1	SUPPORTING INFORMATION			
2	Development of Certified Reference Materials for Diarrhetic Shellfish Poisoning Toxins.			
3	Part 1: Calibration Solutions			
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23 SI-1: Structure Characterization of 19-epi-DTX2 by NMR

1D-¹H spectrum showed resonances consistent with DTX2, notably the vinyl protons at lower 24 field than the OH resonance of the solvent and the proper number of methyl resonances (Figure 25 S1). Proton and ¹³C chemical shifts are shown in Fig. S2A and S2B respectively. The proton 26 27 resonances were identical to published data for DTX2 (1) except for significant deviations in chemical shift between positions C15 and C23, with a maximum difference at C18 and C23, as 28 shown in Fig. S3 for both the ¹H and ¹³C. This is consistent with the published reports of the 19-29 epimer of OA (2). ¹³C resonances were determined from the HSOC (Fig. S4) and HMBC (Fig. 30 31 S5) spectra. Analysis of the HSQC spectrum yielded 5 methyl resonances, 16 methylene resonances and 16 methine resonances. The HMBC spectrum gave chemical shifts for a further 7 32 quaternary carbons. 33 Interpretation of the 2D-¹H TOCSY spectrum (Fig. S6) revealed the following spin systems: 34 1. 4.06, 3.36, 1.93, 1.80, 1.70, 1.67, 1.35 35 2. 5.89, 5.66, 4.53, 3.88, 2.47, 2.39, 2.07, 2.04, 1.91, 1.77, 1.66, 1.13 36 37 3. 4.16, 3.66, 3.04, 2.03, 1.93, 1.83, 1.53 4. 4.08, 3.97, 3.41, 1.82, 1.73, 1.67, 1.60, 1.53, 1.42, 1.37, 1.27, 1.17, 0.96 38 5. 3.69, 3.51, 2.17, 1.78, 1.61, 1.30, 1.26, 0.99 39

Isolated resonances were also observed at 1.73, 1.31 ppm. Vinyl resonance at 5.26 ppm also
showed TOCSY correlations to the methyl resonance at 1.73 ppm and the resonances at 2.04 and
1.77 ppm. The vinyl resonances at 5.37 and 5.05 ppm showed TOCSY correlations to resonances
at 4.16, 3.66 and 3.04 ppm. Analysis of the COSY (Fig. S7), TOCSY (Fig. S6) and HMBC (Fig.
S5) spectra revealed the same connectivity as in DTX2. Figure S8 shows the TOCSY spin

45 systems as bold bonds as well as the key HMBC correlations used to connect the spin systems,







Figure S2. Structure of 19-*epi*-DTX2 showing 1 H (A) and 13 C (B) chemical shift assignments.



- **Figure S3**. Plot of the difference in 13 C chemical shift (top) and 1 H chemical shift (bottom)
- between DTX2 and 19-*epi*-DTX2. Note the largest difference is at the protons and carbons of
 C18 and C23.



Figure S4. ¹H-¹³C HSQC spectrum of 19-*epi*-DTX2 acquired with multiplicity sorting.





Figure S5. ${}^{1}\text{H}{}^{-13}\text{C}$ HMBC spectrum of 19-*epi*-DTX2 acquired with 60 ms mixing time time in CD₃OD.









Figure S7. DQF-COSY spectrum of 19-*epi*-DTX2 in CD₃OD.



Figure S8. TOCSY spin systems shown as bold bonds. Red arrows indicate the key HMBC

- correlations (pointing from carbon to proton) used to connect the spin systems. The blue arrow
- 82 indicates the COSY correlation between the pendant vinyl protons H41 at 5.37 and 5.05 ppm and
- H24 at 4.16 ppm. The numbering of carbons in the molecule is consistent with Hu *et al*, 1992.



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Figure S9. High-resolution full scan (A) and product ion (B) mass spectra of the $[M-H]^-$ ion, m/z803, of OA in CRM-OA-d measured on a Thermo Exactive Orbitrap mass spectrometer equipped with a heated electrospray ionization probe. Data acquired in negative ion mode with a 91 -2.7 kV spray voltage, +360 °C capillary temperature, and +250 °C heater temperature. HCD 92 collision energy in (B) was 60 V.



Figure S10. High-resolution full scan (A) and collision induced dissociation (B) mass spectra of the $[M-H]^-$ ion, m/z 817, of DTX1 in CRM-DTX1-b measured on a Thermo Exactive Orbitrap mass spectrometer equipped with a heated electrospray ionization probe. Data acquired in negative ion mode with a -2.7 kV spray voltage, + 360 °C capillary temperature, and + 250 °C heater temperature. HCD collision energy in (B) was 60 V.





Figure S11. High-resolution full scan (A) and collision induced dissociation (B) mass spectra of the $[M-H]^-$ ion, m/z 803, of DTX2 in CRM-DTX2-b measured on a Thermo Exactive Orbitrap mass spectrometer equipped with a heated electrospray ionization probe. Data acquired in negative ion mode with a -2.7 kV spray voltage, + 360 °C capillary temperature, and + 250 °C heater temperature. HCD collision energy in (B) was 60 V.



Figure S12. Stability of OA solutions in DMF at a range of temperatures and environmentalconditions.



117 Figure S13. High resolution mass spectra of the OA degradation product observed in a DMF

solution exposed to light and oxygen at room temperature for one week. The product was

tentatively identified as the ketone analog formed by elimination of formic acid.



Figure S14. Stability of OA solutions in MeOH at a range of temperatures and environmentalconditions.

133 SI-4: Additional Selected Reaction Monitoring Conditions

Table S1: Selected Reaction Monitoring settings used for quantitative LC-MS/MS analysis ofDSP toxins.

Analyte	<i>m/z</i> Precursor > <i>m/z</i> Product	Declustering Potential (V)	Collision Energy (V)
0.4	803.5>255.1	-80	-65
UA	803.5>113.1	-80	-85
DTV1	817.5>255.1	-70	-70
DIAI	817.5>113.1	-70	-90
DTVI	803.5>255.1	-80	-65
DIAZ	803.5>113.1	-80	-85
Parameter	Setting	Parameter	Setting
Dwell (msec)	125	CUR (psi)	20
GS1 (psi)	50	GS2 (psi)	50
IS (V)	-4500	CAD (V)	-3
EP (V)	-10	CXP (V)	-6

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138 SI-5: LC-FLD of ADAM Derivatized DSP toxin CRMs

139 SI-5.1 ADAM Derivatization

Reactions were carried out under yellow or reduced light. The ADAM precursor, 140 9-anthraldehyde hydrazone, was prepared from 9-anthralaldehyde and hydrazone hydrate 141 according to the method of Nakaya et al. (3). Solutions of 9-anthraldehyde hydrazone (35 mM), 142 *N*-chlorosuccinimide (35 mM) and quinuclidine (70 mM) were prepared in THF. To prepare the 143 ADAM reagent, equivalent volumes (500 µL) of all three solutions were mixed in an amber 144 glass vial and allowed to react at ambient temperature for 1 hour prior to sample derivatization. 145 A 35 μ L aliquot of the sample was placed in a 1.5 mL amber vial with 100 μ L of the ADAM 146 147 reaction reagent. The reaction mixture was placed in a sonication bath (Model 1510, Branson, Danbury, CT, USA) for 10 min at 37 °C and then transferred to a mutlitherm thermal reaction 148 149 chamber (Model H5000-HC, Benchmark Scientific, Sayreville, NJ, USA) for 2.5 hours at 37 °C. Samples were sonicated for an additional 10 min before drying down using a rotary vacuum drier 150 (Model SPD2010-220, Savant Instrument Inc., Holbrook, NJ, USA) and then reconstituted in 151 300 µL n-hexanes: chloroform (1:1) for SPE. 152

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154 SI-5.2 Solid Phase Extraction of Derivatized Samples

Derivatized samples were cleaned up using the SPE procedure published previously (4). Glass 155 SPE tubes (7 mL) equipped with a Teflon frit were packed with 0.5 g of activated silica and 156 placed on a vacuum manifold. The columns were conditioned with 6 mL of chloroform and 3 157 158 mL n-hexanes: chloroform (1:1). The columns were not allowed to become dry from this point on. The ADAM derivatized samples were transferred onto the column and vials were rinsed with 159 160 $2 \times 300 \,\mu\text{L}$ of n-hexanes: chloroform (1:1) and passed slowly through the column at approx. 1 drop/sec. The columns were then washed with 5 mL of n-hexanes: chloroform (1:1) followed by 161 5 mL of chloroform (containing 1.15 % ethanol). Samples were eluted with 5 mL of MeOH: 162 163 chloroform (1.5:8.5). Eluted fractions were evaporated under nitrogen and reconstituted with 2 mL of MeOH. 164

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169 *SI-5.3 Liquid Chromatography with fluorescent detection (LC-FLD)*

170 LC-FLD analysis was carried out on an Agilent 1200 series LC quipped with a fluorescence 171 detector (model no. G1321A) operated with excitation/emission wavelengths of 254/412 nm. A 172 50 x 2 mm Luna C18 HST (2.5 μ m) (Phenomenex, Torrence, CA USA) was eluted isocratically 173 with mobile phase of (A) DIW and (B) 95% acentonitrile each containing 50 mM formic acid 174 and 2 mM ammonium formate at 70% B and 300 μ L/min. The column temperature was 20 °C 175 and an injection volume of 5 μ L was used. 176



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Figure S15: LC-FLD of ADAM derivatized DSP toxin calibration solutions for OA (b), DTX1
(c), DTX2 (d) and a derivatization reagent blank (a).

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183 SI: REFERENCES

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