

Supporting Information
for the Paper

**Mapping of arsenic species and identification of a novel arsenosugar
in giant clams *Tridacna maxima* and *Tridacna derasa* using advanced
mass spectrometric techniques**

Volker Nischwitz and Spiros A. Pergantis*

University of Crete, Department of Chemistry, Environmental Chemical Processes Laboratory,
71003 Voutes-Heraklion, Crete, Greece

* Corresponding author; Tel.: 0030 2810 545084; Fax.: 0030 2810 545001; e-mail:
spergantis@chemistry.uoc.gr

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The abbreviations for the arsenosugars are composed of 4 parts [Nischwitz, V., Pergantis, S.A., <i>Anal. Chem.</i> , 2005, 77 , 5551-5563]:	
1. The number of methyl-groups bound to the arsenic atom: DM for Dimethyl, TM for Trimethyl	
2. Only for the thio-arsenosugars: Thio	
3. The common core for all β -ribofuranosides: AsSugar	
4. The characteristic functional group in the aglycone: e.g. Glycol	
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Standards of arsenic species:

Cacodylic acid (DMA) (puriss., >99%) and monosodium acid methane arsonate (MMA) (sesquihydrate) were obtained from Fluka, Buchs, Switzerland and Chem Service, West Chester, PA, respectively. Arsenobetaine (AsB) (BCR-626) was obtained from the Institute for Reference Materials and Measurements, Geel, Belgium.

Further arsenic species were derived from earlier studies at which they were either isolated from marine organisms (DMArsenoSugar-AminoSulfonate, -Glycol, -Phosphate, -Carboxyl, -Carbamate and -Methoxy) [ref. I, II] or synthesized (TMArsenoSugar-Glycol, -Phosphate, -Sulfonate, -Sulfate, -Methoxy [ref. III], DMArsenoSugar-Adenine [ref. IV], trimethylarsoniopropionate (AsB-2) and trimethylarsoniobutyrate (AsB-3) [ref. V], arsenocholine (AsC), tetramethylarsonium ion (TMA) and trimethylarsine oxide (TMAO) [ref. VI]).

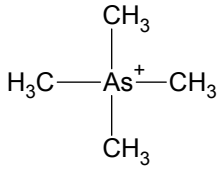
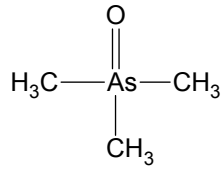
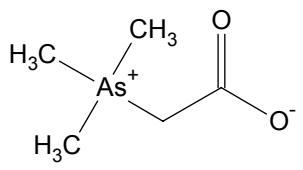
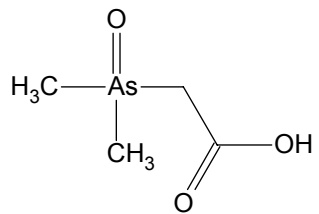
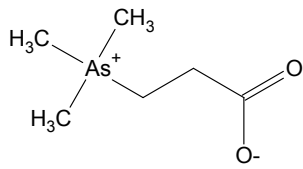
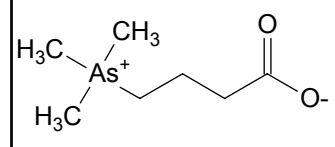
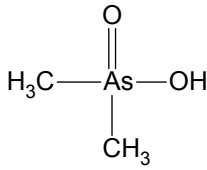
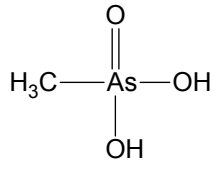
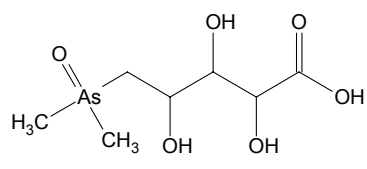
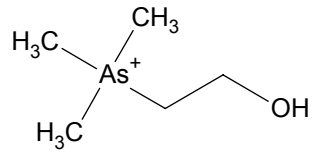
DMArsenoSugarHydroxy and TMArsenoSugarHydroxy were prepared by acidic hydrolysis of DMArsenoSugarMethoxy and TMArsenoSugarPhosphate, respectively. [ref. VII]

An aqueous extract of kelp powder was used as source of DMArsenoSugarSulfonate, DMArsenoSugarSulfate, DMThioAsSugarGlycol, DMThioAsSugarPhosphate, DMThioAsSugarSulfonate and DMThioAsSugarSulfate. [ref. VIII]

DMArsenoSugarMannitol and dimethylarsinoylacetic acid have been identified in extracts of a granulated kelp sample. [ref. IX]

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Tetramethylarsonium ion (TMA)		Trimethylarsine oxide (TMAO)	
Arsenobetaine (AsB)		Dimethylarsinoyl-acetic acid (DMAA)	
Trimethylarsonio-propionate (AsB-2)		Trimethylarsonio-butyrate (AsB-3)	
Dimethylarsinic acid (DMA)		Monomethylarsonic acid (MMA)	
5-Dimethylarsinoyl-2,3,4-trihydroxypentanoic acid (DMArsinoyl-Trihydroxypentanoic acid)		Arsenocholine (AsC)	

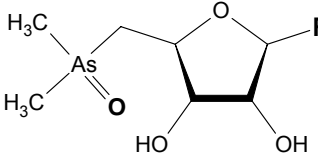
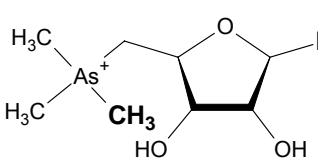
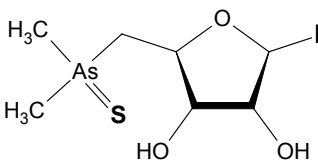
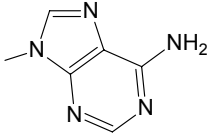
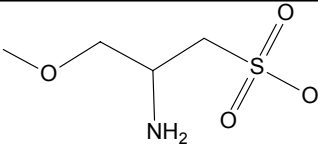
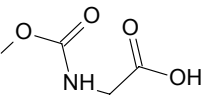
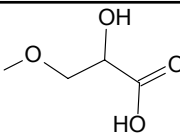
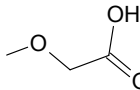
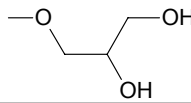

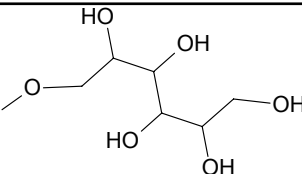
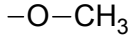
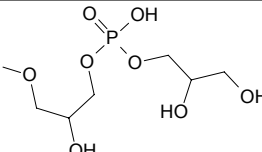
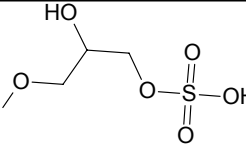
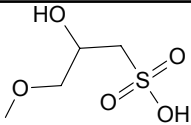
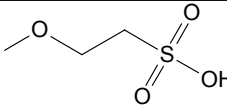
R	DMASugar-	TMASugar-	DMThioAsSugar-
			
Adenine			
Amino-Sulfonate			
Carbamate			
Carboxyl			
Carboxyl-2			
Glycol			
Hydroxy			
Mannitol			
Methoxy			
Phosphate			
Sulfate			
Sulfonate			
Sulfonate-2	(proposed structure in analogy to Carboxyl-2)		

Table A Selected reaction monitoring (SRM) parameters (precursor ion → product ion (collision energy [eV])) for 32 organoarsenic species and glycine betaine.

Species	SRM 1	SRM 2
TMA	135 → 120 (22)	135 → 105 (22)
AsB	179 → 120 (21)	179 → 105 (30)
AsB-2	193 → 120 (28)	193 → 105 (28)
AsB-3	207 → 87 (20)	207 → 121 (20)
AsC	165 → 121 (23)	165 → 105 (23)
TMAO	137 → 107 (26)	137 → 122 (20)
TMA _s SugarGlycol	327 → 193 (34)	327 → 120 (34)
TMA _s SugarHydroxy	253 → 193 (25)	253 → 163 (25)
TMA _s SugarMethoxy	267 → 75 (26)	267 → 120 (38)
TMA _s SugarPhosphate	481 → 327 (30)	481 → 389 (30)
TMA _s SugarSulfonate	391 → 235 (26)	391 → 293 (26)
TMA _s SugarSulfate	407 → 327 (20)	407 → 193 (40)
DMA	139 → 91 (24)	139 → 109 (24)
DMAA	181 → 119 (15)	181 → 139 (15)
DMA _s SugarAdenine	372 → 237 (20)	372 → 178 (28)
DMA _s SugarAminoSulfonate	392 → 97 (30)	392 → 295 (25)
DMA _s SugarCarbamate	356 → 237 (15)	356 → 97 (27)
DMA _s SugarCarboxyl	343 → 97 (28)	343 → 237 (16)
DMA _s SugarGlycol	329 → 97 (25)	329 → 237 (15)
DMA _s SugarHydroxy	255 → 97 (24)	255 → 195 (24)
DMA _s SugarMannitol	419 → 97 (38)	419 → 237 (20)
DMA _s SugarMethoxy	269 → 97 (21)	269 → 105 (32)
DMA _s SugarPhosphate	483 → 97 (40)	483 → 237 (25)
DMA _s SugarSulfonate	393 → 97 (30)	393 → 237 (20)
DMA _s SugarSulfate	409 → 329 (15)	409 → 97 (35)
DMThioAsSugarGlycol	345 → 97 (20)	345 → 253 (10)
DMThioAsSugarPhosphate	499 → 253 (15)	499 → 97 (27)
DMThioAsSugarSulfonate	409 → 97 (25)	409 → 253 (15)
DMThioAsSugarSulfate	425 → 97 (30)	425 → 253 (15)
MMA	141 → 91 (30)	-
DMA _s SugarCarboxyl-2	313 → 97 (20)	313 → 237 (15)
DMA _s rsinoylTrihydroxy-pentanoic acid	271 → 253 (18)	271 → 195 (25)
Glycine-betaine	118 → 58 (30)	118 → 59 (20)

Table B Quantification of arsenic species in extracts of 10 tissue fractions from a giant clam *Tridacna maxima* using anion exchange HPLC-ICPMS (method A). All contents are reported as $\mu\text{g As/g}$ dry weight ($n=1$). Peak assignment is based on HPLC-ES-SRM data and matching retention times with calibration standards in HPLC-ICPMS.

RT ^a	Peaks	TM ₁₋₁	TM ₁₋₂	TM ₁₋₃	TM ₁₋₄	TM ₁₋₅	TM ₁₋₆	TM ₁₋₇	TM ₁₋₈	TM ₁₋₉	TM ₁₋₁₀
2.3	AsB	10.0 ± 0.2	18.4 ± 0.3	11.2 ± 0.2	26.6 ± 0.4	25.6 ± 0.4	24.4 ± 0.4	20.0 ± 0.3	19.8 ± 0.3	14.6 ± 0.2	7.6 ± 0.1
2.4	DMGlycol	14.5 ± 0.6	- ^e	9.5 ± 0.5	-	-	-	-	-	-	-
2.8	TMA	3.2 ± 0.2	0.1 ± 0.1	3.0 ± 0.1	0.15 ± 0.08	0.1 ± 0.1	0.9 ± 0.1	0.47 ± 0.08	0.1 ± 0.1	0.06 ± 0.05	0.06 ± 0.05
3.1	DMA	1.7 ± 0.2	-	1.6 ± 0.2	-	-	-	-	-	-	-
3.5	DMCarboxyl ^b	46.4 ± 0.8	0.6 ± 0.1	45.1 ± 0.7	0.24 ± 0.08	0.8 ± 0.1	10.4 ± 0.2	1.82 ± 0.08	2.8 ± 0.1	0.22 ± 0.05	0.50 ± 0.04
4.2	DMCarbamate	24.6 ± 0.4	0.3 ± 0.1	24.9 ± 0.4	0.11 ± 0.08	0.5 ± 0.1	5.6 ± 0.2	0.60 ± 0.08	1.42 ± 0.09	-	0.31 ± 0.04
5.2	DMSulfonate	9.3 ± 0.2	0.2 ± 0.1	8.2 ± 0.2	-	0.2 ± 0.1	2.0 ± 0.1	0.24 ± 0.09	0.6 ± 0.1	-	0.14 ± 0.05
5.9	Unknown1 ^c	5.9 ± 0.2	0.2 ± 0.1	4.8 ± 0.2	-	0.2 ± 0.1	1.1 ± 0.1	0.21 ± 0.09	0.4 ± 0.1	0.07 ± 0.05	0.11 ± 0.05
8.8	DMSulfate	84 ± 1	0.7 ± 0.1	82 ± 1	0.22 ± 0.08	1.6 ± 0.1	17.2 ± 0.3	1.69 ± 0.08	4.5 ± 0.1	0.10 ± 0.05	0.91 ± 0.04
10.6	Arsenate	8.9 ± 0.5	-	6.3 ± 0.4	-	0.2 ± 0.1	1.7 ± 0.1	0.35 ± 0.07	0.41 ± 0.08	-	0.10 ± 0.04
17.2	DMAdenine	3.6 ± 0.6	-	1.3 ± 0.5	-	0.5 ± 0.4	1.1 ± 0.4	0.4 ± 0.3	0.3 ± 0.3	-	0.2 ± 0.2
30.9	Unknown2	19.8 ± 0.3	-	17.9 ± 0.3	0.11 ± 0.08	0.3 ± 0.1	1.8 ± 0.1	0.39 ± 0.09	0.4 ± 0.1	-	-
Recovery [%] ^d		94 ± 2	92 ± 8	91 ± 1	90 ± 4	87 ± 5	89 ± 3	84 ± 4	91 ± 4	93 ± 4	87 ± 6

^a Retention times [min]; additional peaks at 2.1 min for TM₁₋₁ (3.7 ± 0.2) and for TM₁₋₃ (3.3 ± 0.2) and at 12.0 min for TM₁₋₃ (0.7 ± 0.2)

^b DMPosphate, DMCarboxyl-2

^c Identified as DMA₅SugarSulfonate-2

^d Sum of quantified species in % of total arsenic content in the extract

^e not detected

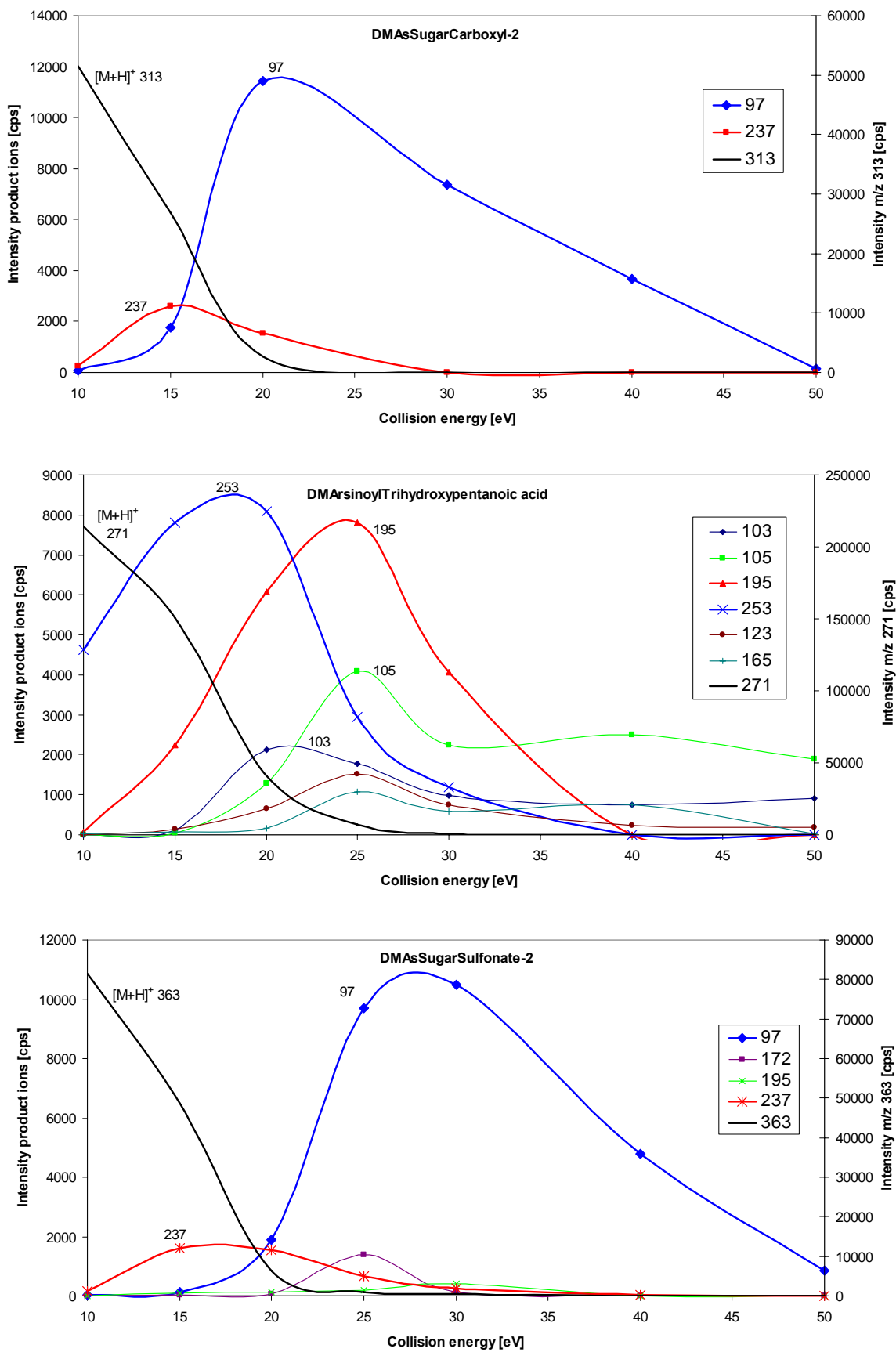


Figure A Collision induced dissociation (CID) breakdown curves for 3 arsenic species detected in *T. maxima* kidney extracts.

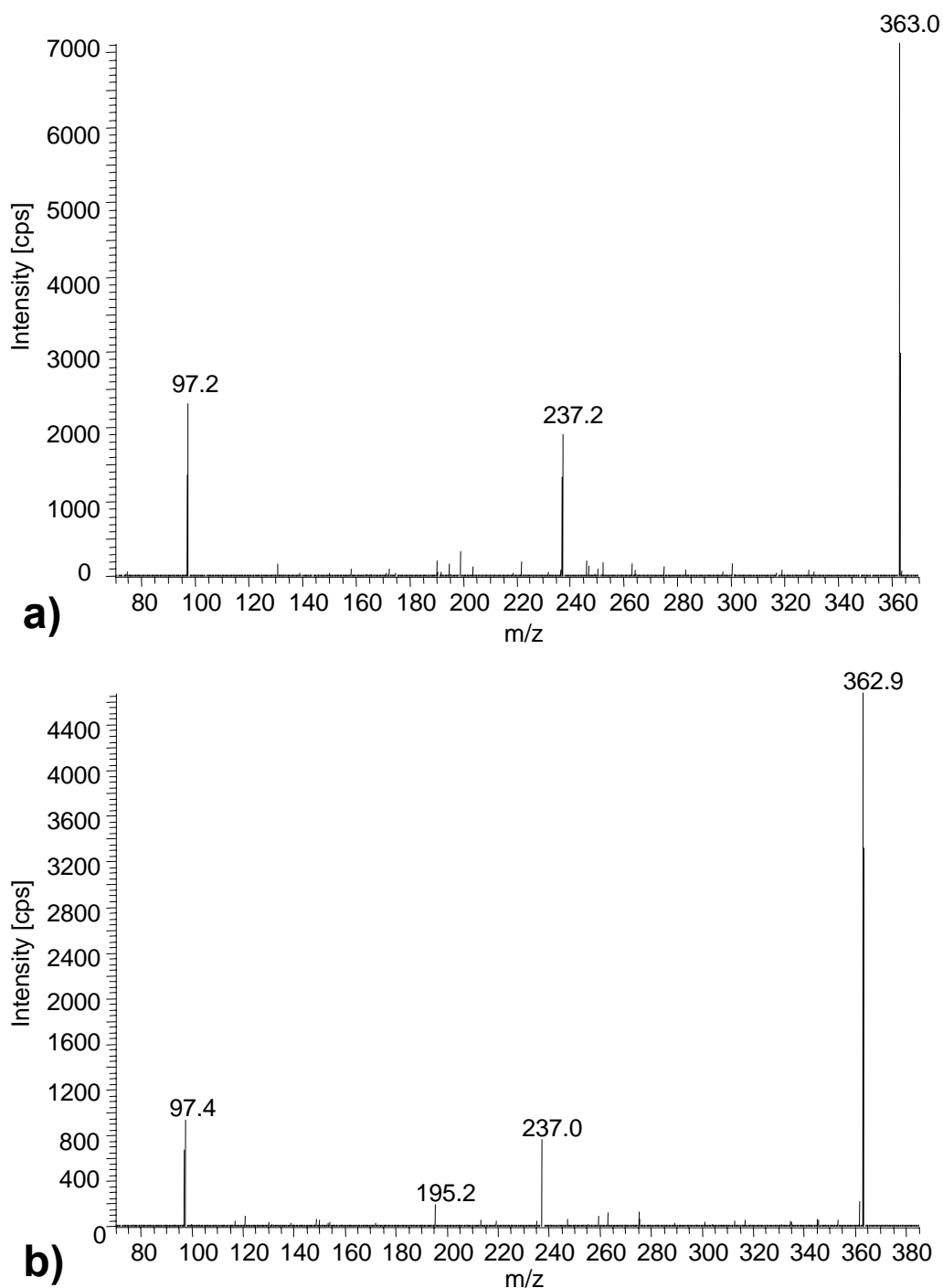


Figure B Product ion spectra of m/z 363 at 20 eV collision energy (*Tridacna* kidney extract) (a) and m/z 380 at 25 eV collision energy (same fraction of *Tridacna* kidney extract as in Fig. 4) (b) obtained from anion exchange HPLC-ES-MS/MS using 20 mM ammonium bicarbonate mobile phase at pH 10 with 10% methanol.

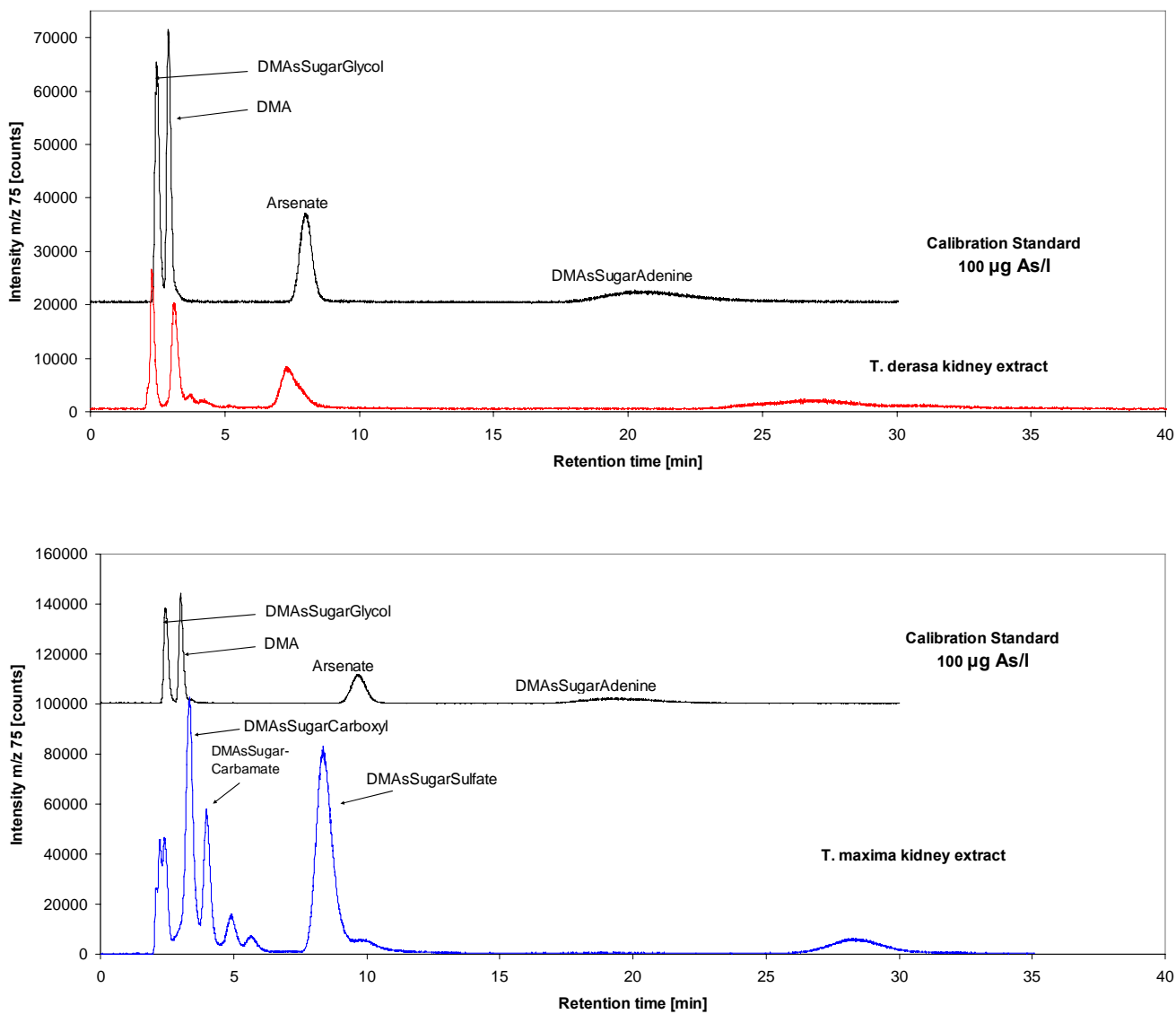


Figure C Anion exchange HPLC-ICPMS chromatograms (PRP-X100 column, 20 mM NH_4HCO_3 , 3% methanol, pH 10; see experimental section for details) of an extract from *T. derasa* kidney (TD_1 , upper graph) and an extract from *T. maxima* kidney (TM_{1-1} , lower graph). The samples were measured on different days, but the same calibration standard was analyzed on both days to indicate small changes in the retention behavior.