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Contribution to toxicity assessment of *Dinophysis acuminata* (Dinophyceae)

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Abstract: Blooms of *Dinophysis* in French coastal waters are implicated in most bans on marketing commercial bivalves. However, the relation between *Dinophysis* cell density and shellfish toxicity is not always consistent. Discrepancies may be due to the simple fact that it is nearly impossible to compare an integral over a few days (shellfish toxin content) and water samples. Furthermore, it seems that cells may have a variable specific toxicity. This work focuses on the variability in cell toxicity taking into account recent findings and using liquid chromatography coupled to mass spectrometry with an ion trap and electrospray interface. Esterified analogues of okadaic acid (DTX-4 and diol-esters) have been identified in cultures of *Prorocentrum lima*, another okadaic acid producer. These analogues are inactive on some protein phosphatases, contrary to okadaic acid, and seem to protect the cell from harmful effects by the toxin and to be enzymatically hydrolyzed during cell lysis. In order to document specific toxicity and to validate the presence of these analogues, *D. acuminata* concentrates were subjected to two separate heating and freeze/thaw procedures, respectively inhibiting or promoting hydrolysis. This paper reports on the high variability of *D. acuminata* specific toxicity and the presence of esters found in half of the samples only.

Keywords: *Dinophysis acuminata* - dinophysistoxins - okadaic acid - liquid chromatography-electrospray ionization mass spectrometry

Introduction

Several species of *Dinophysis* produce toxins known as dinophysistoxins (DTXs,) potentially causing gastroenteritis-type food poisoning once accumulated by filter-feeding molluscs. In France, *Dinophysis* has been the major cause of economic losses suffered by shellfish farmers. The relation between *Dinophysis* spp and mussel toxicity outbreaks is relatively well established along the Atlantic and English Channel coastlines: e.g. in Bay of Seine in 1995 (Marcaillou-Le Baut et al., 2001). However, some cases of high shellfish toxicity with low or zero cell counts in the water, or vice versa have been observed in monitoring surveys. Jackson and Silke (1995) have reported on the difficulty of relating toxic cell count with shellfish toxicity, hence preventing the use of cell counts as a prediction tool for monitoring purposes. Variability in cell toxin content is one of the factors explaining this lack of relation. In Canada, Marr et al. (1992) found that phytoplankton samples in which *Dinophysis* spp predominate do not always contain detectable DTXs. Several publications have since then shown such variability. For *D. acuminata*, the reported toxicity ranges from non-detectable (Lee et al., 1989; Hoshiai et al., 1997) to around 50 pg cell⁻¹ okadaic acid (Andersen et al., 1996; Sato et al., 1996).

Concurrently with these investigations on *Dinophysis*, Hu et al. (1995a,b) have detected okadaic acid-related compounds in *Prorocentrum lima*, another benthic dinoflagellate known to produce DTXs. The terminal acid function of okadaic acid is esterified with chains of various length (diol-esters and DTX4), which renders them inactive on some protein phosphatases (PP2A and PP1, Bialojan & Takaï, 1988). Okadaic acid being a powerful inhibitor of these proteins, the authors have hypothesized that these inactive esters may represent the primary synthesized form of DTXs. Quilliam et al. have actually shown in 1996 that a majority of DTXs in a *P. lima* culture are found in DTX4 form (okadaic acid -sulfated ester), and concluded that a more comprehensive analysis of toxins would be required for a more realistic assessment of phytoplankton toxicity.

This paper reports on the levels and variations of *D. acuminata* toxicity taking into account these recent findings.

Material and methods

Sample collection and treatment

The protocol described by Quilliam (1996) was adapted to natural samples in order to screen for the usual toxins and their esterified forms, if any, via indirect quantification, with and without promoting the enzymatic hydrolysis of esterified forms

Seawater samples were collected during four sampling cruises during *Dinophysis* bloom outbreaks, at a single sampling station located 8 km off the Southern Brittany coast. Sampling depth was chosen according to particles populations measured by an in situ particle size analyzer (Gentien et al., 1995). Water samples (approximately 400 L) were pumped with a peristaltic pump, then prefiltered over a 100- μm screen, with final concentration over a 20- μm sieve (final volume: approx. 300 mL). The second sieve was kept permanently submerged to prevent cell damage. The concentrate was kept under constant low agitation to ensure homogeneity, and divided into three subsamples. One Lugol-fixed subsample was set aside for subsequent cell count. The second subsample (named heated fraction or A) was immersed for 10 min at 100 °C in a water-bath in order to block the enzymatic activity, then cooled down to ambient temperature and immersed in liquid nitrogen. The third subsample (named unheated fraction or B) was immediately immersed in liquid nitrogen. All samples were preserved at -80 °C in the laboratory until analyzed.

Dinophysis cell density

Dinophysis cell density was estimated with a reverse microscope in a 10-mL subsample of the lugol-fixed suspension (Utermöhl, 1958). Except for the occasional presence of a couple of *D. rotundata* cells, the cells observed were related to the *acuminata* group. When needed, the suspension was diluted so as not to exceed few hundred cells in the entire chamber. The standard error on three replicates of three independent samples was estimated to 15%.

Toxin analysis

Prior to extraction, all samples were allowed to thaw for 24 hours at room temperature, before sonication and centrifugation for 20 min at 3000 rpm. The extraction protocol, inspired from Susuki et al (1997), was adapted for use with an automatic sample processor (ASPEC Xli, Gilson) to ensure optimum reproducibility (Mondeguer et al., 2004). After checking for the absence of toxin on the bottom, extraction was carried out on the supernatant, according to the following sequences executed by the automat on SPE cartridge (C18, 500 mg, 3 mL):

- 1) drying under nitrogen atmosphere and compression of silica for 4 min,
- 2) conditioning of cartridge with 20 mL pure methanol followed by 20 mL water at 8 mL min⁻¹,
- 3) deposit of 5 mL supernatant at 2.50 mL min⁻¹,
- 4) wash with 5 mL water at 3 mL min⁻¹,
- 5) elution of toxins with 5 mL pure methanol at 3 mL min⁻¹ and drying with 6 mL pulsed air.

The 5 mL methanol fraction was recovered, evaporated and diluted in 100 μ L pure methanol. A 5 μ L fraction of the final extract was injected into the HPLC/MS. The conditions used in the above protocol (sample dilution, conditioning, wash, analyte elution over cartridge, etc.) were initially designed for *Prorocentrum lima* cultures, and selected for criteria of yield efficiency, reproducibility and signal intensity.

Toxin detection and quantification were carried out with HPLC coupled with ion-trap mass spectrometry (Finnigan-LCQ ion trap), under the following conditions: Column and precolumn Kromasil C18 (250 mm x 2.0 mm I.D. 5 μ m); temperature: 40 °C; Isocratic mobile phase: acetonitrile/water + 0.1% TFA, (75: 25, v/v); flow: 0.2 mL min⁻¹.

Data acquisition was conducted in positive mode, with alternating full MS and full MS2, providing both for universal (full) detection and for highly specific and selective detection (full MS2) in one single injection. The analysis quantifies the daughter ions.

The accuracy and reliability of this automated extraction procedure coupled with LC/MS detection were validated against two reference systems (Algrandi et al., 1992, AFNOR, 1998). According to these standards, the range of linearity was defined as 0.05 to 11.5 ng OA or DTX1 (n = 3, p = 9). The precision of the method as adapted to solid phases and optimized for extraction parameters was checked against a certified reference specimen of digestive gland contaminated with 11.003 μ g. OA g⁻¹ (Quilliam, 1995). Validation tests revealed that the method was capable of detecting 63 pg and of measuring 186 pg OA-equivalent, with a 1% error risk and an experimental standard deviation of 66 pg (Mondeguer et al., 2004).

Results

Table 1 summarizes the characteristics of the sampling points and the results obtained. In spite of the patchiness of *D. acuminata* populations, the samples generally contained significant amounts of toxic cells: four samples only had densities below 100 cells L⁻¹. The hypothesis assuming the *Dinophysis* genus to be the okadaic acid source relies on the fact that no toxins are detected – or as traces only – in samples with very low *Dinophysis* spp. density, and that on other occasions and in spite of the presence of *Dinophysis*, samples were found to be toxin-free (data not shown). Table 1 presents okadaic acid contents alone since no substituted forms (DTX1 or 2) were detected in the concentrates.

Sample	Date	Hour	Depth (m)	Cell Density (cells mL ⁻¹)	OA Content. (pg cell ⁻¹)		Difference B-A (OA pg cell ⁻¹)
					A	B	
1	04 05 99	1530	6	113	5	9	4
2	5 05 99	1400	5	180	24	24	0
3	31 05 99	1830	3	46	6	48	42
4	01 06 99	1000	4	62	17	42	25
5	01 06 99	0800	2	27	0	68	68
6	09 05 00	1700	5.5	360	23	12	-11
7	10 05 00	0900	4.5	835	22	32	10
8	10 05 00	0900	2	130	32	29	-3
9	10 05 00	1100	5	810	34	22	-12
10	10 05 00	1430	12	270	4	11	7
11	10 05 00	1630	2	515	0	33	33
12	10 05 00	1730	3	1200	58	72	14
13	11 05 00	1100	10	520	22	17	-5
14	11 05 00	1410	2	590	57	50	-7
15	11 05 00	1410	2	620	54	66	12
16	12 05 00	1030	15	264	44	43	-1
17	12 05 00	1330	9	1020	57	35	-22
18	12 05 00	1430	10	880	45	46	1
19	13 05 00	10h00	10	970	98	78	-20
20	13 05 00	1130	11	780	63	63	0
21	22 05 00	1530	7	340	27	15	-12
22	22 05 00	1730	4.5	740	51	39	-12
23	23 05 00	0830	3.5	660	28	40	12
24	23 05 00	1030	2.5	575	2	3	1
25	23 05 00	1200	16	420	31	16	-15
26	23 05 00	1700	8	470	61	63	2
27	24 05 00	1100	10	400	62	63	1
28	24 05 00	1300	6	97	158	140	-18
29	24 05 00	1600	7	510	14	9	-5
30	24 05 00	1730	12	154	102	170	68
31	25 05 00	1430	3.5	750	12	14	2
32	26 05 00	1000	8	158	0	1	1

Table 1. Characteristics and okadaic acid (OA) content of phytoplankton samples after heated (A) and unheated (B) treatments. Heated treatment gives free okadaic acid and unheated treatment measures free okadaic acid + sulfated esters.

The fractions subjected to slow thawing without prior heating revealed highly variable okadaic acid contents (Table 1): i.e. values ranging from a few pg up to 160 pg cell⁻¹. The okadaic acid content variability of heated fractions was found to be roughly similar to unheated fractions (Figure 1).

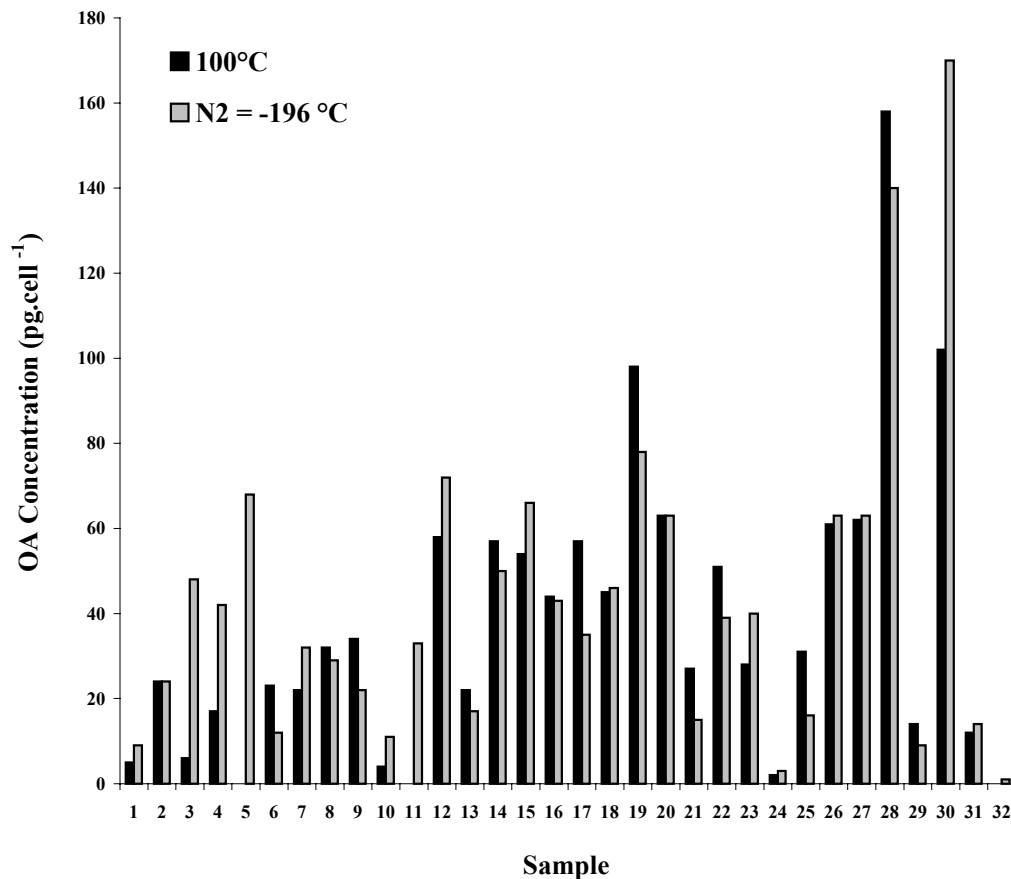


Figure 1. Influence of two different treatments on okadaic acid content of *D. acuminata* concentrates divided in two sub-samples: heated (black) and unheated (grey) fractions.

According to the initial tested assumption, the heated fraction should contain esters, and therefore a lower content in free okadaic acid.

Yet, not only was their okadaic acid content not systematically lower, but it was sometimes much higher, as expressed by differences between the values of unheated and heated subsamples (B-A), occurring in virtually equal numbers of cases (Figure 2): i.e. either positive, or negative (some were non-significant).

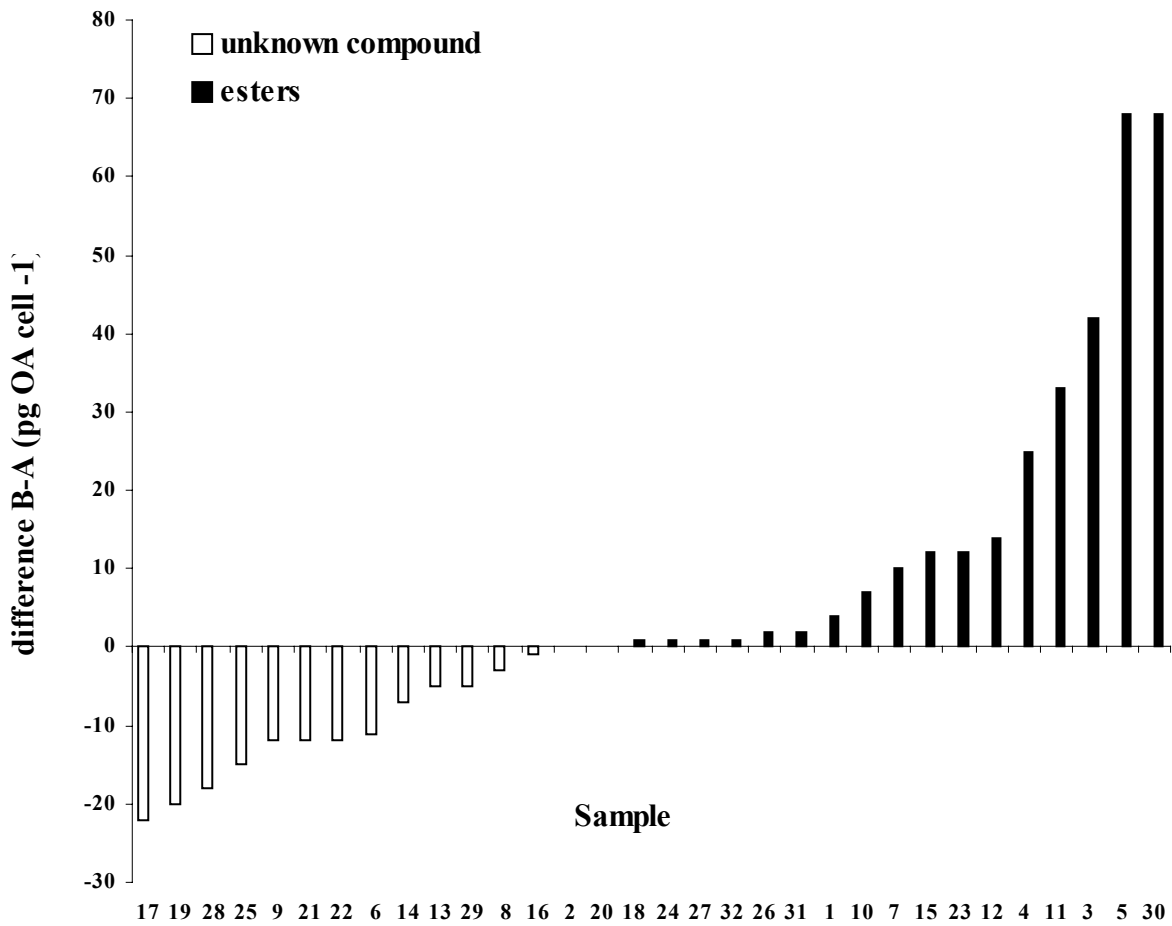


Figure 2. Classification of differences in okadaic acid content between the unheated and heated fractions and inversely: A-B: OA release from unknown compounds (□); B-A: presence of esters (■).

No relation was found with the time or depth of sampling, as illustrated by results from two consecutive sampling days. As an example, on 23 and 24 June 2000, four samples were collected within a few hours: the minimum and maximum values were 3 – 63 pg cell⁻¹ and 9-170 pg cell⁻¹, respectively for unheated and heated procedures.

Discussion

The toxicity of *D. acuminata* in this series of samples shows significant variations over a period of a few days and at a single site, while previously published values reported on samples collected over longer time and space intervals. These results would thus reflect the mean toxicity of a heterogeneous cell population at a given moment. Consequently, when attempting to relate cell density and shellfish toxicity, care should be taken to estimate intrapopulation

specific toxicity. The okadaic acid content measured in unheated fractions are relatively higher than those previously reported, though not exceeding a few tens of pg per cell (Andersen et al., 1996), a result that could be explained by the systematic use of the slow thawing procedure in this study, which promotes the hydrolysis of esterified forms into free okadaic acid.

The presence of esters is confirmed in one-third of the sample. Their level varies from 0 to 100% (virtually no or little free OA in the heated fraction), as described by other authors. Using the same indirect detection procedure, Moroño et al. (2003) also found variable ester concentrations, but none with values exceeding 11.2 pg cell⁻¹. In 2004, Vale showed that it was possible with this technique to recover nearly 90% of the esters contained in a *Dinophysis* concentrate. Finally, Suzuki et al. (2004) showed the presence of several okadaic acid derivatives in *D. acuta*, the most predominant being OA-D8, previously identified by Quilliam (1996).

An examination of differences in toxicity content between unheated and heated fractions however suggests the presence of (an)other form(s) that would be converted into okadaic acid under the action of heat, yet would remain stable under thawing, and could in certain cases be present in amounts greater than esters. This observation refers to the findings of Sato et al. (1996), who reported very high concentrations (230 pg equiv OA cell⁻¹) measured with ELISA tests in isolated *D. acuminata* cells, while a chemical analysis of the same extract yielded a much lower value of 20 pg OA+DTX1 cell⁻¹. These authors further mention a compound different from water-soluble esters (DTX4-like), considered likely to crosslink with anti-OA antibodies and to re-form okadaic acid.

Neither dinophysistoxin-1 nor 2 were detected in this series of samples, an absence previously reported in *D. acuminata* concentrates from the Bay of Seine along the English Channel coast (Marcaillou Le Baut et al., 2001). These results are in accordance with most other published reports (Fernandez et al., 2000) suggesting that *D. acuminata* should be classified among okadaic acid producers. However, some caution is needed, since Sato et al (1996) detected dinophysistoxin-1 in this species at 65 pg cell⁻¹ level.

In conclusion, the analyses of this series reveal that the toxin profile of *D. acuminata* may be more complex than considered previously, and confirms its high variability. Consequently, specific toxicity expressed as a given amount of DTXs (OA and/or DTX1-2) per cell is a highly inadequate measure, particularly for any definition of a shellfish contamination model.

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