50%, on the basis of the recoveries for individual cartridges noted in Table 8 and the assumption of an additive relationship. While NQ recoveries for this particular experiment were not as high as expected, options B and C from Table 9 were selected for further tests. Although option F (PSA500-GAC250) also yielded higher recovery of NQ (38%), triple-stacked options that included

a reverse-phase cartridge were preferred for further studies as an added mechanism to ensure prevention of analyte breakthrough in more concentrated, larger volume or more complex matrix samples; however, this option (or PSA500-GAC500) may be acceptable in some cases to reduce method complexity. Furthermore, in cases where NQ is known or likely to be absent or not of interest, it may be acceptable to use a single PSA500 cartridge (Table 10).

Studies of the 24 compounds' and 2 surrogates' pre- and postextraction stability in laboratory-spiked reagent water samples were carried out using options B and C from Table 11. On the basis of the slightly better results for option B in the stability studies (Tables 24 and 25 in section 3.9), the PRP500-PSA500-GAC500 option was selected as the best multicartridge option for recovery of all compounds, with NQ recovery consistently approximately 50%. There was also a laboratory process—based preference for GAC500 versus GAC250 in stacked options, as the reduced surface area and volume capacity of the GAC250 (3 mL) resulted in an increased time requirement for sample loading and complicated SPE conditioning and extract elution procedures. Therefore, option B was chosen for use in cross-laboratory batch tests of five different laboratory-spiked water samples (Table 29 in section 3.10).

## 3.6 SE method for soils

Current USEPA Method 8330B and ERDC-EL SE methods were modified to enable extraction of all 24 compounds of interest (and 2 surrogates) from environmental soil samples. Preliminary SE trials indicated that traditional USEPA Method 8330B solid matrix extraction methods (that is, extraction using ACN for a single 18 h ultrasonication period) would be insufficient to extract all 24 compounds of interest, supported by previous studies (Walsh et al. 2016; Felt et al. 2016). Therefore, several different solvents, ultrasonication times, and multistage extraction procedures were investigated.

Early trials compared single 18 h extractions of laboratory-spiked samples using either MeOH or ACN, according to USEPA Method 8330B procedures, including spiking, drying, and ultrasonication. The majority of the method development was carried out using a standard soil, ASTM fat clay (CH-1). From the results of early trials, it became clear that using either ACN or MeOH alone would be insufficient to extract all compounds from soils at acceptable recovery rates. Table 12 compares compound recoveries for laboratory-spiked ASTM fat clay soil samples using either MeOH or ACN for either  $1 \times 18$  h or  $2 \times 9$  h ultrasonication periods, with no other modifications to USEPA Method 8330B procedures of drying, sonicating, and filtering. For both MeOH and ACN, splitting the ultrasonication period into two 9 h stages, rather than using a single 18 h period, improved recoveries, especially for some of the IM compounds and daughter products. For NTO, PA, 4-NP, 2,4-DNP, and DNAN, higher recoveries were observed when MeOH was the extraction solvent, while higher recoveries were observed for 2-NP and NQ with ACN. Interestingly, HMX and RDX recoveries were consistently poor in MeOH extractions, while with ACN, these two legacy compounds were recovered at near 100%. Since the majority of IM compounds and daughter products were recovered at higher rates by MeOH than ACN, and since a multi-stage ultrasonication procedure was shown to increase recovery rates overall, further experiments were modified from the  $2 \times 9h$  MeOH option, and were designed mainly to improve recoveries of NQ, NTO, HMX, and RDX.

Several two- and three-stage ultrasonication sequences using various combinations of extraction solvents (including MeOH, 50/50 MeOH/water, and water) were tested, with a summary of the results shown in Table 13. Options A–D were three-stage sequences, where stage one and two were carried out during extraction Day 1 and stage three was carried out overnight from Day 1 into Day 2. Options E-J were two-stage sequences, where stage one of option E–G was carried out during extraction Day 1, and stage two was carried out overnight from Day 1 into Day 2 like with options A–D. Meanwhile, for options H–J, extraction stage one was carried out on Day 1, and stage two was carried out on Day 2, with overnight storage at 4°C in the absence of light. Several multistage ultrasonication sequences resulted in recovery rates for all compounds within the DoD QSM Ver5.2 accepted range (64%-135%) for soils analyzed by USEPA Method 8330B, including options A, B, F, G, I, and J. Each of these options employed a combination of MeOH and water extraction solvents. Only options E and H, which consisted of sequential (6 h-14 h or 8 h-8 h, respectively) MeOH extractions, resulted in low recoveries of NQ, NTO, HMX, and RDX, as expected on the basis of previous results (Table 12). Of the six multistage options that showed acceptable recoveries for all compounds, option F or G was preferred so as to simplify the method and to enable users to perform the procedure within the time constraints of a normal workday.

	Recoveries (%)												
Compound	18 h MeOH	2 × 9 h MeOH	18 h ACN	2 × 9 h ACN									
NQ	<i>30</i> †	<b>41</b> †	<i>36</i> †	<i>50</i> †									
NTO (210 nm)	<i>22</i> †	<i>58</i> †	<b>15</b> †	<i>30</i> †									
НМХ	15†	37†	84	103									
RDX	15†	35†	78	95									
PA (210 nm)	<i>75</i> *	92	<b>64</b> *	76									
4-NP	84	104	82	93									
1,3,5-TNB	87	108	90	110									
2,4-DNP	<b>48</b> †	75*	<b>24</b> †	<b>21</b> †									
1,2-DNB (surr.)	81	103	82	101									
1,3-DNB	86	106	76	97									
2-NP	45	67*	<b>64</b> *	80									
NB	84	105	78	97									
3,5-DNA (254 nm)	93	113	87	104									
DNAN	84	105	75*	89									
Tetryl	82	102	68*	98									
NG	85	106	45†	121									
2,4,6-TNT	87	107	86	119									
4-Am-2,6-DNT	88	111	109	165†									
2-Am-4,6-DNT	88	114	120	172†									
2,6-DNT	84	99	86	114									
2,4-DNT	90	116	92	108									
2-NT	92	112	79	114									
4-NT	88	107	90	115									
3-NT	88	110	89	128*									
PETN	90	111	87	118									

Table 12. Recoveries (%) from spiked ASTM fat clay using either MeOH or ACN to perform SE for either  $1 \times 18$  h or  $2 \times 9$  h ultrasonication periods.

Bold, italicized are IM compounds and degradation products.

Green: 75%-125%; light green (\*): 64%-74% or 126%-135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

	Recoveries (%)													
	3 h-3 h-1	L4 h			6 h-14 h			8 h-8 h						
Compound	A 1. MeOH 2. MeOH 3. H <sub>2</sub> O	B 1. MeO 2. H <sub>2</sub> O 3. H <sub>2</sub> O	C 1. MeOH 2. MeOH 3. MeOH/H <sub>2</sub> O	D 1. MeOH 2. MeOH/H <sub>2</sub> O 3. H <sub>2</sub> O	E 1. MeOH 2. MeOH	F 1. MeOH 2. MeOH/H <sub>2</sub> O	G 1. MeOH 2. H <sub>2</sub> O	H 1. MeOH 2. MeO	I 1. MeOH 2. H <sub>2</sub> O	J 1. MeOH 2. MeOH/H <sub>2</sub> O				
NQ	<i>69</i> *	88	54†	69*	<i>53</i> †	69*	<b>79</b>	52†	74*	66*				
NTO (210nm)	77	94	91	107	<i>37</i> †	<b>9</b> 7	86	<b>36</b> †	81	82				
HMX	65*	92	68*	92	31†	72*	71*	32†	69*	67*				
RDX	69*	99	72*	96	32†	76	77	33†	74*	71*				
PA (210nm)	104	126*	107	125*	116	121	123	110	114	115				
4-NP	103	119	107	125*	120	122	124	116	119	119				
1,3,5-TNB	105	110	110	124	125	122	115	121	110	121				
2,4-DNP	<i>92</i>	116	95	118	81	110	111	<i>79</i>	104	102				
1,2-DNB (surrogate)	104	117	106	124	120	121	122	115	117	118				
1,3-DNB	105	120	108	128*	124	125	125	120	121	123				
2-NP	77	107	79	109	69*	103	104	69*	<i>95</i>	93				
NB	104	116	104	124	119	118	116	117	112	114				
3,5-DNA (254nm)	99	113	99	116	110	111	112	107	107	108				
DNAN	104	115	112	127*	128*	120	121	118	115	120				
Tetryl	102	114	103	123	119	117	117	115	111	119				
NG	94	102	86	105	98	99	101	98	97	98				
2,4,6-TNT	99	109	100	119	117	118	113	113	110	114				
4-Am-2,6-DNT	106	119	113	130*	122	123	122	119	118	120				
2-Am-4,6-DNT	108	123	116	137†	131*	132*	130*	127*	126*	133*				
2,6-DNT	110	116	105	124	117	117	124	133*	121	117				
2,4-DNT	104	119	111	128*	126*	126*	123	117	119	124				
2-NT	105	120	108	127*	127*	127*	125	123	120	126*				

Table 13. Recoveries (%) from spiked ASTM fat clay using several multi-stage SE sequences with ultrasonication for either 3 h–3 h–4 h, 6 h–14 h, or 8 h–8 h with various solvent combinations.

	Recoverie	Recoveries (%)											
	3 h-3 h-14 h							8 h-8 h	8 h-8 h				
Compound	A 1. MeOH 2. MeOH 3. H <sub>2</sub> O	B 1. MeO 2. H <sub>2</sub> O 3. H <sub>2</sub> O	C 1. MeOH 2. MeOH 3. MeOH/H <sub>2</sub> O	D 1. MeOH 2. MeOH/H <sub>2</sub> O 3. H <sub>2</sub> O	E 1. MeOH 2. MeOH	F 1. MeOH 2. MeOH/H <sub>2</sub> O	G 1. MeOH 2. H <sub>2</sub> O	H 1. MeOH 2. MeO	I 1. MeOH 2. H <sub>2</sub> O	J 1. MeOH 2. MeOH/H <sub>2</sub> O			
4-NT	105	116	108	124	126*	127*	124	122	119	125			
3-NT	103	116	110	126*	122	123	121	118	117	118			
PETN	113	132*	109	128*	130*	131*	135*	125*	132*	135*			

*Bold, italicized* are IM compounds and degradation products. Green: 75%–125%; light green (\*): 64%–74% or 126%–135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

Overall, the results shown in Table 13 indicated that water (a more polar solvent than either ACN or MeOH) effectively improved recoveries of NQ, NTO, HMX, and RDX from ASTM fat clay spiked samples and also aided recovery of some other compounds, including 2,4-DNP and 2-NP. While NQ and NTO are highly water soluble and higher recoveries due to extracting with MeOH/water or water were expected for them, the superior ability of water versus MeOH to extract HMX and RDX was not predicted. However, this unexpected result may be because of the observed ability of the water physically to disperse the soil much better than the MeOH, possibly preventing the entrapment of the hydrophobic compounds and resulting in higher observed recoveries.

Because of the unexpected nature of the results when using water as the extraction solvent or cosolvent with MeOH, a similar experiment was carried out to confirm the effects (data not shown). In addition, a comparison of 18 h ultrasonic extractions using either ACN, MeOH, or 50/50 MeOH/water in several different soil types was performed. Six soils with varying physicochemical characteristics were used, described in Table 14, including five field soils (Aberdeen, Jefferson, Memphis, Riley, and Yuma), and the ASTM fat clay used previously. Soil classification is provided, including gravel, sand, and fines percentages. Chemical properties include pH, cation-exchange capacity (CEC), anion-exchange capacity (AEC), total organic carbon (TOC), and percent moisture. The results shown in Table 15 indicated that extraction of the compounds of interest depended on compound chemical properties and soil characteristics and reinforced previous results that showed that extraction using any single solvent or cosolvent mixture would not be sufficient to extract the 24 compounds (and 1,2-DNB surrogate) from any of the six tested soils. (For the ASTM soil, a water-only option was also tested, with mixed results.)

Because differences were observed for different soil types, further SE method development trials were carried out using all six soils. Using the results from the ASTM fat clay trials (Table 13), two two-stage and two three-stage sequential ultrasonication options were tested (Table 16). All four options were successful for each soil type. Only three outliers occurred, for the Yuma sand using option C (3 h MeOH–3 h water–14 h water), which were 1%–3% higher than the accepted 135% (DoD QSM Ver5.2). Thus, this subsequent round of trials confirmed that including

water as an extraction solvent resulted in improved recoveries for NQ, NTO, HMX, and RDX across a broad range of soil types.

Overall, it was shown that a two-stage SE procedure including water as an extraction solvent (Table 16, option A and B) was necessary and sufficient to result in recovery of all compounds of interest (and two surrogates) within current accepted limits (64%–135%) for all soils tested. Results for further tests carried out using option A (for the compounds of interest and two surrogates) appear in sections 3.9 and 3.10. Option A was selected over option B so that both stages in the extraction procedure would include organic extraction solvent (MeOH), which minimized deviation from current USEPA Method 8330B procedures. A three-stage extraction sequence proved unnecessary but may be employed, if desired. For cases in which NQ, NTO, HMX, and RDX are known to be absent or are not of interest in soils similar to those tested, an extraction procedure that includes water as an extraction solvent may not be necessary. In such cases, a one-step ultrasonication may be sufficient, and a two-step procedure may not enhance recovery at a statistically significant or costeffective rate.

						- p		8			
Name	Collection Site	Region	Classification	Gravel (%)	Sand (%)	Fines (%)	рН	CEC (meq Na / 100g)	AEC (meq S / 100g)	TOC (mg/kg)	Moisture (%)
Aberdeen	Aberdeen Proving Ground (Aberdeen, MD)	Mid Atlantic, Southeast	Clay (CL) with sand	2.9	25.4	71.7	7.12	20.6	51.2	6,700	2.1
Jefferson	Jefferson Proving Ground (Madison, IN)	Great Lakes, Midwest	Clay (CL) with sand	0.0	19.7	80.3	4.62	26.2	60.9	1,400	2.1
Memphis	Memphis, TN	Mid-South	Silt (ML)	0.0	0.5	99.5	7.56	11.8	45.3	610	1.1
Riley	Ft. Riley (Riley, KS)	Central Plains, Midwest	Clay; trace of sand	0.0	3.3	96.7	5.96	32.9	53.0	15,000	1.4
Yuma	Yuma Proving Ground (Yuma, AZ)	Pacific, Southwest	Sand with gravel	35	53.2	11.8	6.98	14.7	47.9	ND*	1.6
ASTM Fat Clay (CH-1)	(ASTM-ISR Refere	nce Soils Program)	Clay (CH)	0.0	0.7	99.3	7.45	24.5	24.4	3,600	5.9

Table 14. Characteristic properties for six soils used in SE method development and testing for soils.

\*ND: not detected (Reporting limit: 250 mg/kg)

		Recoveries (%)										
		Aberd	een		Jeffer	son		Memp	ohis			
Compound	ACN	MeOH	MeOH/H <sub>2</sub> O	ACN	MeOH	MeOH/H <sub>2</sub> O	ACN	MeOH	MeOH/H <sub>2</sub> O			
NQ	<i>67</i> *	<i>72</i> *	108	<i>67</i> *	81	106	<i>38</i> †	91	123			
NTO (210 nm)	22	14	88	22	8	108	24	50	101			
НМХ	109	40	96	112	39	96	107	42†	87			
RDX	110	45	108	103	49	103	105	48	97			
PA (210 nm)	58	98	113	11	100	120	84	108	<i>128</i> *			
4-NP	98	99	105	100	109	97	94	103	104			
1,3,5-TNB	109	99	85	106	104	84	104	100	55			
2,4-DNP	21	82	105	60	51	105	15	85	103			
1,2-DNB (surr.)	95	87	87	92	96	84	85	89	91			
1,3-DNB	92	87	88	92	98	85	91	90	89			
2-NP	43	33	45	33	44	6	11	15	38			
NB	108	105	100	97	103	99	105	104	92			
3,5-DNA (254 nm)	15	11	15	20	25	ND	63	4	24			
DNAN	102	103	99	103	108	93	104	98	94			
Tetryl	99	94	72*	98	102	31	104	49	97			
NG	97	92	97	64*	90	86	83	90	86			
2,4,6-TNT	106	102	102	101	108	115	96	118	88			
4-Am-2,6-DNT	108	108	105	97	102	102	102	106	92			
2-Am-4,6-DNT	111	107	101	100	103	101	109	104	95			
2,6-DNT	96	88	86	93	89	80	85	88	89			
2,4-DNT	96	87	117	84	94	83	82	91	86			
2-NT	51	29	33	43	45	4	16	15	43			
4-NT	61	56	58	69*	71*	24	51	44	67*			
3-NT	58	44	48	54	61	12	20	30	55			
PETN	112	110	103	106	111	99	97	106	103			

Table 15. Recoveries (%) from six laboratory-spiked soils using an 18 h ultrasonication period with ACN, MeOH, 50/50 MeOH/water, or water.

*Bold, italicized* are IM compounds and degradation products. ND: not detected. Green: 75%–125%; light green (\*): 64%–74% or 126%–135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

	Recoveries (%)										
		Riley			Yuma			ASTM	Fat Clay		
Compound	ACN	МеОН	MeOH/ H <sub>2</sub> O	ACN	MeOH	MeOH∕ H₂O	ACN	МеОН	MeOH∕ H₂O	H₂O	
NQ	<i>35</i> †	65	100	<i>52</i> †	79	84	<i>62</i> †	66	103	91	
NTO (210 nm)	<i>23</i> †	<i>21</i> †	64	<b>24</b> †	<i>43</i> †	<i>9</i> 5	<b>8</b> †	<b>17</b> †	107	105	
НМХ	94	26†	79	105	34†	84	92	19†	75	70	
RDX	97	32†	88	99	33†	94	92	23†	86	99	
PA (210 nm)	<i>38</i> †	99	108	92	106	130	96	109	122	129	
4-NP	86	101	98	99	104	97	97	123	114	120	
1,3,5-TNB	109	103	55†	99	105	86	93	121	84	37†	
2,4-DNP	<b>42</b> †	<i>51</i> †	99	<i>23</i> †	<i>56</i> †	97	<i>32</i> †	<b>51</b> †	92	96	
1,2-DNB (surr.)	94	93	93	94	87	84	95	113	92	101	
1,3-DNB	97	97	89	91	90	83	91	117	96	87	
2-NP	<i>56</i> †	65	69	<i>36</i> †	<i>23</i> †	<b>37</b> †	<i>59</i> †	<i>53</i> †	ND†	77	
NB	85	86	76	101	105	95	92	117	101	71	
3,5-DNA (254 nm)	57†	54†	59†	11†	8†	9†	45†	52†	ND†	33†	
DNAN	97	101	86	99	101	94	94	113	96	55	
Tetryl	97	93	79	107	70	31†	17†	65	66	35†	
NG	84	100	92	101	95	85	104	124	95	109	
2,4,6-TNT	95	105	84	95	116	112	109	138†	106	70	
4-Am-2,6-DNT	80	85	78	103	106	113	18†	121	110	6†	
2-Am-4,6-DNT	91	93	81	103	104	95	93	117	101	68	
2,6-DNT	87	106	97	71	91	84	87	138†	81	107	
2,4-DNT	96	92	84	89	90	84	95	118	94	69	
2-NT	73	69	68	35†	20†	20†	55†	64	ND†	43†	
4-NT	94	85	78	62†	45†	44†	62†	80	ND†	58†	
3-NT	76	79	74	38†	32†	32†	67	71	ND†	50†	
PETN	97	108	99	100	105	95	100	122	102	31†	

Table 15 (continued).

*Bold, italicized* are IM compounds and degradation products. ND: not detected. Green: 75%–125%; light green (\*): 64%–74% or 126%–135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

	Recoveries (%)															
	Aberdee	n							Jeffersor							
	A		В		С		D		A		В		С		D	
Compound	AVG ( <i>n</i> = 3)	SD														
NQ	<u>98</u>	10	102	5	120	3	117	8	106	10	104	4	119	1	109	6
NTO (210nm)	81	10	80	3	96	2	86	2	83	8	88	1	113	2	96	1
HMX	87	5	80	2	107	1	91	3	83	3	78	2	100	3	87	3
RDX	93	6	88	4	113	2	103	1	94	3	89	2	110	2	99	1
PA (210nm)	105	7	106	8	120	8	117	4	109	5	101	4	114	8	115	9
4-NP	115	8	110	5	123	7	126*	2	121	4	113	5	122	6	124	7
1,3,5-TNB	112	8	100	2	114	6	120	6	111	6	96	3	132*	3	119	2
2,4-DNP	104	7	<u>99</u>	<1	120	3	114	4	<i>95</i>	4	88	3	107	6	96	2
1,2-DNB (surrogate)	123	6	118	4	125*	6	128*	1	121	10	117	6	127*	3	123	8
1,3-DNB	119	9	112	4	125*	8	128*	4	123	5	115	4	124	3	126*	3
2-NP	103	11	96	3	108	6	107	3	100	4	95	3	105	2	105	1
NB	119	9	111	4	125*	8	128*	4	120	6	110	3	119	3	122	3
3,5-DNA (254nm)	89	7	86	4	92	6	96	3	101	6	96	3	101	3	102	5
DNAN	117	10	111	5	123	10	126*	6	121	6	109	2	115	3	124	3
Tetryl	103	10	99	7	113	4	115	3	115	6	109	7	120	5	121	7
NG	113	16	105	10	115	3	119	10	107	11	105	12	112	10	115	14
2,4,6-TNT	116	8	106	6	120	7	123	5	117	1	105	3	112	3	120	5
4-Am-2,6-DNT	124	10	112	4	126*	9	129*	3	120	6	112	3	121	3	123	2
2-Am-4,6-DNT	120	9	111	4	125*	8	128*	3	123	5	114	3	123	3	126*	3
2,6-DNT	116	5	107	4	118	11	124	6	120	4	112	3	119	3	119	3
2,4-DNT	118	6	110	4	125*	9	126*	6	123	8	111	6	117	7	124	6
2-NT	103	7	96	1	103	6	107	4	108	9	101	3	108	4	111	4
4-NT	110	6	98	5	110	9	118	5	115	10	106	2	113	4	114	5
3-NT	103	7	98	3	107	6	112	4	110	7	103	4	110	5	112	5

Table 16. Recoveries (%) from six laboratory-spiked soils using four different two- or three-stage ultrasonication sequences. (A) 6 h MeOH–14-h MeOH/water; (B) 6 h MeOH–14 h water; (C) 3 h MeOH–3 h water-14 h water; (D) 3 h MeOH–3 h MeOH–14 h water. (AVG = average; SD = standard deviation)

		Recoveri	es (%)														
		Aberdee	Aberdeen Jefferson														
	A B C D A B C D																
		AVG		AVG		AVG		AVG		AVG		AVG		AVG		AVG	
Compound		( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD
	PETN	114	8	109	3	124	8	123	3	121	8	114	3	125*	4	124	5

*Bold, italicized* are IM compounds and degradation products. ND: not detected. Green: 75%–125%; light green (\*): 64%–74% or 126%–135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

# Table 16 (continued). (AVG = average; SD = standard deviation)

	Recoveri	ies (%)														
	Memphi	S							Riley							
	A		В		С		D		Α		В		С		D	
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NQ	107	9	111	1	118	9	115	9	90	6	<u>98</u>	9	111	2	109	5
NTO (210nm)	115	15	106	2	125*	2	119	2	100	8	112	3	128*	2	119	4
HMX	98	7	90	1	109	3	103	1	67*	2	66*	4	84	2	72*	3
RDX	99	4	94	<1	114	3	106	2	81	1	78	1	101	3	88	1
PA (210nm)	121	6	111	5	126*	7	121	2	111	5	103	4	124	3	124	2
4-NP	120	4	112	4	126*	5	123	2	115	4	103	3	118	5	120	<1
1,3,5-TNB	122	4	108	<1	121	4	125*	4	113	4	96	1	104	2	120	<1
2,4-DNP	117	7	106	2	123	6	118	1	95	4	89	3	108	1	97	3
1,2-DNB (surrogate)	121	7	118	6	126*	6	127*	4	121	5	111	8	126*	5	124	4
1,3-DNB	123	3	115	2	127*	5	126*	3	121	4	108	1	120	4	125*	1
2-NP	91	4	88	1	101	5	96	4	106	4	96	2	105	5	108	2
NB	124	3	116	1	128*	4	127*	3	110	3	93	<1	103	3	111	<1
3,5-DNA (254nm)	86	4	81	1	88	6	85	3	108	3	100	2	108	5	112	2
DNAN	122	3	112	2	126*	5	125*	1	119	3	102	1	115	3	123	1
Tetryl	86	14	82	5	102	4	97	7	112	8	100	4	112	5	117	5

′G		
= 3)	SD	
118	13	

	Recoveri	es (%)														
	Memphi	S							Riley							
	Α		В		С		D		Α		В		С		D	
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NG	118	11	113	9	117	13	118	8	111	15	103	11	111	10	118	13
2,4,6-TNT	130*	1	119	1	129*	4	129*	4	117	7	100	2	111	1	122	2
4-Am-2,6-DNT	127*	3	119	1	130*	4	129*	2	109	2	93	1	103	2	110	2
2-Am-4,6-DNT	125*	3	117	1	128*	4	127*	2	116	3	99	<1	109	3	118	1
2,6-DNT	121	4	109	2	124	2	121	4	117	10	100	2	113	6	125*	6
2,4-DNT	123	4	115	3	125*	5	126*	2	119	2	106	2	115	6	126*	4
2-NT	99	4	91	2	101	5	100	6	112	5	103	1	112	5	119	2
4-NT	106	4	99	2	110	2	108	4	114	4	104	2	112	3	121	1
3-NT	101	5	93	<1	103	5	101	3	114	3	104	2	112	7	116	4
PETN	115	5	109	7	120	7	117	1	121	7	112	3	122	5	125*	5

*Bold, italicized* are IM compounds and degradation products. ND: not detected. Green: 75%–125%; light green (\*): 64%–74% or 126%–135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

## Table 16 (continued). (AVG = average; SD = standard deviation)

	Recoveri	es (%)														
	Yuma								ASTM Fa	t Clay						
	Α		В		C		D		Α		В		С		D	
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NQ	109	7	106	2	122	4	115	4	101	7	100	5	116	3	113	5
NTO (210nm)	103	3	88	2	111	5	<u>98</u>	1	90	1	92	2	118	1	101	3
HMX	86	4	84	3	105	2	93	2	66*	1	65*	1	92	2	75*	2
RDX	92	3	89	2	111	2	98	<1	72*	1	72*	1	101	1	82	2
PA (210nm)	117	6	107	4	127*	6	123	3	106	5	103	7	119	5	120	2
4-NP	119	5	110	3	132*	10	125*	2	115	6	112	3	124	1	125*	1

	Recoverie	coveries (%)														
	Yuma								ASTM Fa	t Clay						
	A		В		С		D		A		В		С		D	
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
1,3,5-TNB	120	5	111	1	130*	7	127*	2	114	4	103	1	110	1	124	4
2,4-DNP	102	4	<u>98</u>	1	117	3	105	1	92	2	<i>91</i>	3	110	1	100	3
1,2-DNB (surrogate)	122	7	116	7	134*	16	124	5	121	10	115	7	125*	3	125*	1
1,3-DNB	121	4	114	1	134*	9	127*	2	119	4	115	2	125*	1	129*	5
2-NP	85	5	81	4	<u>99</u>	6	<i>93</i>	1	89	2	88	2	102	3	93	5
NB	122	4	116	1	135*	9	128*	2	118	5	113	2	122	<0	127*	5
3,5-DNA (254nm)	75*	3	72*	4	84	9	79	2	96	5	93	1	101	1	103	3
DNAN	120	4	114	2	133*	7	<i>129</i> *	1	117	4	110	2	119	2	127*	4
Tetryl	93	11	93	5	118	6	113	6	99	4	98	2	111	3	111	1
NG	114	11	106	8	125*	17	118	13	106	7	108	9	116	10	114	5
2,4,6-TNT	128*	5	119	2	138†	12	130*	5	116	2	112	4	119	2	124	3
4-Am-2,6-DNT	126*	4	118	1	137†	9	130*	2	121	4	117	4	126*	1	127*	2
2-Am-4,6-DNT	123	4	117	1	136†	9	129*	2	119	4	113	2	122	1	126*	2
2,6-DNT	116	11	109	6	132*	4	121	3	118	4	109	6	123	5	124	4
2,4-DNT	123	2	115	2	135*	8	127*	1	114	4	112	3	122	2	125*	3
2-NT	88	5	83	3	98	9	92	2	105	4	101	1	110	4	111	3
4-NT	97	2	93	3	110	10	104	1	110	6	105	3	114	2	117	5
3-NT	94	4	88	4	101	10	97	1	103	6	101	2	108	<1	111	1
PETN	120	7	113	1	130*	11	123	2	117	5	115	3	123	1	126*	3

Bold, italicized are IM compounds and degradation products.

Green: 75%–125%; light green (\*): 64%–74% or 126%–135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids–lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

#### 3.7 SE method for tissues

Current USEPA Method 8330B and ERDC-EL SE methods were modified to enable extraction of all 24 compounds of interest (and 2 surrogates) from environmental tissue samples. A single ultrasonication period using MeOH was predicted to be sufficient for extraction of the 26 compounds, as the tissues used during method development experiments were not previously lyophilized, and, therefore, contained a significant portion of water. Initial tests led to the adoption of a single-step, 18 h MeOH extraction. Additionally, an extract interference reduction method was needed (adapted from Larson et al. 1999), as all of the tissues (fathead minnow [*Pimephales promelas*], polychaete worm [*Alitta virens*], earthworm [*Eisenia fetida*], and ryegrass [*Lolium perenne*]) suffered from varying degrees of analytical interference because of coeluting nontarget extracted components.

Compound recoveries prior to application of the interference reduction procedure are shown in Table 17 for all four tissue types. The DoD QSM Ver5.2 UCL and LCL accepted limits for solid (soil) matrices analyzed by USEPA Method 8330B were used as a benchmark (64%–135%), as they were during the soil SE method development described previously, since limits have not been established specifically for tissues matrices. A single ultrasonication period of 18 h using MeOH was shown to be sufficient to extract the majority of compounds (17 from earthworm, 22 from polychaete worm, 24 from minnow, and 23 from ryegrass) within the accepted range for solids according to DoD QSM Ver5.2. Tetryl was the only compound generally recovered poorly from all three tested animal tissue types, at <5%. The scientific literature highlights the matrix- and temperature-dependent stability of Tetryl (Harvey et al. 1992; Jenkins 1994; Jenkins and Walsh 1994; Boopathy 2000; Douglas et al. 2009), which was consistent with current and previous results obtained at ERDC-EL. For this reason, it may be beneficial to investigate reduced ultrasonication periods to determine whether the full 18 h period is necessary for efficient recovery of the other compounds of interest or whether Tetryl recovery could be improved by having a shorter time period during which degradation could occur. While no compound stability studies were performed in tissue matrices for the current work, results for water and soil matrices are shown and discussed in section 3.9, with results for Tetryl consistent with referenced literature.

	Recoverie	es (%)						
	Fathead I	minnow	Earthwor	m	Polychaet	te worm	Ryegrass	
Compound	AVG ( <i>n</i> = 4)	SD	AVG ( <i>n</i> = 2)	SD	AVG ( <i>n</i> = 5)	SD	AVG ( <i>n</i> = 3)	SD
NQ	<i>73</i> *	18.0	59†	5.1	<i>140</i> †	134.2	99	6.0
NTO (210 nm)	<i>1,112</i> †	325.3	668†	133.7	<b>395</b> †	122.7	<i>12</i> †	1.4
o-NBA (surr.)	85	6.7	54†	35.3	119	14.5	24†	1.8
HMX	76	2.5	89	27.6	72*	2.9	81	7.2
RDX	73*	1.1	84	18.8	74*	1.4	86	9.9
PA (210 nm)	77	20.7	<b>419</b> †	335.8	<i>60</i> †	4.5	85	9.1
4-NP	102	15.4	110	34.4	79	1.0	89	5.6
1,3,5-TNB	72*	4.0	234†	192.7	74*	2.4	87	6.6
2,4-DNP	79	1.7	96	21.6	77	2.1	110	6.9
1,2-DNB (surr.)	82	0.6	83	2.4	82	0.8	86	7.5
1,3-DNB	74*	0.6	76	2.9	74*	0.8	87	7.0
2-NP	<i>68</i> *	1.5	<i>69</i> *	2.2	<i>68</i> *	1.8	86	6.7
NB	75*	0.8	79	1.3	74*	1.7	87	7.6
3,5-DNA (254 nm)	83	2.2	85	2.6	83	1.0	90	7.0
DNAN	<i>70</i> *	6.8	<i>73</i> *	3.8	<i>70</i> *	2.5	90	6.4
Tetryl	3†	3.2	2†	2.1	4†	1.8	51†	16.5
NG	73*	3.7	74*	0.1	72*	0.6	85	5.5
2,4,6-TNT	73*	3.9	77	1.3	74*	1.3	87	6.7
4-Am-2,6-DNT	75	3.4	78	0.8	73*	0.8	93	8.2
2-Am-4,6-DNT	73*	2.0	77	1.9	74*	1.3	94	7.3
2,6-DNT	91	17.3	85	5.0	82	2.4	88	7.2
2,4-DNT	74*	7.3	72*	4.3	71*	2.4	87	7.4
2-NT	79	1.0	82	0.1	76	2.3	87	7.1
4-NT	76	1.9	80	3.6	70*	3.7	89	5.4
3-NT	79	2.0	80	3.5	80	3.6	89	6.8
PETN	72*	3.2	73*	2.7	83	10.2	92	5.1

Table 17. Recoveries (%) from four laboratory-spiked tissues using one 18 h ultrasonic extraction with MeOH, prior to undergoing chromatographic treatment to reduce analytical interference. (AVG = average; SD = standard deviation)

Bold, italicized are IM compounds and degradation products.

Green: 75%-125%; light green (\*): 64%-74% or 126%-135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

UV chromatograms of both unspiked and spiked tissue samples (not shown) revealed the presence of nontarget matrix components for each tissue type, with the majority occurring toward the beginning of the chromatographic separation (0–16 min), indicating more polar interferents. The presence of these components resulted in observed positive interferences for the more polar compounds of interest (mainly NQ, NTO, and PA), with the worst interference observed for earthworm tissue. Matrix-based analytical interferences occurred in three chromatographic regions for earthworm tissue, which resulted in large positive interference and very high recovery rates for NTO, PA, and 1,3,5-TNB. In addition, NQ and o-NBA surrogate were recovered but not quite within current QSM accepted limits. Compound recovery from polychaete worm tissue was similar to earthworm tissue. However, positive interference was observed for NQ, PA recovery was slightly low, and 1,3,5-TNB analysis was not affected by interference. Positive interference was still observed for NTO but to a lesser extent. For the fathead minnows, positive interference again inhibited accurate determination of NTO. No positive chromatographic interference was observed for ryegrass tissue; however, neither NTO (12%) nor o-NBA (24%) was recovered within current accepted QSM limits. The analytical interferences could have been due to coelution or coextraction of nontarget compounds that interfered with NTO or o-NBA detection during the HPLC analytical method, arising from pH-related or other interactions. Therefore, it was predicted that subjecting plant tissue samples to an appropriate interference reduction procedure might eliminate these interferences as well. Interestingly, Tetryl recovery was approximately 50% from plant tissue, whereas it was <5% for all animal tissue types.

# 3.8 Chromatographic interference reduction method for tissue extracts

Because of the chromatographic results obtained during tissue extraction method development, an interference reduction procedure for tissue matrix interferences was necessary to obtain accurate and reliable results for tissue extract analysis, especially extracts known or suspected of containing the more polar IM compounds like NQ, NTO, and PA. Furthermore, as mentioned briefly in section 2.8, at cold temperatures, some tissue MeOH extracts developed a precipitate. This precipitation issue also occurred for certain extracts that had been prepared for analysis to a ratio of 50/50 MeOH/water. The majority of the precipitation issue was observed for earthworm extracts, to a lesser extent for polychaete worm and minnow extracts, and not at all for ryegrass extracts. In the case of precipitation, samples underwent additional filtering at the point of collection or following dilution with reagent water in preparation for analysis. Additional filtering was performed using either a second 0.45  $\mu$ m PTFE hydrophobic or a 0.20  $\mu$ m Anotop 10 inorganic membrane disk filter. When HPLC-UV data for samples that had undergone additional filtering showed persistent chromatographic interference, further experiments were carried out to determine an effective interference reduction method.

As an alternative to additional filtration alone, either 0.1% or 1% TFA (final v/v) was added to MeOH extracts or 50/50 MeOH/water-prepared analytical samples to encourage "complete" precipitation, followed by additional filtering (if needed). The addition of acid yielded no further benefit than a second filtration step alone (data not shown). Therefore, a chromatographic interference reduction method was adapted from Larson et al (1999). Some changes were made to the method described in the original publication. The original ACN and basic alumina were changed to MeOH and neutral alumina, respectively. Instead of using a combination of alumina and florisil, either alumina or florisil was used initially to enable determination of the responsible component for the effect on HPLC-UV chromatography for each compound of interest. Initial trials of the adapted chromatographic method were performed only on polychaete worms (Table 18). For the experiment shown, treatment with alumina removed all of the analytical interferences, and even resulted in improved recoveries for many other compounds; however, it also caused a reduction in the detected o-NBA surrogate to 39%. Meanwhile, treatment with florisil proved similar; however, both NTO and o-NBA recoveries were reduced to levels below the accepted DoD QSM Ver5.2 lower limit of 64%.

Because of the preliminary success with the alumina and florisil methods, the same tests were repeated, using all four tissue matrices, shown in Table 19, with similar results. Further interference reduction methods were investigated, including mixed combinations of neutral alumina and florisil as well as basic alumina and silica. For these tests, MeOH extracts were obtained via 18 h ultrasonication of unspiked tissue. Post-extraction spiked MeOH extracts underwent chromatographic treatment as described in section 2.8. Compound recoveries (%) for each tissue type using each of the five chromatographic methods are shown in Tables 18–23.

	Polychaet	te worm—r	ecoveries (	(%)
	Alumina d	column	Florisil co	lumn
Compound	AVG ( <i>n</i> = 2)	SD	AVG ( <i>n</i> = 2)	SD
NQ	117	5.0	<i>129</i> *	13.7
NTO (210 nm)	93	16.2	<i>26</i> †	4.2
o-NBA (surrogate)	39†	1.4	24†	27.9
НМХ	82	2.0	90	3.3
RDX	109	5.7	95	16.3
PA (210 nm)	88	2.7	90	3.9
4-NP	92	2.1	97	4.1
1,3,5-TNB	90	0.7	96	1.4
2,4-DNP	95	0.5	96	4.6
1,2-DNB (surrogate)	91	2.3	96	2.4
1,3-DNB	91	1.1	96	2.1
2-NP	<i>70</i> *	3.4	<b>71</b> *	2.6
NB	88	1.4	94	2.7
3,5-DNA (254 nm)	97	2.7	97	3.4
DNAN	78	1.5	85	0.6
Tetryl	4†	0.3	5†	1.6
NG	95	3.6	96	9.1
2,4,6-TNT	92	0.1	97	2.1
4-Am-2,6-DNT	87	0.7	91	4.3
2-Am-4,6-DNT	90	0.6	92	4.9
2,6-DNT	88	1.3	95	1.4
2,4-DNT	88	0.2	90	2.5
2-NT	98	4.1	94	13.9
4-NT	89	2.9	91	1.6
3-NT	98	3.6	103	2.6
PETN	84	4.5	85	1.4

Table 18. Recoveries (%) from laboratory-spiked polychaete worm tissue MeOH extracts following either neutral alumina or florisil chromatographic treatment for interference reduction. (AVG = average; SD = standard deviation)

Bold, italicized are IM compounds and degradation products.

Green: 75%-125%; light green (\*): 64%-74% or 126%-135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)
	Recoveries (%)											
	Fathead	minnow					Earthwor	m				
	No chromato interferen reductior	ographic nce n	Alumina	column	Florisil co	blumn	No chromato interferen reductior	ographic nce 1	Alumina	column	Florisil co	blumn
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NQ	<b>398</b> †	33.6	238†	20.3	<i>280</i> †	30.1	88	4.0	<i>203</i> †	77.8	<i>162</i> †	21.9
NTO (210nm)	<b>4,363</b> †	422.6	<i>4,441</i> †	81.1	<b>5,659</b> †	723.5	<i>3,095</i> †	346.1	3,532†	1,845.6	<b>5,994</b> †	1,205.5
NTO (315nm)	<i>21</i> †	<b>8.</b> 7	114	10.6	<i>140</i> †	8.0	<i>16</i> †	3.0	129*	10.1	<i>162</i> †	5.1
o-NBA (surrogate)	93	4.3	82	4.3	98	4.0	52†	3.8	79	6.8	82	1.8
HMX	87	2.6	88	5.7	96	1.6	102	5.0	60†	6.0	54†	2.1
RDX	88	2.3	93	2.5	96	5.2	94	6.3	108	4.4	88	3.2
PA (210nm)	7 <b>0</b> *	7.4	78	3.5	81	6.6	<i>719</i> †	36.8	<i>165</i> †	135.6	<b>360</b> †	44.2
PA (315nm)	<i>71</i> *	1.7	68*	2.4	<i>73</i> *	2.3	<i>51</i> †	2.0	<b>0</b> †	0.0	<i>7</i> †	0.4
4-NP	101	1.9	97	3.1	104	4.5	57†	2.4	<b>48</b> †	6.3	<b>48</b> †	3.5
1,3,5-TNB	52†	0.6	64*	6.1	54†	1.5	53†	1.5	59†	2.9	57†	3.4
2,4-DNP	66*	1.9	<i>62</i> †	3.0	73*	4.2	<b>63</b> †	5.7	<i>71</i> *	4.9	67*	5.0
1,2-DNB (surrogate)	80	0.1	81	3.7	82	4.3	76	0.6	78	6.6	84	6.3
1,3-DNB	89	1.4	90	3.5	94	1.4	76	1.3	79	0.9	78	2.4
2-NP	92	1.6	80	2.8	87	1.6	73*	1.9	67*	1.6	72*	4.1
NB	78	1.2	75*	6.4	77	0.7	73*	1.2	77	3.8	77	3.7
3,5-DNA (210nm)	92	1.7	92	3.8	99	7.3	79	4.6	82	1.3	84	7.4
3,5-DNA (254nm)	97	2.2	104	2.3	104	3.5	85	2.0	89	3.9	93	5.5
DNAN	107	3.2	107	3.9	109	6.1	76	2.4	89	2.9	92	2.2
Tetryl	2†	2.8	0†	0.0	0†	0.0	7†	6.4	11†	2.1	12†	2.3
NG	42†	3.5	36†	7.2	44†	4.5	39†	4.6	41†	2.8	45†	2.7
2,4,6-TNT	78	2.1	90	3.4	87	4.6	71*	2.7	87	1.6	86	2.8

 Table 19. Recoveries (%) from laboratory-spiked tissues MeOH extract, following no interference reduction treatment, or treatment with either

 neutral alumina or florisil. (AVG = average; SD = standard deviation)

	Recoverie	es (%)										
	Fathead	minnow					Earthwor	m				
	No chromato interferei reductior	ographic nce 1	Alumina column		Florisil co	Florisil column		ographic nce 1	Alumina	column	Florisil co	olumn
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
4-Am-2,6-DNT	67*	0.5	66*	3.3	64*	5.5	64†	2.3	69*	4.2	68*	3.0
2-Am-4,6-DNT	83	2.2	82	3.7	86	1.8	73*	2.5	79	2.8	77	9.5
2,6-DNT	88	6.0	96	4.3	100	0.4	72*	4.3	74*	2.7	72*	2.7
2,4-DNT	88	4.8	88	1.4	94	3.0	72*	2.7	75*	1.3	76	2.3
2-NT	91	3.2	91	5.0	97	6.8	77	5.7	75*	3.1	94	9.1
4-NT	86	1.2	78	7.5	88	3.4	71*	2.9	73*	7.8	70*	3.4
3-NT	85	2.4	88	2.1	97	1.3	71*	1.6	75	3.3	74*	2.7
PETN	67*	5.6	59†	1.6	65*	6.0	52†	7.5	60†	5.9	60†	4.0

*Bold, italicized* are IM compounds and degradation products. Green: 75%-125%; light green (\*): 64%-74% or 126%-135%; red (†): <64% or >135%

(DoD QSM Ver5.2 for Solids-lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

### Table 19 (continued). (AVG = average; SD = standard deviation)

	Recoverie	es (%)										
	Polychae	te worm					Ryegrass	;				
	No chromato interferen reductior	No chromatographic interference reduction AVG		column	Florisil co	Florisil column		No chromatographic interference reduction		Alumina column		blumn
	AVG		AVG		AVG		AVG		AVG		AVG	
Compound	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD	( <i>n</i> = 3)	SD
NQ	<i>2,135</i> †	188.2	324	206.0	889	7.4	159	4.1	88	6.0	53	29.3
NTO (210nm)	<i>1,613</i> †	202.6	1,955	372.4	3,060	<i>99.2</i>	1,612	92.2	<b>984</b>	258.5	1,084	1,094.1
NTO (315nm)	<i>20</i> †	1.9	109	11.6	174	11.2	8	5.7	86	27.9	363	493.2

	Recoveries (%) Polyopage											
	Polychae	te worm					Ryegrass	;				
	No chromato interferer reductior	ographic nce 1	Alumina	column	Florisil co	blumn	No chromato interferen reductior	ographic nce 1	Alumina	column	Florisil co	blumn
Compound	AVG ( <i>n</i> = 3)	SD	AVG $(n=3)$	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG $(n=3)$	SD	AVG $(n=3)$	SD
o-NBA (surrogate)	24+	19.4	78	6.9	90	5.9	29+	8.4	137+	43.2	113	96.0
HMX	68*	6.7	71*	2.0	71*	2.2	75	6.4	85	5.0	68*	23.1
RDX	77	0.9	77	2.6	76	1.4	97	3.3	88	6.8	89	18.8
PA (210nm)	75*	5.0	83	8.5	106	5.1	<i>141</i> †	27.4	57†	7.0	89	52.8
PA (315nm)	67*	0.4	65*	3.7	64*	4.8	<i>78</i>	4.4	76	14.2	65*	21.5
4-NP	75*	0.6	75	1.3	78	1.2	<u>83</u>	4.5	81	3.5	78	12.3
1,3,5-TNB	47†	0.8	47†	0.4	46†	5.8	75	3.6	86	6.6	75*	20.2
2,4-DNP	65*	2.7	<b>59</b> †	13.0	<b>69</b> *	5.1	<i>37</i> †	9.6	109	19.3	<i>92</i>	11.9
1,2-DNB (surrogate)	70*	8.6	74*	3.4	72*	5.3	69*	1.1	74*	5.1	70*	9.2
1,3-DNB	70*	2.4	75	1.5	73*	2.2	78	2.0	78	2.8	72*	12.7
2-NP	75*	3.4	66*	4.5	67*	1.8	<i>73</i> *	1.8	<b>63</b> †	3.5	66*	11.4
NB	60†	1.5	59†	1.2	62†	4.1	74*	3.7	66*	4.3	66*	16.8
3,5-DNA (210nm)	78	1.2	82	1.5	85	7.1	72*	1.8	70*	2.8	67*	15.5
3,5-DNA (254nm)	81	3.0	81	4.6	86	5.9	72*	0.4	74*	5.1	71*	16.2
DNAN	<i>79</i>	3.2	<i>73</i> *	1.6	77	5.0	83	1.1	82	8.1	<b>69</b> *	9.8
Tetryl	0†	0.0	0†	0.0	0†	0.0	20†	2.6	17†	2.2	18†	0.9
NG	68*	4.9	67*	7.4	63†	2.4	104	4.9	104	9.1	86	19.1
2,4,6-TNT	64†	1.7	64*	5.6	60†	5.9	75*	4.6	75*	3.5	70*	13.0
4-Am-2,6-DNT	44†	0.4	44†	5.6	48†	1.2	61†	16.8	45†	4.8	42†	9.4
2-Am-4,6-DNT	65*	1.2	64†	6.6	68*	2.8	72*	7.8	73*	4.2	65*	11.9
2,6-DNT	80	1.5	75*	4.1	79	3.7	75*	3.7	72*	3.1	67*	20.2
2,4-DNT	70*	1.2	73*	3.9	75	2.8	76	4.0	75*	3.4	73*	14.6
2-NT	90	2.2	77	6.1	93	5.0	77	2.5	84	6.7	71*	13.4

		Recoverie	es (%)											
		Polychae	te worm					Ryegrass						
	No chromatographic interference reduction			column	Florisil column No chromatographic interference reduction				Alumina	column	Florisil column			
Compound		AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	
4-	NT	86	4.8	69*	6.1	75*	3.8	73*	6.7	75	4.2	62†	12.1	
3-	NT	81	4.4	79	3.5	82	1.9	68*	2.9	73*	1.4	64*	9.1	
PE	TN	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$			2.5	82	7.4	74*	7.2	72*	10.7			

**Bold, italicized** are IM compounds and degradation products.

Green: 75%-125%; light green (\*) : 64%-74% or 126%-135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids–lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

Each chromatographic treatment option was effective for fathead minnow tissue (Table 20). A reduction of NQ recovery was observed for neutral alumina-florisil combinations (to 77%–81%), compared to a positive NQ interference introduced by basic alumina (150% recovery). With the exception of a slightly higher than acceptable recovery for NTO using the silica gel option (141%), NTO detected at 315 nm (detection wavelength selection discussed below) was within DoD QSM Ver5.2 accepted limits (64%–135%) for each treatment option.

For earthworm tissue (Table 21), again several chromatographic treatment options were successful at reducing or eliminating matrix-related interference. For the three neutral alumina-florisil combinations, all compounds were recovered within current DoD QSM Ver5.2 accepted limits. The silica gel option resulted in acceptable recoveries for all compounds as well, with the exception of NTO, which was slightly high at 138%. Treatment with basic alumina resulted in three out-of-spec recoveries, including *o*-NBA surrogate (146%), Tetryl (58%), and NG (169%).

For polychaete worm tissue (Table 22), analytical interferences were previously observed for NQ, NTO, and o-NBA surrogate. All five chromatographic interference reduction options resulted in o-NBA recoveries of approximately 100%. NTO interference was also greatly reduced. NTO recoveries were only slightly above the accepted limit (135%) for mixed neutral alumina-florisil (136%) and silica gel (140%) and within limits for two stacked alumina-florisil options (132%–134%). Unlike for other tissue types, polychaete worm tissue also suffered from analytical interference of NQ. All five chromatographic treatments were able to reduce this interference as well, with the silica gel option yielding the best NQ recovery result (92%), followed by basic alumina (80%) and mixed neutral alumina-florisil (72%). Neither stacked neutral aluminaflorisil option completely removed the NQ interference. However, because each chromatographic interference removal procedure tested demonstrated specific successes and failures, all remain options for specific applications if needed.

For ryegrass plant tissue (Table 23), analytical interferences were observed for NQ, NTO, *o*-NBA surrogate, and 2,4-DNP in the original untreated extracts. With each of the five chromatographic options, the NQ and the 2,4-DNP interferences were completely removed. In addition, all but the mixed neutral alumina-florisil were able to remove the *o*-NBA interference to yield recoveries within or just above the accepted DoD QSM Ver5.2 limit. Meanwhile, NTO interference was reduced by all but the basic alumina option. On the basis of the results from all four tissue types using each of the five chromatographic options, the silica gel option was chosen as the preferred option. This choice was made because of (1) the ability to remove the majority of interferences from the majority of tissue matrices, (2) the generally higher recoveries, and (3) the simplicity as compared to a combination of neutral alumina-florisil, which typically yielded similar recovery results.

### Table 20. Recoveries (%) from postextraction spiked fathead minnow tissue MeOH extract, following five different chromatographic treatment options for matrix-related analytical interference reduction. (AVG = average; SD = standard deviation)

	Fathead	athead minnow—recoveries (%)										
	Neutral flo	alumina   risil	Flo neutral	risil   alumina	Mixed ne alumina-	eutral -florisil	Basic alu	ımina	Silica gel			
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 2)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD		
NQ	81	6.7	81	0.7	77	9.8	<i>150</i> †	<b>8.</b> 7	<i>92</i>	4.7		
NTO (210nm)	<i>2,280</i> †	747.3	<i>1,766</i> †	41.0	<i>1,797</i> †	157.0	<i>3,924</i> †	189.3	<i>2,054</i> †	179.0		
NTO (315nm)	130*	1.7	116	2.0	119	9.4	127*	10.6	<i>141</i> †	5.3		
o-NBA (surrogate)	100	3.6	93	0.1	96	5.3	105	2.8	95	5.4		
HMX	104	9.9	94	1.5	94	6.3	100	4.1	98	3.3		
RDX	99	3.5	97	1.5	94	9.7	106	1.1	99	2.9		
PA (210nm)	119	29.2	115	31.0	127*	60.9	101	3.7	124	14.1		
PA (315nm)	96	3.3	97	<b>0.</b> 7	<u>99</u>	2.5	121	7.2	<u>99</u>	3.6		
4-NP	108	2.2	107	0.5	110	1.7	106	4.3	112	2.0		
1,3,5-TNB	102	4.5	97	4.6	97	3.4	97	4.5	106	4.1		
2,4-DNP	106	<b>9.</b> 7	<i>91</i>	2.7	<u>98</u>	7.6	96	1.5	94	4.2		
1,2-DNB (surrogate)	101	3.3	97	3.7	97	6.8	102	4.6	100	4.6		
1,3-DNB	100	6.5	96	0.8	101	4.9	99	3.2	98	2.6		
2-NP	106	25.8	90	0.9	104	18.5	97	2.5	<i>92</i>	4.3		
NB	95	4.8	96	5.1	93	11.3	97	4.1	96	1.0		
3,5-DNA (210nm)	88	7.1	91	8.3	89	11.5	111	3.8	91	3.2		
3,5-DNA (254nm)	94	0.9	94	1.0	96	2.1	106	8.1	97	1.7		
DNAN	104	3.5	100	10.8	98	10.8	101	8.1	104	3.7		
Tetryl	92	3.8	102	2.4	102	7.8	89	2.7	96	3.6		
NG	91	5.5	100	3.9	104	9.0	111	5.7	88	11.6		
2,4,6-TNT	98	8.6	100	7.1	102	4.8	99	7.0	95	6.5		
4-Am-2,6-DNT	94	7.1	93	0.5	97	3.5	92	3.9	88	3.8		

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	Fathead	head minnow-recoveries (%)										
	Neutral floi	alumina   risil	Flo neutral	risil   alumina	Mixed ne alumina-	eutral -florisil	Basic alu	mina	Silica gel			
Compound	AVG ( <i>n</i> = 3)	AVG ( <i>n</i> = 3) SD (		SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD		
2-Am-4,6-DNT	102	10.5	97	3.9	104	6.4	97	5.6	99	7.4		
2,6-DNT	106	12.6	120	2.2	98	8.5	104	5.8	96	6.2		
2,4-DNT	97	1.7	98	2.2	99	5.9	97	7.1	101	3.8		
2-NT	94	0.8	95	0.6	100	1.3	99	5.8	99	2.6		
4-NT	101	14.0	94	0.3	100	5.0	96	7.3	97	4.6		
3-NT	96	3.7	98	6.1	101	2.6	98	8.9	97	5.8		
PETN	101	96         3.7           101         7.7		0.4	98	6.0	96	6.9	94	8.2		

*Bold, italicized* are IM compounds and degradation products. Green: 75%–125%; light green (\*): 64%–74% or 126%–135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

	Earthwor	arthworm—recoveries (%)								
	Neutral flo	alumina   risil	Flo neutral	risil   alumina	Mixed ne alumina-	eutral -florisil	Basic alu	imina	Silica gel	
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NQ	95	1.2	67*	1.2	84	<b>9.</b> 7	<u>98</u>	1.0	116	4.9
NTO (210nm)	<i>1,814</i> †	19.8	<i>2,081</i> †	180.4	<i>1,971</i> †	69.4	<i>746</i> †	255.7	<i>2,244</i> †	54.6
NTO (315nm)	121	5.7	115	10.5	117	2.6	82	<b>8.</b> 7	<i>138</i> †	6.2
o-NBA (surrogate)	86	15.9	89	12.8	76	7.3	146†	17.0	78	5.5
HMX	89	9.6	91	11.8	86	7.8	91	2.8	81	7.1
RDX	76	33.7	96	3.2	96	3.0	107	7.4	99	0.1
PA (210nm)	<i>170</i> †	103.3	<i>175</i> †	82.1	<i>149</i> †	70.8	86	10.4	112	63.4
PA (315nm)	<i>93</i>	6.9	84	16.9	80	9.7	87	23.2	67*	14.8
4-NP	113	27.3	<i>92</i>	9.2	90	0.7	101	0.7	89	10.4
1,3,5-TNB	96	10.9	92	3.7	92	4.4	98	0.4	89	8.5
2,4-DNP	<i>95</i>	1.5	96	2.4	96	1.5	104	12.3	<i>95</i>	4.1
1,2-DNB (surrogate)	97	2.8	100	0.7	98	1.0	104	9.6	101	2.8
1,3-DNB	95	1.5	98	1.2	96	2.3	100	0.4	96	2.0
2-NP	101	14.4	91	3.1	<i>91</i>	1.5	85	3.7	88	3.6
NB	99	2.0	98	6.4	98	3.0	131*	15.3	98	2.7
3,5-DNA (210nm)	97	14.3	94	4.4	88	3.0	100	0.6	90	4.4
3,5-DNA (254nm)	94	2.6	96	1.6	94	2.2	112	5.5	94	2.6
DNAN	104	14.2	102	6.2	95	3.2	113	8.2	<u>98</u>	6.4
Tetryl	89	13.0	103	11.4	106	4.2	58†	2.7	104	6.1
NG	70*	32.3	111	16.7	107	20.8	169†	11.6	116	19.2
2,4,6-TNT	95	5.5	99	5.1	103	7.0	108	1.4	104	8.3
4-Am-2,6-DNT	104	10.5	98	5.0	97	1.6	104	7.8	95	0.9
2-Am-4,6-DNT	105	7.4	97	1.3	98	3.0	104	9.5	100	2.7

# Table 21. Recoveries (%) from postextraction spiked earthworm tissue MeOH extract, following five different chromatographic treatment options for matrix-related analytical interference reduction. (AVG = average; SD = standard deviation)

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	Earthworm—recoveries (%)												
	Neutral flo	alumina   risil	Florisil   neutral alumina		Mixed ne alumina-	eutral -florisil	Basic alu	imina	Silica gel				
	AVG ( <i>n</i> = 3) SD		AVG ( <i>n</i> = 3)	SD	AVG (n=3) SD		AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD			
2,6-DNT	96	8.9	90	2.5	93	6.7	102	2.4	85	1.0			
2,4-DNT	99	4.8	99	3.2	96	3.6	101	1.1	97	2.9			
2-NT	98	4.1	101	4.1	101	1.1	117	2.1	98	2.8			
4-NT	97	2.9	104	9.7	94	5.9	99	4.6	100	0.8			
3-NT	99	5.1	100	3.7	100	4.0	109	9.0	97	2.1			

6.5

89

95

2.0

95

6.4

89

Compound

PETN

99

1.4

*Bold, italicized* are IM compounds and degradation products. Green: 75%-125%; light green (\*): 64%-74% or 126%-135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids-lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

5.4

	Polychaete worm-recoveries (%)									
	Neutral	alumina 	Flo	risil I	Mixed ne	eutral	Basic alu	ımina	Silica gel	
	flo	risil	neutral	alumina	alumina	-florisil				
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NQ	<i>140</i> †	100.4	<i>183</i> †	123.6	72*	5.4	80	5.3	92	6.3
NTO (210nm)	<b>9</b> 77†	218.1	<b>848</b> †	49.0	<b>836</b> †	25.5	<i>1,860</i> †	642.5	<b>939</b> †	61.5
NTO (315nm)	134*	2.1	132*	7.6	<i>136</i> †	5.1	<i>148</i> †	5.2	<i>140</i> †	7.3
o-NBA (surrogate)	102	2.0	100	0.7	102	3.6	105	6.2	98	6.8
HMX	98	3.2	95	2.5	97	3.5	92	1.7	91	8.2
RDX	99	3.7	101	3.1	100	3.5	95	1.8	95	6.7
PA (210nm)	<u>98</u>	5.2	102	11.9	122	7.1	<i>93</i>	1.7	101	8.9
PA (315nm)	<u>98</u>	1.3	<u>99</u>	0.7	<u>99</u>	2.4	95	6.6	94	7.3
4-NP	<u>99</u>	1.7	<u>99</u>	0.4	100	3.5	96	1.6	95	6.3
1,3,5-TNB	97	2.8	96	1.5	107	16.0	93	2.0	94	8.3
2,4-DNP	<u>98</u>	3.9	97	2.3	90	9.5	94	1.9	<u>99</u>	8.0
1,2-DNB (surrogate)	104	3.9	102	1.5	92	9.4	97	2.3	101	7.9
1,3-DNB	99	1.1	99	2.2	100	3.4	97	5.1	96	8.0
2-NP	<i>91</i>	4.3	90	4.7	<u>89</u>	4.1	103	7.6	89	5.2
NB	98	2.6	99	0.4	98	5.1	91	2.5	97	9.9
3,5-DNA (210nm)	96	7.9	100	5.5	99	8.7	108	0.1	90	9.1
3,5-DNA (254nm)	103	3.9	102	2.6	104	7.1	105	11.1	93	8.6
DNAN	<u>98</u>	9.3	100	6.5	<i>92</i>	1.4	<i>91</i>	4.8	94	8.6
Tetryl	101	2.6	97	0.7	97	3.6	89	0.4	95	8.4
NG	106	5.6	95	3.7	98	16.7	103	2.0	98	13.6
2,4,6-TNT	101	1.5	98	1.0	103	7.1	95	1.3	95	6.7
4-Am-2,6-DNT	99	2.7	98	5.5	101	9.5	91	1.6	93	9.2
2-Am-4,6-DNT	97	3.3	95	7.4	96	0.8	95	2.1	91	8.0

## Table 22. Recoveries (%) from postextraction spiked polychaete worm tissue MeOH extract, following five different chromatographic treatment options for matrix-related analytical interference reduction. (AVG = average; SD = standard deviation)

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	Polychae	Polychaete worm-recoveries (%)											
	Neutral flo	alumina   risil	Florisil   neutral alumina		Mixed neutral alumina-florisil		Basic alu	imina	Silica gel				
Compound	AVG ( <i>n</i> = 3)	AVG ( <i>n</i> = 3) SD		SD	AVG (n=3) SD		AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD			
2,6-DNT	97	2.4	100	3.9	96	4.3	105	6.0	103	15.2			
2,4-DNT	99	3.3	94	3.4	99	5.5	94	2.9	96	7.3			
2-NT	100	3.0	99	4.1	102	8.8	103	3.3	92	6.8			
4-NT	99	1.4	96	9.5	99	1.1	94	0.9	93	6.8			
3-NT	106	4.7	104	4.2	106	5.3	101	1.8	96	8.6			
PETN	100	0.5	95	4.8	99	2.9	95	7.8	90	8.5			

*Bold, italicized* are IM compounds and degradation products. Green: 75%-125%; light green (\*): 64%-74% or 126%-135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids-lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

	Ryegrass	Ryegrass—recoveries (%)										
	Neutral flo	Neutral alumina florisil AVG A		risil   alumina	Mixed ne alumina-	eutral -florisil	Basic alu	ımina	Silica gel			
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD		
NQ	97	4.7	82	7.6	88	0.7	100	3.5	90	7.0		
NTO (210nm)	<i>573</i> †	147.7	<i>576</i> †	63.6	<i>500</i> †	45.4	<i>1,081</i> †	1,272.6	<b>595</b> †	<b>29.</b> 7		
NTO (315nm)	114	3.2	112	3.0	113	3.9	<i>174</i> †	146.1	130*	4.9		
o-NBA (surrogate)	138†	29.6	127*	15.0	171†	6.3	107	8.4	146†	8.0		
HMX	98	2.7	93	5.5	101	4.0	99	5.0	96	9.9		
RDX	100	5.2	100	2.7	101	3.1	101	2.9	99	2.8		
PA (210nm)	96	12.1	84	3.1	101	2.8	90	15.6	96	5.4		
PA (315nm)	100	3.5	101	1.8	101	2.5	87	12.5	<i>93</i>	3.4		
4-NP	101	1.0	100	3.1	<u>98</u>	1.8	<u>98</u>	4.2	96	3.9		
1,3,5-TNB	99	3.6	100	6.1	98	2.7	96	3.4	96	5.8		
2,4-DNP	<u>98</u>	2.4	106	10.8	104	4.3	95	6.1	96	4.8		
1,2-DNB (surrogate)	108	1.3	112	5.8	106	2.4	104	1.7	103	9.3		
1,3-DNB	100	2.4	101	1.4	99	2.1	99	2.1	97	4.2		
2-NP	<i>92</i>	4.2	84	3.6	<b>95</b>	4.7	84	15.1	88	4.1		
NB	103	1.4	103	4.0	100	0.5	97	3.9	103	2.6		
3,5-DNA (210nm)	101	2.9	113	6.2	96	1.2	98	20.6	97	4.0		
3,5-DNA (254nm)	101	1.4	103	3.3	97	4.1	99	25.7	94	1.6		
DNAN	102	4.7	115	7.3	96	1.9	92	5.0	100	3.7		
Tetryl	100	3.6	97	3.6	94	2.8	95	8.4	96	7.5		
NG	109	8.3	100	2.1	94	2.4	109	7.3	86	13.3		
2,4,6-TNT	110	8.2	105	8.5	98	4.1	100	2.8	97	3.3		
4-Am-2,6-DNT	98	1.7	99	3.7	98	2.6	96	5.4	104	9.9		
2-Am-4,6-DNT	98	0.4	100	6.0	97	2.6	99	2.0	95	6.1		

# Table 23. Recoveries (%) from postextraction spiked ryegrass tissue MeOH extract, following five different chromatographic treatment options for matrix-related analytical interference reduction. (AVG = average; SD = standard deviation)

	Ryegrass	-recoverie	es (%)							
	Neutral flo	alumina   risil	Flo neutral	risil   alumina	Mixed ne alumina-	eutral -florisil	Basic alu	imina	Silica gel	
	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
2,6-DNT	98	3.1	97	8.2	96	2.0	93	9.8	108	8.5
2,4-DNT	98	2.0	101	1.7	101	5.5	99	2.3	95	3.8
2-NT	99	3.0	101	5.7	102	3.3	93	23.3	97	2.6
4-NT	101	0.3	99	1.1	99	4.2	97	7.3	96	2.7
3-NT	107	6.1	98	7.1	99	9.8	92	16.4	95	7.0

101

107

5.4

99

1.1

7.3

97

2.3

Compound

PETN

100

*Bold, italicized* are IM compounds and degradation products. Green: 75%-125%; light green (\*): 64%-74% or 126%-135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids-lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

3.8

During method development for tissue matrix—related interference reduction, comparative analysis of NTO and of PA using detection wavelengths of 210 nm and 315 nm showed that much of the analytical interferences observed at 210 nm were absent at 315 nm; therefore, use of 315 nm was preferred for determinations of NTO and PA in tissue extracts. A detection wavelength of 315 nm may also be preferred for NTO and PA in water and soil matrices as a measure to combat the contribution of interferent compounds in munition compound determinations. In addition, lower MDLs were obtained for both NTO and PA in water and soil by detecting at 315 nm rather than 210 nm (Table 6, section 3.1).

Finally, pre-extraction spiked tissue sample MeOH extracts were treated with the silica gel interference reduction option. Overall, the results (Table 24) showed that using a single 18 h MeOH extraction in combination with silica gel treatment proved sufficient to yield recoveries for most compounds within current DoD QSM limits. The only compound recovered poorly from all four tissue matrices was Tetryl, with the highest recovery obtained for plant tissue, which again may point to the matrixdependent stability of this compound. The silica gel treatment removed or reduced the majority of interferences previously observed for each tissue type. Nontarget chromatographic peaks remained in the chromatographic region surrounding NTO, o-NBA surrogate, and 1,3,5-TNB (for earthworm). Of these remaining observed interferences, only the 1,3,5-TNB determination was affected. For polychaete worm, the 1,3,5-TNB and 4-Am-2,6-DNT recoveries were somewhat low but had been shown in previous experiments to be recovered within QSM Ver5.2 accepted limits. Therefore, 18 h MeOH extraction in combination with silica gel chromatographic treatment for interference reduction was selected for future work.

		Recoverie	s (%)	
Compound	Fathead minnow	Polychaete worm	Earthworm	Ryegrass
NQ	113	<i>73</i> *	86	68*
NTO (210 nm)	<i>8,477</i> †	<i>5,318</i> †	<i>4,503</i> †	<i>149</i> †
NTO (315 nm)	87	113	98	133*
o-NBA (surrogate)	101	82	111	67*
HMX	81	66*	79	99
RDX	86	66*	93	85
PA (210 nm)	79	<i>73</i> *	<i>37</i> †	79
PA (315 nm)	66*	66*	<i>68</i> *	71*
4-NP	83	71*	<i>72</i> *	84
1,3,5-TNB	84	52†	3,670†	83
2,4-DNP	110	<i>72</i> *	81	93
1,2-DNB (surrogate)	93	72*	82	73*
1,3-DNB	76	65v	65*	86
2-NP	105	81	74*	94
NB	84	94	83	90
3,5-DNA (210 nm)	183†	139†	73*	90
3,5-DNA (254 nm)	69*	75*	80	73*
DNAN	123	<i>72</i> *	86	76
Tetryl	16†	6†	4†	62†
NG	116	132*	87	125
2,4,6-TNT	119	100	80	80
4-Am-2,6-DNT	78	57†	74*	88
2-Am-4,6-DNT	102	67*	88	85
2,6-DNT	110	71*	79	85
2,4-DNT	108	67*	69*	88
2-NT	129*	101	68*	89
4-NT	95	73*	69*	80
3-NT	64*	64*	68*	86
PETN	109	74*	67*	91

Table 24. Recoveries (%) from pre-extraction spiked tissue MeOH extracts, following chromatographic treatment with silica gel for matrix-related analytical interference reduction. (AVG = average; SD = standard deviation)

Bold, italicized are IM compounds and degradation products.

Green: 75%-125%; light green (\*): 64%-74% or 126%-135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

### 3.9 Stability study

Extraction and analysis hold-time studies were carried out in reagent water (direct injection and SPE) and three soils (Riley, Yuma, and ASTM fat clay, described in Table 14) spiked with the 17 legacy and 7 IM compounds of interest and two surrogates. Currently accepted pre-extraction hold times are 7 days for waters and 14 days for solid matrices, and postextraction hold time for analysis is 40 days for samples analyzed by USEPA Method 8330B, which references USEPA SW-846 section 4.1 guidelines for semi-volatile organics (USEPA 2014). As these hold times were adopted for methods including only legacy munition compounds, experiments were carried out to test whether the limits would also be appropriate for the seven compounds of interest and *o*-NBA surrogate added under SERDP-ER2722. Soils with varying physicochemical properties were tested in order to observe any potential matrix-related variability.

Results for the pre-extraction hold-time experiments for water are shown in Table 25. For direct-injection water samples, the majority of compounds were recovered within current DoD QSM limits on Day 0 and Day 7, so most compounds were stable under all three storage conditions for at least seven days. However, NTO recovery was <57% on Day 7 for samples stored at room temperature in clear containers. For unknown reasons, NTO determinations were generally very high in this study, possibly suffering from analytical interference not observed elsewhere in the current project. Still, the recoveries for NTO stored at room temperature in clear containers were extremely low compared to those for samples stored under the other two conditions. Therefore, NTO determinations in samples stored in light for extended periods might not be reliable.

In general, SPE option B proved more effective at NQ recovery than option A, and option B was selected during SPE method finalization. Apart from possible NTO instability, only NQ showed possible instability at room temperature. NQ recoveries were lower for room temperature than for cold samples and were lower in samples stored in clear containers than those stored in amber containers. Thus, NQ determinations in samples stored at room temperature (especially in light) for extended periods might not be reliable.

Results for the postextraction hold-time experiments for water are shown in Table 26. Again, for unknown reasons, NTO determinations were generally very high in this study, possibly due to analytical interference. Overall, though, NTO recoveries decreased from Day o to Day 40, and were lower in room temperature samples, especially those stored in clear containers. However, all of the NTO recoveries were nominally within current DoD QSM limits. Therefore, determination of NTO might be affected by possible instability issues.

On the other hand, NQ determinations in SPE samples differed significantly depending on the SPE cartridge option used to extract and preconcentrate the water sample. For SPE option A, recoveries for NQ were much higher (about 100%) in Day 40 samples, whereas the average recovery on Day 0 was very low (19%). There could be a positive analytical interferent, such as a degradation product, accumulating over the course of the 40-day storage period and enhancing the NQ recovery determination. Alternately, for SPE option B, recoveries for NQ were fairly stable over the 40-day period.

In addition, instability of the three NTs was observed in direct-injection samples and SPE samples, with recoveries for each of them generally decreasing with increased temperature and light exposure. Recoveries of these compounds were within current DoD QSM limits for Day 40 samples stored in cold temperatures, indicating that the instability could be reduced by avoiding elevated temperatures and keeping with current recommendations to store at  $\leq 4^{\circ}$ C.

Results for the pre-extraction hold-time experiments for soil are shown in Table 27. For unknown reasons, in the Day O soil samples, a few compounds were recovered at lower than usual levels, especially for the Riley field soil. In addition, NTO, HMX, and RDX recoveries were recovered at levels below current limits (<64%) at Day 14. This lower recovery was observed for two of the soils for HMX and RDX, and for all three soils for NTO, indicating that pre-extraction holding-time limits might need shortening for soils where these compounds are known or suspected to be present. Overall, though, the majority of compounds were recovered within current DoD QSM limits at both Day 0 and Day 14, including all other IM compounds tested.

Results for the postextraction hold-time experiments for soil are shown in Table 28. Again, for unknown reasons, NTO determinations were

generally very high in this study, possibly because of analytical interference.

	Recover	ries (%)																						
							Waters	Day 7 extr	action and	d analysis			•						-					
	Waters	Day 0 extr	action and	d analysis			Cold/Da	ark					Room te	emperatur	e / Dark				Room te	emperatur	e / Light			
	SPE A (e	evap)	SPE B (e	evap)	Direct i	njection	SPE A (r	no evap)	SPE B (r	no evap)	Direct in	ijection	SPE A (r	no evap)	SPE B (I	no evap)	Direct in	njection	SPE A (n	io evap)	SPE B (r	no evap)	Direct in	jection
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NQ	<i>19</i> †	1.4	<i>42</i> †	1.9	74	13.6	<b>14</b> †	2.9	<b>34</b> †	4.3	75	20.8	<i>15</i> †	2.1	<i>26</i> †	1.1	88	9.8	<i>15</i> †	0.7	<i>21</i> †	6.4	80	9.9
NTO (210nm)	<i>261</i> †	2.5	<b>286</b> †	14.1	<i>138</i> †	2.8	<i>305</i> †	37.8	<i>311</i> †	4.2	79	19.2	<i>293</i> †	17.7	<i>303</i> †	6.3	80	3.8	<i>300</i> †	11.7	<i>229</i> †	62.9	<i>29</i> †	22.9
NTO (315nm)	<i>270</i> †	2.4	<b>292</b> †	15.2	99	14.2	<i>306</i> †	40.9	<i>307</i> †	2.4	78	25.0	<i>291</i> †	16.4	<b>299</b> †	2.3	<i>68</i> *	17.1	<i>300</i> †	<i>9.9</i>	<b>227</b> †	66.6	<i>53</i> †	8.9
o-NBA (surrogate)	107	0.9	112	7.0	106	10.6	106	16.1	104	4.7	100	23.0	99	5.3	99	2.1	101	4.2	103	3.4	66*	30.3	107	1.8
HMX	91	0.8	98	5.1	88	9.6	104	10.2	108	2.3	95	0.6	100	3.9	104	0.7	100	4.7	104	3.2	103	10.5	99	1.5
RDX	77	4.3	88	7.8	<i>68</i> *	10.3	93	20.9	102	2.4	91	3.1	78	19.0	101	3.3	90	0.3	96	0.7	95	14.4	116	10.0
PA (210nm)	81	1.5	93	5.4	84	5.8	101	21.7	111	0.3	97	5.6	88	21.6	107	5.4	116	6.5	104	3.5	101	17.0	<i>125</i> *	15.8
PA (315nm)	93	1.1	100	5.4	83	4.9	107	6.9	109	0.9	97	0.3	101	6.0	107	2.1	103	7.6	108	1.8	106	9.1	96	0.5
4-NP	90	0.8	97	5.0	92	6.3	110	7.0	114	1.0	109	1.0	105	6.4	111	0.3	105	5.2	110	4.4	111	10.4	112	1.6
1,3,5-TNB	89	1.3	95	4.9	87	7.0	104	6.5	105	1.2	102	1.2	99	6.1	102	0.8	104	1.0	105	2.8	102	9.9	107	7.3
2,4-DNP	85	0.6	90	4.3	96	2.5	106	18.7	114	0.8	119	7.9	95	19.5	112	0.7	<i>136</i> †	4.7	110	5.3	112	12.4	<i>125</i> *	5.7
1,2-DNB (surrogate)	87	0.8	79	7.0	97	5.3	114	8.9	115	3.6	116	10.3	109	5.4	113	1.6	115	3.7	115	2.5	113	11.4	116	7.0
1,3-DNB	84	0.8	77	6.7	87	6.2	104	5.6	106	1.0	104	3.1	100	5.5	104	1.6	104	1.7	105	2.1	98	16.8	106	1.0
2-NP	<i>5</i> †	3.0	<i>0</i> †	0.0	87	0.8	88	5.9	93	1.9	87	5.6	82	5.5	89	0.6	92	5.7	81	13.2	87	<i>12.8</i>	98	3.0
NB	90	2.3	97	7.9	79	9.7	104	7.1	106	0.6	106	5.1	98	6.1	104	0.2	108	6.0	104	1.3	102	10.9	107	4.8
3,5-DNA (210nm)	1†	1.1	0†	0.0	88	9.4	93	8.0	101	1.4	107	12.2	81	0.1	98	1.9	101	6.7	91	3.5	93	16.2	109	2.0
3,5-DNA (254nm)	1†	0.8	0†	0.0	87	10.8	93	7.5	102	2.2	98	3.6	86	5.3	99	1.6	96	2.0	93	4.4	93	16.3	97	3.5
DNAN	88	0.5	92	6.5	<i>72</i> *	5.4	98	5.2	98	1.4	109	0.1	92	6.3	96	1.7	100	2.5	97	3.1	89	22.0	113	14.8
Tetryl	89	1.8	95	4.6	86	16.1	100	5.7	103	1.0	96	8.0	95	5.5	100	0.7	86	6.7	101	0.3	97	9.3	75	16.2
NG	86	0.6	85	5.9	39†	10.3	97	5.7	97	1.0	93	10.5	91	4.8	95	3.2	92	4.4	99	3.3	98	8.8	89	11.4
2,4,6-TNT	87	0.9	91	5.5	60*	12.2	103	6.2	103	1.8	101	2.2	98	5.8	100	1.7	96	5.3	100	7.6	98	16.5	99	3.9
4-Am-2,6-DNT	90	0.2	97	4.9	85	8.0	101	7.8	106	1.0	104	10.7	97	6.6	102	0.3	110	2.7	102	0.9	101	11.1	102	3.8
2-Am-4,6-DNT	92	2.4	99	4.9	88	6.9	101	6.6	105	1.6	108	15.4	97	6.5	102	1.7	109	2.8	105	3.9	99	15.4	106	8.6
2,6-DNT	74*	2.6	38†	10.0	100	11.8	96	4.8	96	1.8	68*	6.4	92	6.3	95	1.5	71*	1.6	96	2.0	86	20.5	81	7.4
2,4-DNT	84	0.8	70*	7.2	91	13.5	100	6.6	99	1.9	78	7.5	94	6.4	96	3.2	69*	13.4	99	1.2	81	28.0	77	4.3
2-NT	1†	0.5	1†	0.0	65*	19.5	86	8.5	96	2.6	81	17.8	78	4.5	93	1.1	84	6.7	84	5.9	84	21.1	94	13.7
4-NT	4†	1.8	1†	0.3	88	9.9	92	6.4	100	4.8	99	5.5	88	6.7	95	3.3	94	11.7	92	4.5	82	26.8	93	18.3
3-NT	2†	1.3	1†	0.1	75*	11.7	91	7.6	103	2.3	92	10.2	86	5.8	99	3.2	96	1.7	91	4.4	86	27.0	98	3.8
PETN	92	1.4	88	14.7	73*	12.8	103	7.4	107	0.8	108	24.7	98	7.9	102	5.7	101	8.3	104	2.0	103	12.9	107	3.0

Table 25. Extraction hold-time stability for laboratory-spiked reagent water samples over seven days. (AVG = average; SD = standard deviation)

Bold, italicized are IM compounds and degradation products. Green: 75%–125%; light green (\*): 64%–74% or 126%–135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

	Recov	eries (%)																										
	Watar			n and a	nahraia		Water	s Day 0 e	extractio	n, Day 4	10 analy	sis																
	waters	s Day U (	extractio	n anu a	naiysis		Cold/	Dark									Room	tempera	ature / D	ark			Room	tempera	iture / L	ight		
	SPE A	(evap)	SPE B	(evap)	Direct injecti	on	SPE A	(evap)	SPE A (no ev	ap)	SPE B	(evap)	SPE B (no eva	ap)	Direct injectio	on	SPE A	(evap)	SPE B	(evap)	Direct injectio	on	SPE A	(evap)	SPE B	(evap)	Direct injecti	on
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NQ	<i>19</i> †	1.4	<i>42</i> †	1.9	<b>74</b> *	13.6	102	16.8	<b>17</b> †	2.9	<i>37</i> †	5.0	<i>38</i> †	1.7	89	20.0	100	14.3	<b>42</b> †	4.1	83	12.9	98	21.0	<i>39</i> †	4.5	77	3.9
NTO (210nm)	<i>261</i> †	2.5	<i>286</i> †	14.1	<i>138</i> †	2.8	<i>126</i> *	2.0	123	1.2	<i>134</i> *	5.1	<i>134</i> *	9.0	<i>136</i> †	3.3	<i>128</i> *	2.3	<i>142</i> †	0.0	<i>136</i> †	3.3	<i>127</i> *	2.9	<i>139</i> †	3.2	120	2.7
NTO (315nm)	270†	2.4	<i>292</i> †	15.2	99	14.2	<i>128</i> *	2.6	123	0.8	<i>133</i> *	4.9	<i>131</i> *	9.0	<i>130</i> *	4.5	<i>130</i> *	3.0	<i>142</i> †	0.0	<i>131</i> *	3.1	<i>128</i> *	3.2	<b>140</b> †	3.0	<i>126</i> *	1.2
o-NBA (surrogate)	107	0.9	112	7.0	106	10.6	109	1.1	111	1.9	112	1.5	118	4.9	117	4.6	106	0.5	109	5.1	117	5.9	105	3.3	108	4.3	118	1.4
HMX	91	0.8	98	5.1	88	9.6	81	30.6	108	0.8	117	4.3	116	7.8	96	2.4	113	2.4	125	1.9	99	2.4	112	2.7	123	2.6	98	3.1
RDX	77	4.3	88	7.8	<i>68</i> *	10.3	116	2.4	113	1.7	120	4.7	120	8.4	106	3.3	118	2.7	127*	2.4	107	5.4	117	3.3	<i>126</i> *	2.6	105	3.5
PA (210nm)	81	1.5	93	5.4	84	5.8	88	2.1	81	1.6	93	5.0	88	1.7	114	10.6	89	1.1	99	2.0	111	4.0	88	2.2	99	2.1	<i>139</i> †	7.0
PA (315nm)	93	1.1	100	5.4	83	4.9	94	1.6	83	1.5	102	1.6	92	2.6	97	3.1	96	1.4	108	2.2	113	3.7	95	2.2	107	1.8	135*	4.0
4-NP	90	0.8	97	5.0	92	6.3	106	1.6	100	1.1	111	4.5	108	6.9	107	3.2	108	2.3	119	2.2	109	1.2	107	2.7	117	2.3	109	1.5
1,3,5-TNB	89	1.3	95	4.9	87	7.0	109	2.2	109	1.4	111	5.8	115	7.2	111	3.7	110	1.6	116	0.9	115	0.3	108	2.0	113	1.9	116	3.1
2,4-DNP	85	0.6	90	4.3	96	2.5	110	1.5	108	1.9	112	5.8	119	5.3	99	8.9	111	2.4	120	4.1	100	10.4	110	2.1	118	4.5	106	12.0
1,2-DNB (surrogate)	87	0.8	79	7.0	97	5.3	108	1.2	114	2.1	97	7.6	120	9.6	110	3.0	106	2.2	99	8.3	102	7.8	105	1.7	98	8.0	107	10.8
1,3-DNB	84	0.8	77	6.7	87	6.2	106	2.1	110	0.4	102	19.7	116	6.3	104	3.6	109	2.2	104	0.2	101	6.8	109	3.0	99	5.8	103	3.5
2-NP	<b>5</b> †	3.0	<i>0</i> †	0.0	87	0.8	<b>7</b> †	3.8	103	0.6	<i>0</i> †	0.0	106	3.1	91	40.7	<i>6</i> †	3.3	<i>0</i> †	0.0	82	52.9	<i>6</i> †	3.5	<i>0</i> †	0.0	92	42.0
NB	90	2.3	97	7.9	79	9.7	116	2.8	105	0.2	116	0.1	112	7.0	111	6.1	116	2.4	121	7.5	127*	0.1	110	1.4	123	0.2	120	3.6
3,5-DNA (210nm)	1†	1.1	0†	0.0	88	9.4	0†	0.0	102	0.8	0†	0.0	104	3.0	89	68.7	0†	0.0	0†	0.0	75	65.1	0†	0.0	0†	0.0	129*	1.6
3,5-DNA (254nm)	1†	0.8	0†	0.0	87	10.8	0†	0.0	117	0.7	0†	0.0	119	4.0	81	66.7	0†	0.0	0†	0.0	111	2.0	0†	0.0	0†	0.0	74*	63.7
DNAN	88	0.5	92	6.5	<i>72</i> *	5.4	104	2	89	0	104	1	94	4	92	6	106	2	110	2	93	4	102	2	109	2	91	2.6
Tetryl	89	1.8	95	4.6	86	16.1	110	2	111	1	113	4	117	7	101	4	114	2	117	2	78	9	111	3	111	3	49†	7.1
NG	86	0.6	85	5.9	39†	10.3	115	1.6	121	4.2	112	6.5	129*	8.4	101	5.7	107	3.6	117	4.5	96	10.3	105	1.3	115	4.8	94	8.2
2,4,6-TNT	87	0.9	91	5.5	60*	12.2	113	2.2	112	3.6	115	4.0	120	7.4	110	4.2	114	2.3	122	2.0	107	4.3	112	2.4	120	2.1	109	2.9
4-Am-2,6-DNT	90	0.2	97	4.9	85	8.0	110	2.1	106	2.4	114	4.9	116	8.5	100	7.2	111	1.9	108	13.1	98	9.8	107	3.2	102	15.6	106	7.1
2-Am-4,6-DNT	92	2.4	99	4.9	88	6.9	112	2.3	106	2.2	116	4.8	117	9.0	107	6.6	112	4.1	114	11.7	102	11.2	110	2.2	117	0.1	116	7.3
2,6-DNT	74*	2.6	38†	10.0	100	11.8	99	1.0	111	3.3	49†	11.1	113	1.2	80	19.9	103	1.5	57*	1.0	68*	15.4	104	6.8	59*	0.4	70*	14.4
2,4-DNT	84	0.8	70*	7.2	91	13.5	94	4.8	104	4.7	80	3.3	106	7.5	102	3.1	99	4.7	73*	40.4	93	12.2	97	1.3	79	28.0	103	7.5
2-NT	1†	0.5	1†	0.0	65*	19.5	0†	0.0	110	2.1	0†	0.0	110	3.8	74*	65.1	0†	0.0	1†	1.5	73*	63.2	0†	0.0	1†	1.7	108	1.8
4-NT	4†	1.8	1†	0.3	88	9.9	4†	2.9	109	2.6	0†	0.0	108	4.1	67*	58.7	2†	0.0	1†	0.3	52†	47.0	2†	0.0	1†	1.0	54†	47.5
3-NT	2†	1.3	1†	0.1	75*	11.7	2†	1.3	108	1.3	0†	0.0	104	6.5	65*	56.4	2†	1.2	0†	0.2	59*	51.7	2†	1.2	0†	0.2	57†	49.4
PETN	92	1.4	88	14.7	73*	12.8	110	2.5	112	3.8	109	15.8	117	8.2	117	14.0	99	5.7	132*	30.2	115	7.8	97	3.6	134*	31.4	120	11.2

Table 26. Analysis hold-time stability for laboratory-spiked reagent water samples over 40 days. (AVG = average; SD = standard deviation)

 Bold, italicized are IM compounds and degradation products.

 Green: 75%-125%; light green (\*): 64%-74% or 126%-135%; red (†): <64% or >135%

 (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

	Recove	eries (%)																						
					•-		Soils Da	ay 14 ext	traction a	nd analy	sis													
	Solis D	ay 0 extra	action an	a analys	IS		Cold/D	ark					Room t	emperat	ure / Dar	k			Room t	emperat	ure / Ligi	ht		
	Riley		Yuma		ASTM F	C	Riley		Yuma		ASTM F	-C	Riley		Yuma		ASTM F	ic .	Riley		Yuma		ASTM F	÷C
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NQ	<i>65</i> *	4.7	<i>65</i> *	4.1	<i>64</i> *	4.1	98	8.3	147†	10.1	102	6.9	90	6.9	116	10.0	95	3.3	98	4.9	119	14.0	103	8.2
NTO (210nm)	<i>260</i> †	7.4	<i>73</i> *	1.5	78	4.9	33†	3.3	35†	1.4	34†	2.7	31†	2.2	35†	2.6	35†	1.5	31†	1.8	38†	3.8	30†	1.3
o-NBA (surrogate)	108	1.0	100	1.7	97	4.1	91	6.2	99	1.2	95	1.8	90	3.2	96	2.0	93	0.8	90	7.9	101	1.0	97	3.4
HMX	41†	0.6	71*	3.2	57*	14.2	43†	1.2	64*	2.3	39†	2.1	38†	4.6	63*	1.7	38†	2.9	47†	1.9	68*	2.3	41†	4.1
RDX	91	1.4	113	3.4	112	2.3	53†	4.1	64*	7.7	44†	1.3	53†	1.0	63*	3.9	42†	3.0	56†	4.9	65*	5.9	44†	7.9
PA (210nm)	56†	0.5	81	3.3	59*	0.4	83	8.0	119	1.3	111	0.6	67*	12.2	125	1.5	109	2.6	69*	12.3	131*	0.6	114	0.6
4-NP	<i>125</i> *	1.9	<i>125</i> *	1.0	<i>127</i> *	1.4	115	0.7	119	0.9	120	0.9	103	5.8	114	0.7	117	0.8	113	3.0	121	0.4	122	2.1
1,3,5-TNB	102	2.7	115	0.7	112	3.2	97	0.6	109	0.7	103	0.3	87	3.9	107	1.4	98	1.8	89	4.5	114	0.6	103	1.1
2,4-DNP	92	4.0	102	5.3	103	1.0	90	3.7	103	1.5	101	0.5	75	3.7	100	3.0	96	1.6	83	1.3	105	1.4	102	2.4
1,2-DNB (surrogate)	125*	4.3	124	2.0	124	5.1	121	2.7	122	3.8	119	0.7	110	5.8	115	1.3	118	2.8	117	4.0	122	2.8	122	2.0
1,3-DNB	117	1.7	118	0.6	120	1.2	108	0.3	111	1.0	112	0.6	95	4.4	106	1.0	110	1.3	104	2.4	112	0.3	113	2.0
2-NP	98	3.9	80	0.3	81	1.3	90	0.4	82	2.2	80	0.4	81	3.3	78	1.4	76	0.7	89	1.5	85	1.9	81	1.4
NB	101	1.2	113	2.3	115	0.7	87	1.4	110	0.2	110	0.6	64*	4.7	105	0.9	102	2.2	67*	1.9	112	1.2	107	1.6
3,5-DNA (210nm)	114	1.7	91	0.6	103	9.8	99	3.1	96	4.8	98	4.4	92	5.1	89	1.7	95	2.9	103	8.0	96	4.9	101	2.3
DNAN	103	2.8	104	1.0	109	3.3	102	2.6	112	1.4	107	4.1	89	5.7	107	1.5	102	2.5	98	0.2	113	3.0	105	2.6
Tetryl	114	3.9	111	0.6	112	0.7	100	1.3	104	4.4	104	3.5	68*	7.6	66*	3.2	75	2.7	63*	4.7	70*	2.1	78	2.0
NG	123	4.2	118	3.5	120	6.1	101	0.9	105	4.7	106	1.2	6/*	5.2	102	3.7	106	2.1	/1*	3.0	106	3.9	109	2.6
2,4,6-INI	118	2.0	121	2.7	122	3.6	104	1.6	108	2.1	108	0.7	82	4.0	91	2.3	90	3.3	84	3.7	97	0.8	94	2.6
4-Am-2,6-DNT	105	1.9	115	1.8	117	1.4	87	2.1	111	1.8	112	1.8	61*	4.0	100	0.6	105	2.8	64*	3.2	112	10.9	108	1.6
2-AIII-4,0-DINI	116	2.4	112	3.3	115	0.8	115	1.3	112	1.5	115	1.1	ŏ∠ 74+	4.1	104	1.2	114	1.9	115	2.2	114	5.7	110	3.4
2,0-DNT	112	0.0	115	1.0	117	1.0	106	1.6	110	2.5	112	0.0	02	0.4	107	0.0	109	2.1	100	 	112	2.0	112	2.6
2,4-DNT	112	2.0	95	<u>1.0</u>	102	1.0 0.1	107	0.7	102	2. <u>1</u>	108	1.5	100	2.0 5 1	90	0.3	106	2.1	108	2.1 2.6	103	2.0	100	2.0
2-NT	113	2.0	96	<u> </u>	102	9.1	109	13	102	2.1	100	13	96	55	99	13	107	1.0	105	15	102	2.0	108	2.1
3-NT	113	0.1	96	15	105	9.2	107	0.8	101	39	107	0.7	97	55	98	12	105	15	105	2.6	103	11	110	2.0
PETN	116	4.6	111	4.6	118	3.9	109	1.9	109	2.4	110	1.9	91	5.5	102	2.5	111	5.4	99	2.8	110	2.4	115	1.9

Table 27. Extraction hold-time stability for three laboratory-spiked soil samples over 14 days. (AVG = average; SD = standard deviation)

Bold, italicized are IM compounds and degradation products. Green: 75%–125%; light green (\*): 64%–74% or 126%–135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

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	Recover	'ies (%)																						
	Soils Da	v 0 extrac	tion and	analveie			Soils Da	y 0 extrac	tion, Day	40 analys	is													
	5013 Da			anaiyəiə			Cold/Da	ırk			-		Room te	mperatur	e / Dark				Room te	emperatur	e / Light			
	Riley		Yuma		ASTM F	С	Riley		Yuma		ASTM F	0	Riley		Yuma		ASTM F	C	Riley		Yuma		ASTM FC	)
Compound	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NQ	<i>65</i> *	4.7	<i>65</i> *	4.1	<i>64</i> *	4.1	85	4.1	104	3.9	94	0.8	84	4.0	102	3.1	92	2.2	83	4.0	102	8.9	93	2.2
NTO (210nm)	<i>260</i> †	7.4	<i>73</i> *	1.5	78	4.9	<i>148</i> †	4.7	<i>149</i> †	1.4	<i>143</i> †	3.5	<i>149</i> †	5.9	<i>153</i> †	0.9	<i>146</i> †	4.1	<i>144</i> †	4.0	<i>150</i> †	1.7	<b>144</b> †	3.1
o-NBA (surrogate)	108	1.0	100	1.7	97	4.1	87	8.8	97	0.9	91	1.1	87	8.3	99	2.9	90	0.9	86	8.6	97	1.3	91	2.0
HMX	41†	0.6	71*	3.2	57v	14.2	35†	2.5	60*	2.5	39†	1.0	36†	3.8	60*	3.5	39†	0.9	35†	3.5	61*	2.5	39†	2.0
RDX	91	1.4	113	3.4	112	2.3	<i>60</i> *	1.7	<b>74</b> *	2.5	98	4.7	60*	1.9	<i>72</i> *	3.2	<i>51</i> †	0.9	<i>57</i> *	6.7	<i>71</i> *	2.8	<i>50</i> †	0.4
PA (210nm)	56†	0.5	81	3.3	59*	0.4	88	2.3	104	3.4	98	4.7	111	4.5	152†	3.1	143†	5.9	120	3.1	153†	3.6	142†	6.2
4-NP	<i>125</i> *	1.9	<i>125</i> *	1.0	<i>127</i> *	1.4	107	1.2	108	2.9	108	4.0	107	1.5	107	1.9	106	4.1	106	0.8	107	2.1	107	3.9
1,3,5-TNB	102	2.7	115	0.7	112	3.2	96	1.8	109	1.0	102	1.9	96	0.9	109	1.7	103	1.8	95	2.0	109	0.4	103	1.7
2,4-DNP	92	4.0	102	5.3	103	1.0	87	3.3	98	0.7	96	2.7	86	2.7	100	2.4	96	2.0	88	2.0	99	1.2	<i>95</i>	2.0
1,2-DNB (surrogate)	125*	4.3	124	2.0	124	5.1	115	1.3	117	1.7	115	2.6	119	0.3	117	0.7	119	1.8	118	0.2	119	0.9	118	2.0
1,3-DNB	117	1.7	118	0.6	120	1.2	106	2.2	109	1.0	110	1.9	107	0.8	110	1.7	110	1.9	106	1.5	110	0.6	110	1.3
2-NP	98	3.9	80	0.3	81	1.3	85	1.6	<i>71</i> *	2.1	<i>71</i> *	2.0	85	1.9	<i>72</i> *	1.4	<i>71</i> *	1.4	85	1.4	<i>71</i> *	0.3	<i>71</i> *	1.7
NB	101	1.2	113	2.3	115	0.7	89	1.6	103	0.9	102	2.1	92	0.3	105	2.1	102	2.4	91	0.7	105	0.4	103	1.3
3,5-DNA (210nm)	114	1.7	91	0.6	103	9.8	112	1.5	93	1.9	110	1.0	113	0.8	94	1.2	111	1.1	112	1.7	94	0.7	110	0.7
DNAN	103	2.8	104	1.0	109	3.3	76	0.3	82	1.0	78	3.0	77	0.8	82	1.3	79	1.6	77	0.6	82	0.9	80	1.9
Tetryl	114	3.9	111	0.6	112	0.7	94	2.4	52†	2.6	55†	1.4	35†	7.5	0†	0.0	0†	0.0	4†	0.3	0†	0.0	0†	0.0
NG	123	4.2	118	3.5	120	6.1	86	5.6	98	2.7	97	2.5	101	2.7	94	7.3	99	5.4	109	0.2	78	3.8	84	7.8
2,4,6-TNT	118	2.0	121	2.7	122	3.6	98	2.9	135*	4.7	130*	3.4	106	0.9	112	0.3	110	1.7	98	0.9	109	0.3	105	4.6
4-Am-2,6-DNT	105	1.9	115	1.8	117	1.4	94	1.4	112	1.1	113	1.9	87	1.3	99	1.1	105	10.0	97	9.1	99	1.1	99	0.6
2-Am-4,6-DNT	108	2.4	114	3.3	115	0.8	98	1.7	109	1.1	107	1.5	87	0.5	93	1.2	97	10.4	96	8.2	93	0.9	92	1.0
2,6-DNT	116	0.6	113	1.6	116	1.8	96	1.7	89	3.4	91	7.1	100	8.1	95	2.1	94	4.9	95	6.2	96	2.7	95	1.4
2,4-DNT	112	0.7	115	1.0	117	1.8	100	4.5	111	2.5	110	5.1	99	4.2	107	4.0	109	8.4	103	2.8	106	1.2	108	2.3
2-NT	112	2.0	95	3.3	102	9.1	104	1.9	92	1.5	104	1.5	107	5.5	91	1.8	103	1.5	106	2.1	93	2.3	103	1.1
4-NT	113	3.1	96	4.1	102	9.3	104	2.3	95	2.2	105	2.9	106	4.6	95	2.0	105	1.7	106	1.3	94	0.9	105	0.6
3-NT	113	0.9	96	1.5	105	9.2	106	3.2	92	1.0	103	0.9	106	2.2	92	1.1	104	0.7	105	2.4	92	1.4	104	0.4
PETN	116	4.6	111	4.6	118	3.9	106	1.9	110	1.0	110	3.8	110	2.1	110	2.6	110	2.0	111	2.3	111	3.5	111	2.6

Table 28 Analysis hold-time stability	v for three laboratory-spiked soil sam	noles over $40 \text{ days}$ (AVG = a	verage: SD = standard deviation)
Tuble 20. Analysis nota time stability	y for three laboratory spined son san		voluge, ob Standard deviation)

Bold, italicized are IM compounds and degradation products. Green: 75%–125%; light green (\*): 64%–74% or 126%–135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

In contrast to the NQ determinations described above, NTO recoveries were fairly stable from Day 0 to Day 40, with no storage condition– dependent trends. Only three compounds were detected at reduced levels at Day 40, including HMX, RDX, and Tetryl. HMX recoveries did not seem to depend on storage condition, while RDX recoveries were slightly lower for room temperature samples than those stored at colder temperatures. Tetryl recoveries were much lower for room temperature samples than for cold samples; however, even the cold samples were detected at 50% of the levels observed on Day 0. As in the soil pre-extraction stability study, the majority of compounds in the analysis hold-time stability study were recovered within current DoD QSM limits at both Day 0 and Day 40, including all other IM compounds tested.

Compound recovery results obtained for Day o of the water stability study also yielded information on compound volatility, as several compounds, specifically 2-NP, 3,5-DNA, 2,6-DNT and 2-/3-/4-NT, were recovered at lower levels on Day o than on Day 7. The compounds were detected at their usual levels within current DoD QSM limits on Day 7 but were nearly absent in samples from Day 0. Since samples prepared on Day 0 were stored for extraction on Day 7, if the compounds were lost from the original water samples, they would not have been detected in the Day 7 extracts. Because they were, the loss of these more volatile compounds from Day 0 extracts likely occurred when the extracts obtained by SPE on Day 0 underwent evaporative concentration under nitrogen. This phenomenon was also observed during SPE method development. Therefore, in general, and especially when any of these compounds are known or suspected to be present in a sample, evaporative concentration of extracts is not recommended.

### 3.10 Interlaboratory batch analysis

A cross-laboratory batch study was carried out at ERDC-EL and ERDC-CRREL using five water sources (Table 8) and six soil sources (Table 14). Multiple source samples were used to observe any potential matrix-related variability of extraction and analysis method efficiency. Both high-level (direct-injection) and low-level (SPE) water samples were used. Samples were extracted using the extraction methods and primary HPLC-UV method described in section 2.

Compound recovery results for direct-injection and SPE water extracts are shown in Table 29. For samples that underwent extraction and

preconcentration by SPE, the majority of compounds were recovered from all five water source samples at similar levels to those observed during SPE method development (Table 10, option B). As expected, NQ recoveries averaged approximately 50% for SPE extracts at ERDC-EL; however, much higher recoveries were obtained for NQ at ERDC-CRREL, so further investigation is merited. In general, at ERDC-EL, recoveries from reagent water tended to be slightly higher than usual, and at ERDC-CRREL, recovery of Tetryl from all water sources tended to be lower than usual. At both laboratories, it became apparent that using a detection wavelength of 315 nm rather than 210 nm for NTO might be preferred, to avoid positive interferences that appear at 210 nm for field samples. In addition, both laboratories observed low recoveries for NTO, 4-NP, 2-NP, NB (ERDC-EL only), 3,5-DNA and Tetryl (ERDC-CRREL only), and 4-Am-2,6-DNT in tap water samples (pH 9.2) that underwent SPE. Because of the consistency between laboratories, the high pH of the water samples, and the chemical properties of the compounds affected (acidic or unstable at high pH), the low recoveries may have been due to insufficient pH reduction during the second SPE elution step (using 2% HCl/MeOH).

In the same manner, during the direct injection cross-laboratory study, much lower recoveries for NTO, and in some cases *o*-NBA surrogate (ERDC-CRREL), were observed for field water samples than for reagent water. As this trend appeared to be pH related, with NTO recovery being lower for higher-pH water samples (see Table 27), further experiments were carried out in which the pH of samples to be analyzed by direct injection was adjusted, as discussed in section 3.4.

Compound recovery results for cross-laboratory soil studies are shown in Table 30. The majority of compounds were recovered within current DoD QSM limits, with some compounds having inconsistent recoveries from one soil to another, especially for NQ, NTO, *o*-NBA, and HMX. Some discrepancies were observed between the recoveries obtained by ERDC-EL and ERDC-CRREL, but for the most part, all compounds were recovered within current limits from most soils.

ERDC-EL	Recoveries	s (%)																						
					SPE (0.	10 mg/L	)				-				Direct i	njection	(2 mg/L)		-					
					Reager	nt water	Tap wat	ter	Housto Shippir Channe (North Morgar Point) v	n ng of n's water	Yazoo F water	River	Rayville well wa	e, LA Iter	Reager	it water	Tap wa	ter	Housto Shippir Channo (North Morgar Point) v	n ng of n's water	Yazoo F water	River	Rayville well wa	≽, LA iter
Compound	Blank (reagent water)	Blank spike (reagent water)	Matrix spike (HSCN)	Matrix spike duplicate (HSCN)	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NQ	0	23	32	36	50	8.6	1	0.5	47	3.0	50	<i>12.</i> 4	43	9.4	99	3.3	<i>92</i>	6.6	88	1.9	114	8.4	84	4.6
NTO (210nm)	0	127	113	136	154	17.0	27	14.8	158	18.1	212	42.5	129	17.2	115	0.9	3	0.2	11	0.5	15	0.7	2	0.1
NTO (315nm)	0	124	97	120	141	14.6	0	0.3	110	11.3	131	24.1	117	15.4	116	0.7	3	0.5	10	0.6	14	0.8	2	0.3
o-NBA (surrogate)	133	94	93	109	113	10.7	111	29.1	96	11.2	105	10.5	112	10.7	102	0.3	100	0.9	105	2.1	103	3.1	104	2.4
HMX	0	89	106	107	123	22.0	105	19.0	103	3.7	103	22.3	107	6.5	98	1.2	96	1.0	100	0.9	100	1.2	103	5.3
RDX	0	94	109	110	123	19.4	109	12.8	106	4.9	111	19.5	109	8.2	102	2.6	99	1.1	104	0.3	103	1.1	102	2.7
PA (210nm)	0	91	94	120	126	15.2	107	28.0	109	3.7	107	18.8	114	8.2	88	5.8	96	13.2	99	12.8	96	8.4	98	3.3
PA (315nm)	0	86	97	114	114	16.1	103	18.6	98	8.0	96	12.2	102	5.7	89	1.8	91	0.9	94	0.4	92	1.1	92	1.3
4-NP	0	86	103	105	123	23.6	39	5.3	103	0.4	106	17.2	105	4.1	94	1.2	94	0.8	97	1.0	95	2.0	94	2.0
1,3,5-TNB	0	95	103	110	140	22.8	116	23.1	109	1.2	104	19.0	118	8.8	100	1.4	100	0.8	105	2.2	103	4.2	103	2.0
2,4-DNP	0	89	104	101	122	20.6	104	22.8	101	6.1	100	15.6	108	5.7	101	2.5	101	1.4	105	1.8	104	3.6	100	2.2
1,2-DNB (surrogate)	110	89	106	107	124	27.6	97	23.3	104	5.6	108	21.1	113	9.1	97	1.9	96	0.5	99	2.3	99	1.0	99	2.5
1,3-DNB	0	97	100	114	132	12.5	114	25.0	109	8.0	105	10.7	118	13.5	100	0.8	100	1.2	103	0.3	102	0.5	101	1.1
2-NP	0	93	100	105	131	17.4	20	3.3	109	2.1	106	10.4	114	2.9	104	0.2	97	1.0	108	0.9	107	0.6	105	0.9
NB	0	91	101	108	133	24.1	52	7.2	105	1.9	113	21.0	114	9.5	99	1.3	99	0.9	102	0.9	101	0.5	99	0.2
3,5-DNA (210nm)	0	95	103	114	128	9.5	90	10.9	106	5.1	102	3.8	117	8.6	103	1.0	104	2.2	108	1.3	104	1.3	104	0.3
3,5-DNA (254nm)	0	96	102	110	130	9.4	108	16.6	110	5.9	103	2.3	117	10.1	104	0.7	104	1.1	107	0.9	106	0.2	105	0.3
DNAN	0	104	102	121	132	1.6	109	20.0	107	8.7	106	4.8	128	8.0	105	2.8	104	2.3	109	1.4	105	0.6	107	3.4
Tetryl	0	95	91	93	132	23.3	75	15.5	88	3.4	85	19.6	96	10.8	91	1.9	85	5.9	97	1.0	94	1.8	98	2.1
NG	0	94	105	104	129	25.0	127	34.8	100	7.5	113	21.4	141	7.3	94	2.7	97	6.9	101	5.4	96	2.4	98	3.3
2,4,6-TNT	0	94	101	108	129	19.3	109	29.2	99	4.9	95	15.1	122	12.9	100	4.8	103	3.6	105	2.8	102	3.5	100	1.0
4-Am-2,6-DNT	0	92	96	97	143	31.1	5	1.4	95	6.1	116	34.3	112	12.1	97	1.3	95	1.5	103	1.5	100	0.2	99	2.2
2-Am-4,6-DNT	0	96	99	106	136	22.2	106	19.7	100	8.3	109	15.2	118	15.0	97	2.0	99	1.2	102	3.0	99	1.6	98	3.1
2,6-DNT	0	92	96	113	117	6.0	91	23.9	94	4.9	89	8.3	103	11.2	97	1.3	96	2.7	104	1.9	100	3.2	96	1.0
2,4-DNT	0	103	99	114	121	9.5	113	22.4	113	6.9	100	1.3	112	9.9	101	1.7	102	1.6	105	1.7	104	0.9	103	0.8
2-NT	0	95	96	106	112	4.2	105	15.1	104	7.1	94	6.8	114	11.7	102	0.4	104	2.8	105	1.1	104	0.8	100	1.4
4-NT	0	96	93	106	108	13.9	95	24.6	107	7.3	86	10.6	109	11.4	100	0.7	101	1.7	104	5.0	104	1.8	101	4.3
3-NT	0	92	92	101	110	10.9	102	27.3	108	8.0	94	5.9	110	13.4	99	1.4	99	3.0	101	0.2	99	0.7	100	2.7
PETN	0	87	101	102	117	20.5	94	14.6	101	11.2	100	26.7	97	8.6	94	3.5	95	4.3	101	2.6	96	5.6	95	3.8

Table 29. Cross-laboratory batch analysis tests using five different water sources and the SPE method developed under SERDP ER-2722. (AVG = average

age:	SD =	= stan	dard	dev	iation	۱
чвс,	30-	- stai	lualu	uev	auon	,

ERDC-CRREL	Recoveries	s (%)																						
					SPE (0.	.10 mg/L	.)								Direct i	njection (	(2 mg/L)							
					Reager	nt water	Tap wat	ter	Housto Shippir Channe (North Morgar Point) V	n ng el of Nater	Yazoo I water	River	Rayville well wa	e, LA ter	Reager	nt water	Tap wat	ter	Housto Shippir Channe (North Morgar Point)	n ng el of n's water	Yazoo F water	River	Rayville well wa	≽, LA ter
Compound	Blank (reagent water)	Blank spike (reagent water)	Matrix spike (Yazoo)	Matrix spike duplicate (Yazoo)	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NQ	0	92	<i>73</i> *	82	86	7.6	<i>0</i> †	0.0	82	4.7	<i>67</i> *	10.4	<i>62</i> *	3.0	76	15.6	88	2.2	104	2.9	106	1.8	103	3.0
NTO	0	<i>169</i> †	<i>161</i> †	<b>164</b> †	124	2.4	<i>0</i> †	0.0	<i>140</i> †	17.4	<b>144</b> †	12.6	<i>138</i> †	6.4	120	26.2	<i>2</i> †	0.5	<i>8</i> †	1.6	<b>11</b> †	2.0	<b>6</b> †	1.4
o-NBA (surrogate)	117	0†	0†	0†	0†	0.0	0†	0.0	0†	0.0	0†	0.0	0†	0.0	286†	159.3	30†	24.7	62*	50.6	23†	19.4	16†	13.0
HMX	0	125*	108	120	115	5.4	113	3.1	119	4.9	110	9.6	122	6.2	100	3.4	98	3.7	97	3.2	99	3.5	96	3.2
RDX	0	118	106	117	83	45.9	111	1.3	116	8.5	103	7.6	123	2.6	115	4.8	117	4.9	107	6.8	105	6.6	106	7.9
PA	0	116	109	118	103	3.5	109	0.4	119	12.4	100	10.3	<u>124</u>	4.6	<i>98</i>	0.9	99	1.6	97	0.0	98	0.5	96	0.3
4-NP	0	116	96	110	107	5.0	<i>33</i> †	1.2	<u>112</u>	0.9	105	2.2	114	2.2	101	3.1	97	3.3	<u>98</u>	2.8	<u>99</u>	3.1	<u>96</u>	2.9
1,3,5-INB	0	116	89	108	107	5.9	106	4.7	101	5.4	65*	11.0	115	6.1	102	4.9	99	5.1	98	4.6	100	4.8	97	4.6
	0	126*	104	119	122	<i>9.</i> 6	126*	2.8	122	1.4	115	2.7	125	2.5	99 001+	8.0	164+	4.0	<u>91</u>	<b>5.2</b>	162+	4.3	150+	3.9
	04	116	105	110	103	6.5 5.1	99	3.2	102	4.8	93	9.9	102	5.4	231 <u>7</u>	22	164T	2 2 2	1027	200	1037	2.0	1597	2.1
2.1/0	0	112	102	102	100	5.1	10+	4.0	103	5.0	95	10.4 0.6	107	3.1 26	112	3.3 12 /	<u> </u>	120	104	121	105	3.2 121	102	 
	0	113	102	107	99	4.9	96	4.0	106	53	90	3.0 8.8	101	37	101	28	99	32	98	26	<u> </u>	27	97	29
3.5-DNA	0	115	104	108	103	6.0	44+	2.0	105	5.0	107	9.3	106	5.9	100	2.0	98	3.0	98	2.0	99	2.1	96	2.5
DNAN	0	117	106	109	100	4.5	98	3.9	108	7.2	<u> </u>	10.4	102	3.0	 99	<u> </u>	98	1.4	97	1.2	98	1.7	96	1.6
Tetryl	0	118	64*	78	108	11.1	55†	3.6	60*	18.8	44†	3.4	42†	23.1	131*	54.4	83	42.1	102	39.1	101	38.2	106	39.0
NG	0	124	95	112	104	6.4	113	11.6	115	13.3	117	10.6	128*	14.7	98	20.4	88	8.6	90	5.7	98	15.2	107	13.5
2,4,6-TNT	0	114	99	111	100	4.2	96	3.7	103	8.4	78	10.5	114	4.4	103	0.9	103	2.7	101	2.3	103	2.3	99	1.9
4-Am-2,6-DNT	0	113	98	104	102	6.6	5†	1.6	91	4.5	95	8.9	104	5.9	100	2.5	98	3.1	98	2.5	99	2.7	96	2.8
2-Am-4,6-DNT	0	115	104	108	102	5.7	61*	2.2	105	5.9	97	9.7	105	6.2	101	3.4	99	3.7	98	3.3	99	3.5	97	3.4
2,6-DNT	0	114	108	113	93	5.3	86	3.0	101	7.6	86	8.9	93	1.4	101	2.8	99	3.5	99	2.7	100	3.1	96	2.4
2,4-DNT	0	114	109	114	95	5.2	90	3.5	105	8.9	89	9.5	96	4.1	101	2.8	99	3.8	98	1.7	100	3.4	96	2.9
2-NT	0	111	103	106	94	4.0	92	3.5	102	5.7	90	8.6	97	2.0	100	2.6	98	2.8	97	2.4	99	2.6	96	2.6
4-NT	0	113	104	107	96	3.6	93	3.1	103	7.1	91	8.6	96	1.5	101	3.9	99	3.6	98	3.3	99	3.7	97	3.6
3-NT	0	117	104	107	102	4.6	101	3.5	113	7.9	99	11.2	105	2.4	99	2.0	98	2.2	97	1.8	98	1.8	96	2.0
PETN	0	115	96	111	108	8.2	103	4.1	107	4.1	100	3.5	114	3.6	101	3.4	99	4.2	98	4.2	100	4.0	95	3.0

Bold, italicized are IM compounds and degradation products. Green: 75%-125%; light green (\*): 64%-74% or 126%-135%; red (†): <64% or >135% (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

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ASTM Fat Clay

### Matrix spike Blank Blank spike Matrix spike duplicate AVG AVG AVG AVG AVG AVG (*n* = 3) SD (*n* = 3) SD (n = 3)SD (*n* = 3) SD (n = 3)SD (*n* = 3) SD Compound (hydromatrix) (hydromatrix) (Aberdeen) (Aberdeen) NQ 94 90 0.7 93 1.7 4.0 1.4 1.8 0 *99* 100 1.6 85 79 98 82 124 **74**\* NTO (210nm) 0 76 *129*\* 3.1 104 0.9 97 0.5 76 0.8 *72*\* 1.8 105 2.2 120 NTO (315nm) 0 *69*\* *68*\* *64*\* 0.9 *66*\* 1.7 98 1.5 84 2.0 100 1.2 90 0.7 o-NBA (surrogate) 116 108 100 97 95 75\* 2.2 102 0.7 86 105 1.7 93 1.1 1.0 1.2 106 90 2.6 2.7 86 2.6 1.0 89 83 70\* 1.0 55† 56† HMX 0 91 1.0 96 94 106 96 2.0 RDX 0 89 3.8 76 1.2 93 1.5 66\* 2.2 1.5 65\* 92 PA (210nm) 0 86 84 6.2 2.4 101 1.5 87 7.5 90 4.4 82 4.7 *99* 82 98 94 92 88 79 98 89 0.9 PA (315nm) 0 2.3 85 1.2 94 2.0 2.9 1.4 92 0 100 97 95 91 1.4 90 2.7 94 1.5 86 1.0 99 2.0 2.0 4-NP 1,3,5-TNB 107 98 96 93 82 1.5 92 1.5 0 1.1 84 1.0 100 1.1 0.4 106 100 98 80 6.7 91 2.7 2.4-DNP 0 104 *95* 1.6 82 1.8 *99* 1.1 2.0 103 98 1,2-DNB (surrogate) 99 0.5 103 98 121 0.6 96 0.5 100 2.3 95 1.8 103 2.2 97 105 103 100 0 99 0.3 98 0.7 102 0.8 94 0.9 107 1.4 99 0.7 1,3-DNB 109 103 101 94 1.6 85 0.9 0 2-NP 0.4 90 0.1 84 1.0 91 1.7 93 102 100 NB 0 107 97 0.7 86 0.3 100 1.5 77 0.8 105 1.2 96 0.5 3,5-DNA (210nm) 0 113 102 101 93 2.7 92 1.3 77 3.8 93 3.2 90 0.5 92 0.9 0 101 99 93 0.3 0.9 76 93 88 0.7 91 1.0 3.5-DNA (254nm) 111 91 1.6 1.5 108 104 94 112 101 0.1 0 113 101 1.8 100 3.0 105 4.0 1.5 1.6 DNAN 95 96 91 90 84 89 3.4 Tetryl 0 2.8 97 0.8 89 1.7 1.0 99 1.1 90 96 90 2.6 99 4.7 NG 0 110 94 4.7 94 2.1 97 2.5 1.6 107 113 94 94 2.1 94 2.8 2,4,6-TNT 0 95 4.0 94 0.4 99 2.3 86 3.2 107 98 101 98 96 73\* 95 4-Am-2,6-DNT 0 1.5 87 1.7 99 1.0 0.8 103 1.3 1.2 2-Am-4,6-DNT 0 101 102 99 98 2.3 94 2.0 2.4 1.5 103 2.6 97 0.6 101 86 107 102 102 0 104 102 4.5 102 5.1 107 4.4 99 3.7 109 4.0 4.4 2,6-DNT 105 95 89 2.8 96 2,4-DNT 0 102 96 0.8 96 0.9 100 1.2 0.6 106 0.2 100 89 0 107 101 94 1.2 92 93 1.2 0.8 92 2-NT 0.3 80 1.7 1.4

102

99

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92

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2.7

1.6

108

104

# Table 30. Cross-laboratory batch analysis tests using six different soil sources and the two-stage SE method developed under SERDP ER-2722.(AVG = average; SD = standard deviation)

Spiked soil (20 mg/kg)

Jefferson

Aberdeen

Riley

Yuma

Memphis

82

76

2.3

1.7

91

90

3.9

2.8

89

86

1.4

1.8

88

88

1.1

1.0

ERDC-EL

Recoveries (%)

0

0

4-NT

3-NT

PETN	0	99	105	92	94	3.1	94	3.1	96	4.4	88	2.7	99	7.4	98	0.9
	T															
ERDC-CRREL	Recoveries (%)	-														
					Spiked s	soil (20 m	g/kg)									
					Aberdee	en	Jefferso	n	Memph	is	Riley		Yuma		ASTM F	at Clay
Compound	Blank (hydromatrix)	Blank spike (hydromatrix)	Matrix spike (Yuma)	Matrix spike duplicate (Yuma)	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD	AVG ( <i>n</i> = 3)	SD
NQ	0	<i>186</i> †	120	117	<i>148</i> †	<i>3.5</i>	89	3.7	<i>133</i> *	23.6	<i>147</i> †	7.3	108	3.9	87	2.3
NTO	0	58*	<i>38</i> †	81	<i>29</i> †	3.1	<b>74</b> *	9.5	<i>37</i> †	7.7	<i>40</i> †	6.9	76	10.7	<i>53</i> †	5.4
o-NBA (surrogate)	119	123	90	97	44†	2.6	92	5.3	90	41.4	44†	3.1	98	2.6	95	2.6
HMX	0	126*	90	90	96	1.0	85	1.7	101	4.5	73*	0.5	91	0.2	71*	2.1
RDX	0	140†	115	118	114	3.5	103	2.7	120	7.5	99	2.0	112	0.9	97	2.5
PA	0	<i>141</i> †	123	127*	117	5.3	110	2.3	122	9.0	100	1.7	115	1.4	107	3.3
4-NP	0	131*	122	131*	119	5.7	111	2.6	120	10.3	113	1.7	117	1.4	110	3.9
1,3,5-TNB	0	125*	120	129*	116	5.3	100	2.4	119	10.5	102	1.8	117	1.4	105	3.9
2,4-DNP	0	114	97	105	109	4.0	94	3.7	103	6.2	88	0.7	103	1.4	92	2.2
1,2-DNB (surrogate)	101	128*	121	129*	121	6.3	112	1.8	118	10.8	112	1.6	116	2.0	110	3.4
1,3-DNB	0	125	119	128*	118	5.5	110	1.3	113	11.8	111	0.9	115	2.9	109	4.2
2-NP	0	117	98	100	108	3.2	<i>98</i>	0.2	87	7.6	103	0.2	95	3.3	88	1.1
NB	0	133*	105	112	109	3.9	100	0.5	84	8.6	108	0.4	97	3.2	96	2.8
3,5-DNA	0	127*	119	130*	117	5.6	101	2.6	118	10.8	95	1.8	116	1.4	108	3.9
DNAN	0	121	120	130*	119	5.8	110	2.2	119	11.1	110	1.7	117	1.1	109	4.0
Tetryl	0	35†	58*	133v	103	20.0	119	16.3	73*	33.3	121	4.7	99	14.4	75	4.7
NG	0	152†	111	121	114	9.1	99	0.4	116	22.3	101	8.0	103	4.0	123	17.0
2,4,6-INI	0	137†	123	123	113	6.8	106	1.3	106	13.6	100	1.3	111	4.2	105	4.0
4-Am-2,6-DNT	0	127(*	120	131(*	118	5.9	103	2.2	120	10.6	92	2.4	116	1.3	111	4.0
2-Am-4,6-DNT	0	124	120	130(*	120	5.8	110	2.5	121	10.9	105	2.0	11/	1.2	111	3.9
2,6-DNT	0	0†	121	131(*	11/	6.3	109	1.6	112	12.7	112	1.0	115	3.9	109	5.4
2,4-DNT	0	111	120	131(*	11/	6.6	109	2.2	112	13.3	110	1.4	114	3.9	109	4./
2-NT	0	133(*	108	115	110	3.9	102	0.1	88	10.6	109	0.2	99	4.2	97	3.8
4-NT	0	129(*	107	113	114	4.1	102	0.8	84	12.2	107	0.6	98	4.9	93	4.3
3-NT	0	135(*	106	112	109	4.0	100	1.0	81	11.3	105	0.8	95	5.1	92	3.3
PETN	0	123	124	132(*	120	6.9	112	2.3	122	12.1	115	2.9	117	2.0	112	4.8

 Bold, italicized are IM compounds and degradation products.

 Green: 75%-125%; light green (\*): 64%-74% or 126%-135%; red (†): <64% or >135%

 (DoD QSM Ver5.2 for Solids—lowest LCL: 4-Am-2,6-DNT, 64%; highest UCL: Tetryl, 135%)

# 4 Conclusions and Implications for Future Research and Implementation

The primary objective of this project was to fulfill the requirements of SERDP SON ERSON-17-02 by developing new methods of extraction, preconcentration, and analytical separation and quantitation of 17 legacy as well as 7 IM compounds, daughter products of IM, and other munition compounds absent from USEPA Method 8330B. The goal was to produce a single, standardized method for simultaneous analysis of legacy and IM compounds in environmental matrices. HPLC-UV and HPLC-MS methods were developed as well as extraction methods for waters (direct-injection and SPE), soils (two-stage SE), and tissues (one-stage SE). An interference reduction procedure was also developed for tissues. The development of these methods should produce cost savings with a minimal increase in environmental monitoring efforts and no modification of existing sample collection procedures.

During this project, two separate HPLC-UV column separation methods were developed, enabling orthogonal confirmation of analyte concentration down to low  $\mu$ g/L levels in water, soil or sediment, and tissue samples. Preconcentration of water samples using SPE can lower analyte detection levels to sub  $\mu$ g/L levels with sufficient concentration factors applied. A less sensitive HPLC-MS scan mode method was developed, which can be used as an alternative to a secondary column confirmation method. Sample preparation and solvent gradients were refined for optimal method performance. Furthermore, the SPE and SE methods (and interference reduction method for tissues) developed for this project yield compound recoveries for all analytes of interest and surrogates within currently accepted limits, with the recoveries for many of the compounds consistently better than current methods.

The development of these extraction and analysis methods to simultaneously analyze environmentally copresent legacy and IM compounds was necessary to enable future monitoring of co-occurring components at firing ranges, demilitarization facilities, manufacturing facilities, and environmental sites, where munitions are tested, produced, and detected (Felt et al. 2013; Walsh et al. 2014). The primary and secondary HPLC-UV methods reduce total analysis time by approximately 25% versus using four different analytical methods as has previously been required (two methods for legacy and two different methods for IM). In addition, the labor and supply cost savings because of coextraction and sample preparation is at least 50%, as only a single extraction procedure and analytical sample preparation are required versus up to four (including confirmation analyses), because the sample preparation for the primary and secondary methods are the same. Furthermore, the products of this work will contribute to the ability to conduct fate and transport studies for IM compounds by providing a standardized method for quantitation of IM and legacy materials needed to determine long-term effects.

The method development work executed under SERDP ER-2722 will continue under ESTCP ER19-D1-5078 (started February 2019) to validate the methods. The validation will consist of round-robin comparison studies, carried out with the cooperation of USEPA, with the end goal of producing an addendum to the current USEPA Method 8330B. Eight participants, including DoD laboratories (ERDC-EL and -CRREL), another federal laboratory (USGS National Water Quality Laboratory, NELAP certified), a munition manufacturing laboratory (American Ordnance), and several ELAP- or NELAP-certified, or both, commercial laboratories are participating in the validation studies. The standardization and publication of the validated methods will enable commercial and DoD laboratories to perform the standardized methodology on a broad commercial scale.

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## **List of Acronyms**

ACN	acetonitrile				
AEC	anion exchange capacity				
4-Am-2,6-DNT	4-amino-2,6-dinitrotoluene				
2-Am-4,6-DNT	2-amino-4,6-dinitrotoluene				
ASTM	American Society for Testing and Materials				
BIP	blow in place				
CEC	cation exchange capacity				
CH₃	methyl				
Comp-B	Composition-B				
CRREL	Cold Regions Research and Engineering Laboratory				
3,5-DNA	3,5-dinitroaniline				
DNAN	2,4-dinitroanisole				
1,2-DNB	1,2-dinitrobenzene				
1,3-DNB	1,3-dinitrobenzene				
2,4-DNP	2,4-dinitrophenol				
2,4-DNT	2,4-dinitrotoluene				
2,6-DNT	2,6-dinitrotoluene				
DoD	Department of Defense				
EL	Environmental Laboratory				
ER	Environmental Restoration				
ERDC	Engineer Research and Development Center				
ESTCP	Environmental Security Technology Certification Program				
FA	formic acid				
GAC	granular activated carbon				
Н	hydrogen atom, proton				
HOAc	acetic acid				
HCI	hydrochloric acid				
НМХ	octahydro-1,3,5-7-tetranitro-1,3,5,7-tetrazocine				
HPLC	high-performance liquid chromatography				
IM	insensitive munition				
LCL	lower control limit				
LDR	linear dynamic range				
MDL	method detection limit				
MeOH	methanol				
meq	mole equivalent				
MS	mass spectrometer or mass spectrometry				
NB	nitrobenzene				
ND	not detected				
NG	nitroglycerine				
2-NP	2-nitrophenol				
4-NP	4-nitrophenol				
NQ	nitroguanidine				

2-NT	2-nitrotoluene
3-NT	3-nitrotoluene
4-NT	4-nitrotoluene
NTO	3-nitro-1,2,4-triazol-5-one
o-NBA	ortho-nitrobenzoic acid
PA	picric acid
PETN	pentaerythritol tetranitrate
PTFE	polytetrafluoroethylene
QSM	quality systems manual
RDX	hexahydro-1,3,5-trinitro-s-triazine
SERDP	Strategic Environmental Research and Development Program
SPE	solid phase extraction
SE	solvent extraction
Tetryl	N-methyl-N-(2,4,6-trinitrophenyl) nitramide
TFA	trifluoroacetic acid
TIC	total ion chromatogram
1,3,5-TNB	1,3,5-trinitrobenzene
2,4,6-TNT	2,4,6-trinitrotoluene
TNT	trinitrotoluene
TDS	total dissolved solids
TOC	total organic carbon
UCL	upper control limit
UL	upper limit
USEPA	United States Environmental Protection Agency
UV	ultraviolet
UXO	unexploded ordnance
v/v	volume/volume
vol/vol	volume/volume

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No standard method exists for determining levels of insensitive munition (IM) compounds in environmental matrices. This project resulted in new methods of extraction, analytical separation and quantitation of 17 legacy and 7 IM compounds, daughter products of IM, and other munition com-pounds absent from USEPA Method 8330B. Extraction methods were developed for aqueous (direct-injection and solid-phase extraction [SPE]), soil, sediment, and tissue samples using laboratory-spiked samples. Aqueous methods were tested on 5 water sources, with 23 of 24 compounds recovered within DoD QSM Ver5.2 limits. New solvent extraction (SE) methods enabled recovery of all 24 compounds from 6 soils within QSM limits, and a majority of the 24 compounds were recovered at acceptable levels from 4 tissues types. A modified chromatographic treatment method removed analytical interferences from tissue extracts. Two orthogonal high-performance liquid chromatography-ultraviolet (HPLC-UV) separation methods, along with an HPLC–mass spectrometric (HPLC-MS) method, were developed. Implementing these new methods should reduce labor and supply costs by approximately 50%, requiring a single extraction and sample preparation, and 2 analyses rather than 4. These new methods will support environmental monitoring of IM and facilitate execution of risk-related studies to determine long-term effects of IM compounds.							
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