

# Accumulation of red tide toxins in larger size fractions of zooplankton assemblages from Massachusetts Bay, USA

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**ABSTRACT:** Phytoplankton toxins undergo trophic transport and accumulation in marine food webs, causing vectorial intoxication of upper-level consumers such as fishes, seabirds, and marine mammals. An entry point for phytoplankton toxins into these pelagic trophic pathways is frequently the herbivorous zooplankton. During the 1995 spring–summer red tide season in Massachusetts Bay, we examined accumulation of paralytic shellfish poisoning (PSP) toxins from the dinoflagellate *Alexandrium* spp. in various plankton size fractions (20–64, 64–100, 100–200, 200–500, and >500 µm), and identified the relative composition of the zooplankton in these size fractions. Toxin levels were estimated by both high-performance liquid chromatography (HPLC) and a receptor-binding assay, the latter based on sample toxic potency. Although no PSP toxicity was detected in nearshore shellfish by routine monitoring programs using the mouse bioassay, positive responses were detected in zooplankton size fractions with the more sensitive HPLC and the receptor assay methods. The toxin signal was disproportionately concentrated in the larger zooplankton size fraction, frequently dominated by large copepods such as *Calanus finmarchicus* and *Centropages typicus*, which comprised only a small portion of total zooplankton abundance in quantitative samples obtained with 100 µm mesh nets. By comparison, signal levels were low or undetectable in the smaller size fraction, which contained the overwhelmingly most-abundant zooplankters such as protists, copepod nauplii and copepodites and adults of small copepods such as *Oithona similis*, *Paracalanus parvus*, and *Pseudocalanus* spp. The larger toxin-accumulating copepods could provide a direct trophic linkage for vectorial intoxication of baleen whales that are known to feed upon such copepods.

**KEY WORDS:** Red tide · Harmful algal blooms · Zooplankton · Copepods · Algal toxins · Vectorial intoxication · Massachusetts Bay · Gulf of Maine

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## INTRODUCTION

Marine phytoplankton toxins can cause finfish mortality and shellfish toxicity, as well as poisoning or death of humans, marine mammals and seabirds that

ingest contaminated fish or shellfish (Anderson & White 1992, Baden & Trainer 1993, Turner & Tester 1997, Turner et al. 1998b). Accumulation and transport of these toxins through feeding interactions in pelagic food webs can also cause vectorial intoxication of consumers at higher trophic levels that are not direct consumers of toxic phytoplankton. The pioneer-

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ing work of White (1977, 1979, 1980, 1981) on accumulation of paralytic shellfish poisoning (PSP) toxins in various compartments of zooplankton communities indicated that food-web transport of these toxins could cause fish kills in the Bay of Fundy. The dinoflagellates White classified as *Gonyaulax excavata* are now viewed as members of the genus *Alexandrium* (Moestrup & Larsen 1990). There have been few subsequent field studies on

vectorial transport of dinoflagellate toxins to upper-level consumers via zooplankton (Tester et al. 2000).

A major obstacle to studies of vectorial intoxication through zooplankton pathways has been the difficulty of measuring phytoplankton toxins in the ppb range found in field zooplankton populations. Recent advances in analytical techniques such as high-performance liquid chromatography (HPLC) (Oshima 1995), immunoassays and receptor-binding assays (Van Dolah et al. 1994, Cembella et al. 1995), capillary electrophoresis (Shea 1997), and LC-tandem mass spectrometry (Lefebvre et al. 1999) allow detection of phyco toxins or their activity at concentrations low enough for detection in field populations of zooplankton.

During the 1995 spring–summer bloom period of *Alexandrium* spp. in Massachusetts Bay, we measured PSP toxin levels in various size fractions of field zooplankton populations. Since 2 species of *Alexandrium* (*A. tamarense* and *A. fundyense*) can co-occur in the southern Gulf of Maine and Massachusetts Bay (Anderson 1997), and these are not readily distinguishable without considerable effort, we hereafter use only the genus name to refer to toxic *Alexandrium* spp. in our study area. PSP toxin levels were determined using HPLC and a saxitoxin receptor-binding assay (Doucette et al. 1997), employed here in a field study for the first time. Our goal was to measure PSP accumulation in zooplankton, which are the entry points to food webs that are thought to have caused fatal vectorial intoxication of marine mammals in these waters (Geraci et al. 1989).

## METHODS

**Study site.** The site chosen for our study, Massachusetts Bay, is an open embayment offshore of Boston, defined by Cape Ann at the northern boundary and Cape Cod at the southern boundary, with Stellwagen Bank lying at the outer edge of the Bay (Fig. 1). Stellwagen Bank is a shallow, highly productive offshore embankment of about 20 m depth that is known as a habitat for several species of endangered marine mam-

Table 1. Cruises, dates in 1995, and stations for collection of zooplankton size fractions for toxin analyses, and stations for which toxins were recorded with both HPLC and receptor-binding assay (RBA)

| Cruise | Date      | Size-fraction collection stations | Toxins with HPLC + RBA |
|--------|-----------|-----------------------------------|------------------------|
| 95MB1  | 26–27 Apr | 1, 6, 11, 15, 20, 23, 25, 27, 31  | 31                     |
| 95MB2  | 10 May    | 1, 6, 9, 11, 15, 19               | 1, 9, 11, 15, 19       |
| 95MB4  | 6 Jun     | 2                                 | 2                      |
| 95MB5  | 20 Jun    | 2, 7, 10, 11, 15                  | 10, 11, 15             |

als. It has recently been designated as a marine sanctuary. The areas just south of Boston Harbor, known as the 'South Shore', experience outbreaks of PSP shellfish toxicity every several years during the spring and early summer months. The last significant outbreak was recorded in 1993, although toxic cells are present in the bay annually.

The source of *Alexandrium* blooms in Massachusetts Bay has been suggested to be upstream in the Casco Bay/Kennebec River area of southern Maine (Franks & Anderson 1992, Anderson, 1997). *Alexandrium* populations are transported by the Western Maine Coastal Current, which runs from northeast to southwest, impacting the coastlines of southern Maine, New Hampshire and Massachusetts, with inputs into Massachusetts Bay entering near Cape Ann.

**Sampling.** Samples for zooplankton size fractionation were collected on 4 cruises during April, May, and June 1995 (Fig. 1, Table 1). Temperature and salinity data were collected using a Seabird SeaCat CTD profiler #19 and initially processed using Seabird's software. Final processing to produce selected contours of salinity, temperature, and cell concentration was completed using Matlab routines that included the Massachusetts Bay coastline. Satellite data of sea-surface temperature (SST) were also downloaded through NOAA's Coastwatch node and analyzed during the bloom season from April–June 1995.

Samples for *Alexandrium* abundance were collected with Niskin bottles at all stations, from the surface, and at 10 and 20 m depths. Two-liter samples were sieved onto 20 µm mesh Nitex screens, and the phytoplankton retained on the screens were backwashed and preserved with borate-buffered formalin (5% final concentration). Samples were stored in the dark, and refrigerated until analyses. Since *Alexandrium* cells were usually located primarily in the upper mixed layer (0 to 10 m) (Franks & Anderson 1992), the 20 m samples were archived and not analyzed.

Zooplankton were usually collected at 3 stations on each transect, generally at the most offshore, the most inshore, and an intermediate station (Fig. 1). At each station where zooplankton size fractions were obtained,

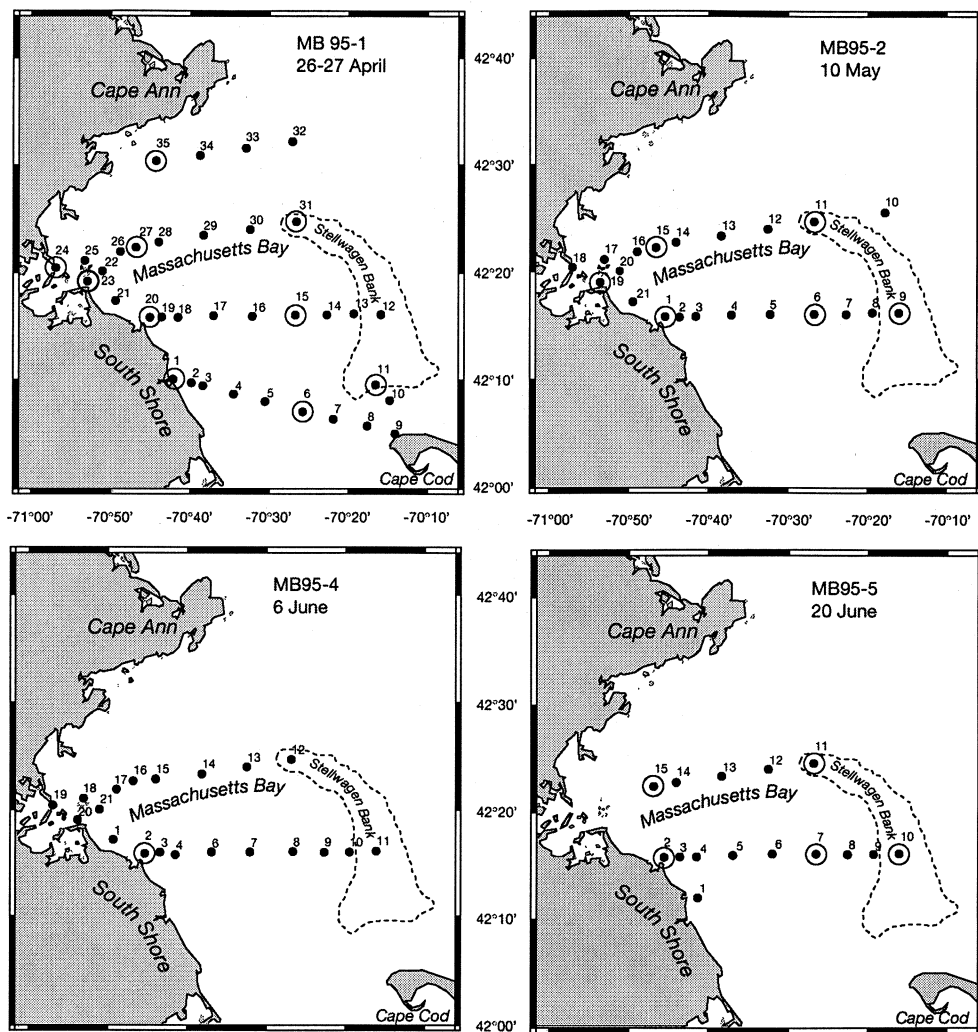


Fig. 1. Station locations for cruises in Massachusetts Bay during 1995 spring-summer bloom season of *Alexandrium* spp. Zooplankton/toxin stations are circled

we sampled for PSP toxins in, and taxonomic composition of, the microzooplankton (20 to 64 and 64 to 100  $\mu\text{m}$ ) and mesozooplankton (100 to 200, 200 to 500, > 500  $\mu\text{m}$ ) fractions.

For microzooplankton, surface water was collected by running water from a deck hose into a 20  $\mu\text{m}$  mesh net hanging over the side of the boat. Water was passed through this net for approx. 15 to 30 min at each station, obtaining total samples estimated to represent >100  $\text{m}^3$ .

For collection of mesozooplankton, 2 horizontal net tows were made with 0.5 m diam., 100  $\mu\text{m}$  mesh nets from just below the surface. This approach was adopted so that net tows would come from the same depth strata as the water for dinoflagellates and microzooplankton. First, a quantitative timed tow was made using a net equipped with a flowmeter (General Oceanics Model 2030), and all zooplankton were pre-

served immediately in 5 to 10% formalin:seawater solutions for quantitative taxonomic analyses. Care was taken to ensure that flowmeters were still turning upon retrieval, indicating that there was no net clogging. If clogging occurred, the tow was repeated for shorter time periods, until there was no clogging. Generally, the quantitative tows lasted <2 min. Immediately after the quantitative tow, a second non-quantitative tow was made with the same net for collection of mesozooplankton for toxin analyses. These tows were made without regard to flowmeter readings, and generally lasted 10 min, whether there was net clogging or not. The purpose of the qualitative tows was to collect sufficient zooplankton biomass for accurate detection of a PSP toxin signal, if present. The long towing time required to obtain sufficient zooplankton biomass usually resulted in net clogging, precluding quantitative measurements of animals  $\text{m}^{-3}$  or zooplank-

ton biomass  $m^{-3}$  for these tows. However, plankton from these tows were processed quantitatively in order to provide quantitative measurements of PSP toxin levels and percentage composition comprised by various zooplankton taxa in different size fractions (see below).

**Zooplankton size-fractionation for toxin and community composition analyses.** The catch from the qualitative tows was fractionated through sequential meshes of 500, 200, and 100  $\mu m$  to obtain zooplankton fractions of  $>500$ , 200 to 500, and 100 to 200  $\mu m$ . The slurry of plankton in these size fractions was then rinsed and consolidated by squirting with 20  $\mu m$  screened natural seawater (which would not have contained the 30 to 40  $\mu m$  diam. *Alexandrium* cells), scraped off meshes with spatulas, and packed into aliquots of 1 ml in plastic syringes with the tips cut off. The plungers of the syringes were then used to expel the 1 ml aliquot of wet plankton into preweighed plastic centrifuge tubes containing 1 ml of 0.1 N HCl. Tubes with plankton size fractions in acid were then frozen for toxin analyses ashore. The remainder of the slurry of each size fraction was preserved in 5 to 10% formalin-seawater solutions for microscopic analyses of the relative taxonomic composition of each size fraction.

The microplankton caught by the 20  $\mu m$  mesh net filtering water from the deck hose was poured onto a 20  $\mu m$  mesh sieve, and then backwashed with filtered seawater. This  $>20$   $\mu m$  suspension was split into 2 aliquots of approx. 100 to 500 ml each. One of these was filtered onto a 47  $\mu m$  diam. Whatman GF glass-fiber filter, which was then placed in a preweighed plastic centrifuge tube in 1 ml of 0.1 N HCl and frozen for toxin analyses. The other aliquot was screened through sequential sieves of 100, 64, and 20  $\mu m$ . The  $>100$   $\mu m$  fraction was discarded, and the remaining 64 to 100 and 20 to 64  $\mu m$  size fractions were backwashed into 15 ml centrifuge tubes with filtered seawater. Aliquots of 0.5 ml were removed from each fraction, added to 100 ml jars, and preserved in Utermöhl's solution (Guillard 1973), after bringing the total volume to approx. 100 ml with filtered seawater. These samples were later examined ashore for relative taxonomic composition of the microplankton in each fraction. The remaining 14.5 ml of each aliquot were filtered onto glass-fiber filters, which were placed in preweighed plastic centrifuge tubes in 1 ml of 0.1 N HCl for toxin analyses.

**Zooplankton abundance and community-composition analyses.** Mesozooplankton from quantitative tows were transferred to 70% ethanol solutions and split with a Folsom plankton splitter to obtain aliquots of approx. 300 to 500 animals each. Animals in these aliquots were counted and identified to the lowest possible taxonomic level with a dissecting microscope. Abundances were calculated as animals  $m^{-3}$ . Meso-

zooplankton in size fractions from qualitative tows were similarly split, counted, and identified, and abundances were presented as relative percentages of animals per aliquot. Microzooplankton in Utermöhl's preserved samples were examined in 1 ml aliquots in a Sedgwick-Rafter cell with a compound microscope, counted and identified, and presented as relative percentages of animals observed. Qualitative observations on presence and relative abundance of dinoflagellates were also made during examination of these aliquots.

***Alexandrium* abundance.** An immunofluorescence labeling protocol was used to identify and quantify *Alexandrium* cells with a genus-specific antibody probe (M8751-1; Adachi et al. 1993, Sako et al. 1993) known to react with cell-surface antigens of *Alexandrium* spp. from the Gulf of Maine region. This probe has been shown to label cultured *Alexandrium* cells with an acceptable level of consistency regardless of physiological condition.

To stain the cells, the preserved field samples for *Alexandrium* counts were further concentrated by settling and removal of the overlying seawater (final volume = 2 ml from a 2 l water sample). A 1 ml aliquot of the concentrate was transferred to a siliconized microcentrifuge tube, where the particulate material was pelleted by centrifugation (swinging bucket type;  $5000 \times g$  for 5 min). The supernatant was removed and 5% normal goat serum (NGS; Sigma #G-9023) in 0.02M PBS added as the blocking agent. After incubation for 30 min on a LabQuake Supplier shaker, the material was pelleted, the supernatant removed and 100  $\mu l$  of primary monoclonal antibody (M8751-1 in 5% NGS) were added and incubated with mixing for 30 min. After washing 3 times with 0.5% NGS in 0.02 M PBS by centrifugation to remove the excess primary antibody, 100  $\mu l$  of a secondary polyclonal antibody, goat anti-mouse IgG (heavy + light chain) conjugated to fluorescein isothiocyanate (FITC; Molecular Probes Inc. #F-2761; diluted 1:300 v/v with 5% NGS/PBS), were added and incubated for another 30 min. The stained samples were washed twice with 0.02M 5% NGS/PBS, resuspended in 100  $\mu l$  PBS, and 25  $\mu l$  80% glycerol/PBS were added. The sample was loaded into a Palmer-Maloney counting chamber and examined using a Zeiss epifluorescence microscope equipped with a FITC filter set (excitation = BP450–490; emission = LP 520). The full slide was counted for the presence of the bright green stain on the surface of the target *Alexandrium* cells, while red fluorescence, indicative of chlorophyll *a*, was also observed internally. Positive controls using cultured *Alexandrium* cells were stained with each batch of samples to ensure that the procedure worked properly.

**Toxin analyses.** Samples for toxin analyses were sonified, heated to 97°C for 5 min in a water bath for toxin hydrolysis, centrifuged, and the supernatants were passed through C-18 Sep-Pak cartridges. Aliquots were removed for both HPLC and receptor-binding assay analyses. The saxitoxin receptor assay is a rapid, high-throughput, competitive binding assay based on the interaction of saxitoxin and all other PSP derivatives with their biological target, the voltage-dependent sodium channel (Doucette et al. 1997). Values generated by this assay for extracts of *Alexandrium* cultures correspond closely to those obtained by HPLC, and the current work provides a similar comparison for estimating PSP toxin activity in size fractions of field material dominated by either *Alexandrium* spp. or various zooplankters. The assay protocol and calculations of sample values used herein follow those described by Doucette et al. (1997), except that a microplate scintillation counter (Wallac Inc., Gaithersburg, Maryland), rather than a conventional liquid scintillation counter, was employed to assess the assay's radioactive endpoint. Briefly, the receptor-binding assay is performed in a microplate filtration format and involves the incubation of a rat brain synaptosome preparation containing the receptors with <sup>3</sup>H-STX and unlabeled PSP toxins contained in a standard or sample. After removal of unbound toxin by washing, the remaining bound radioactivity (i.e., <sup>3</sup>H-STX), which is inversely and quantitatively related to the concentration of PSP-like activity in a sample, is determined by microplate scintillation counting. Toxin concentrations are expressed in terms of saxitoxin equivalents (STX equiv).

The receptor assay was used as an initial screen for the presence of PSP-like activity in the size-fractionated samples. Since extracts with a toxicity of ca 50 nM STX equiv or less were run undiluted on the assay, such samples having a highly acidic pH of <1 tended to yield false positive results and were not considered further. If an extract of any size fraction for a given station had a toxicity >50 nM STX equiv, then all size fractions from that station were also analyzed by HPLC. Although samples for toxin accumulation in zooplankton were collected at 21 stations, by the criteria given above, toxin levels were measured with both the HPLC and the receptor-binding assay in all size fractions from a total of 10 stations (Table 1).

HPLC analysis followed the methods described in Anderson et al. (1994), except that analysis for C toxins was not performed since the samples were previously hydrolyzed. Toxin concentrations of the individual derivatives were converted into STX equivalents (382 for dcGTX2 and 935 for dcGTX3) using the factors given by Oshima et al. (1992) for direct comparison to receptor assay results.

## RESULTS

### Shellfish toxicity and *Alexandrium* distributions

Shellfish toxicity was not recorded within Massachusetts Bay in 1995, but toxicity was recorded in shellfish from adjacent waters of southern Maine, New Hampshire and northern Massachusetts (D. Whittaker & J. Hurst pers. comm.).

*Alexandrium* was present in Massachusetts Bay during all cruises, but at most stations cells were either undetectable (detection limit = 1 cell l<sup>-1</sup>) or present in low concentrations (<25 cells l<sup>-1</sup>). Less than 10 cells l<sup>-1</sup> were recorded for Cruise 95MB1 (data not shown), but during Cruise 95MB2 on 10 May 1995, a population of *Alexandrium* was detected offshore near the northwest corner of Stellwagen Bank (Stn 11), with a maximum abundance of approx. 50 cells l<sup>-1</sup> (Fig. 2a). This low level population was near a very weak front associated with slightly warmer and slightly fresher water (8°C; 31.0 to 31.5 psu, Fig. 2b,c) than the ambient Gulf of Maine waters. The likely source of these cells was from the Casco Bay region, carried to the south in the Western Maine Coastal Current (Anderson 1997). A sea-surface temperature (SST) image on 9 May 1995 indicated a downwelling condition which compressed the warmer waters of the Western Maine Coastal Current against the coast during this time (Fig. 3 left panel). Rapid delivery of toxic cells from the north into Massachusetts Bay can occur under these conditions (Franks & Anderson 1992, Anderson 1997). By 28 May, SST imagery revealed that surface waters north of Cape Ann had generally warmed from 8.5–10.5°C to 12–13°C (note larger area covered by red and orange in Fig. 3, right panel, compared to area of light blue and yellow in Fig. 3, left panel), and that the coastal current had widened to the east, most likely keeping any *Alexandrium* populations away from nearshore (Fig. 3, right panel; Keafer & Anderson 1993). During this late May period, strong upwelling conditions were detected nearshore off southern Maine, visible as an orange patch of 10 to 11°C surface water immediately offshore from the Maine-New Hampshire state line, shown in Fig. 3, right panel. During subsequent cruises, cell concentrations were near detection limits, except for several offshore stations located near the southern flank of Stellwagen Bank where approx. 25 cells l<sup>-1</sup> were recorded during Cruise 95MB4 on 6 June 1995 (data not shown). This was probably another very weak pulse of cells originating from the north that passed through the offshore areas of Massachusetts Bay. The northeast winds needed to drive these cells into inner portions of the Bay did not occur.

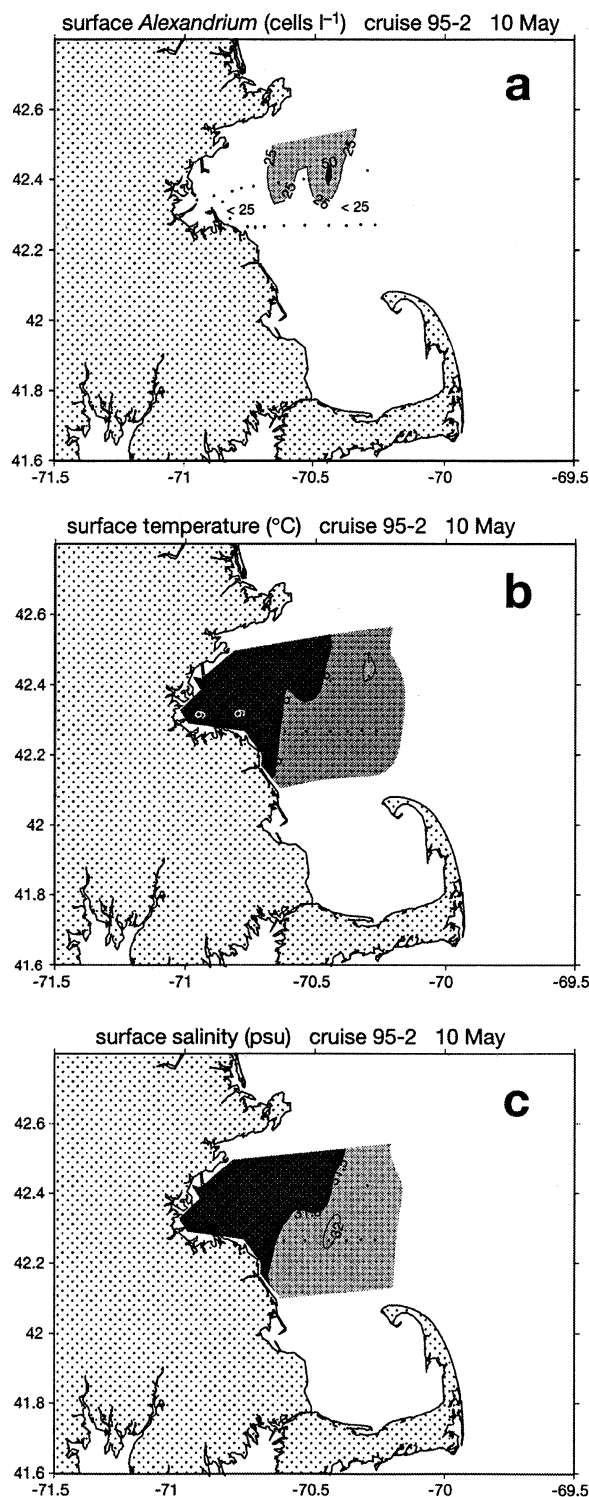


Fig. 2. Surface *Alexandrium* spp. cell concentrations and hydrographic conditions during Cruise 95MB-2. (a) Maximum abundance of toxic cells during bloom season was ca 50 cells  $l^{-1}$  near northwest corner of Stellwagen Bank. (b) Surface temperature was ca 8°C, and (c) salinity ca 31.25 psu

### Zooplankton and toxins

Size-fractionated material from 10 stations (5 fractions per station for a total of 50 samples) was assessed for PSP toxin levels by both the receptor-binding assay (Fig. 4a) and HPLC (Fig. 4b). A toxin signal was detected by at least 1 of the 2 methods in 23 of the 50 size-fractionated samples examined, with HPLC and the receptor-binding assay both reporting toxin on approx. 85% of the 23 positive samples.

Analyses of size fractions by the receptor-binding assay (Fig. 4a) and HPLC (Fig. 4b) revealed PSP toxin signals primarily in the smallest (20 to 64  $\mu m$ ) and the 2 largest (200 to 500 and  $>500 \mu m$ ) fractions. These represented *Alexandrium* cells and larger metazoan grazers, respectively. Toxin was in most cases absent from the 2 intermediate fractions (64 to 100 and 100 to 200  $\mu m$ ). Also, since the Fig. 4 ordinates are on a logarithmic scale, it should be pointed out that toxin levels measured by HPLC were considerably higher in the  $>500 \mu m$  size fraction (82 to 289 pmol STX equiv  $g^{-1}$  wet wt, mean = 202, for the 8 stations where toxins were detected) than in the 200 to 500  $\mu m$  size fraction (23 to 122 pmol STX equiv  $g^{-1}$  wet wt, mean = 71, for the 6 stations where toxins were detected). Receptor-binding assay toxin data were 8 to 210 pmol STX equiv  $g^{-1}$  wet wt, mean = 122, for the 9 stations where toxins were detected in the  $>500 \mu m$  size fraction, and 42 to 241 pmol STX equiv  $g^{-1}$  wet wt, mean = 142, for the 7 stations where toxins were detected in the 200 to 500  $\mu m$  size fraction. In terms of overall toxicity, Stns 9, 11 and 15 on Cruise 95MB2 and Stns 10, 11 and 15 on Cruise 95MB5 exhibited the highest levels measured. Cruise 95MB2 was notable in that *Alexandrium* cell concentrations were the highest of any determined during the 1995 sampling (approx. 50 cells  $l^{-1}$ ) and, likewise, PSP toxin levels in the 20 to 64  $\mu m$  size fraction containing these dinoflagellates were 1 to 2 orders of magnitude above any others measured in this study ( $4$  to  $40 \times 10^3$  pmol STX equiv  $g^{-1}$  wet wt). Toxin levels in the 20 to 64  $\mu m$  size fraction obtained during Cruise 95MB5 were considerably lower, consistent with the undetectable levels of *Alexandrium* cells at that time. However, on Cruise 95MB5, detectable toxin signals were present in the 200 to 500 and  $>500 \mu m$  size fractions.

### Zooplankton abundance and size-fraction community composition

Total zooplankton abundance from quantitative tows is presented in Table 2. Included are the 10 stations for which toxin levels were obtained with both HPLC and the receptor-binding assay (see Figs. 5 to 14).

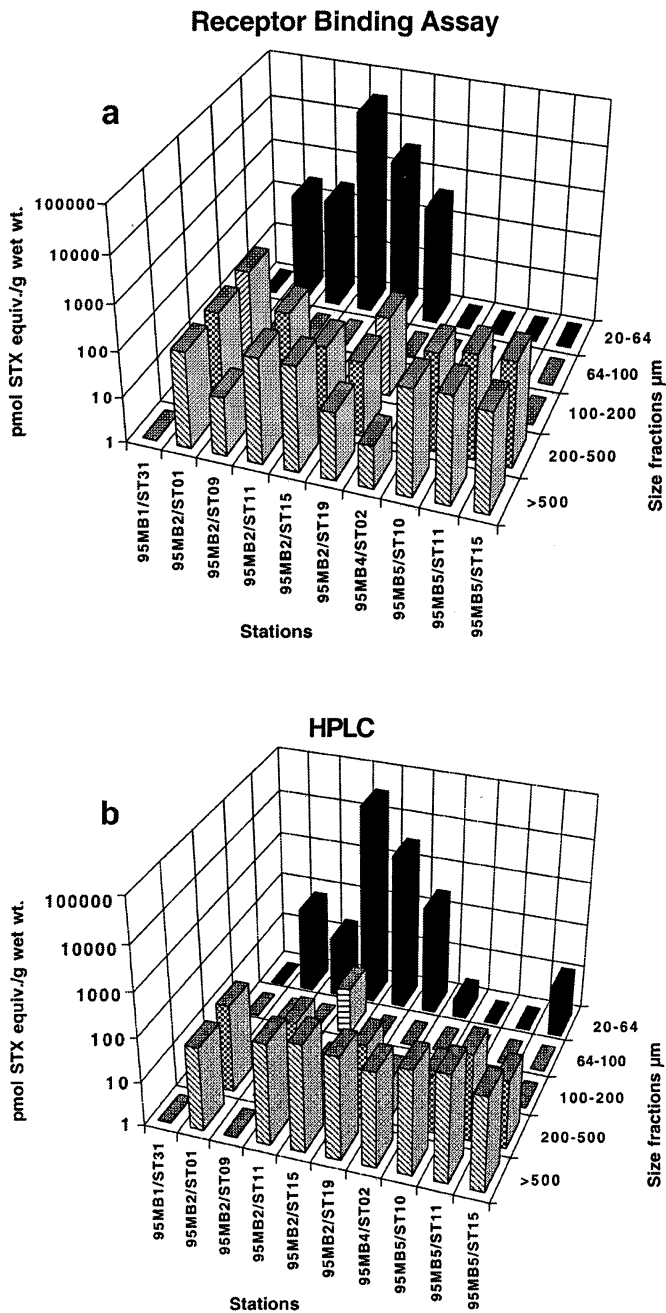


Fig. 4. Results of paralytic shellfish poisoning toxin determinations for size-fractionated zooplankton samples collected during 1995 Massachusetts Bay Cruises 1, 2, 4 and 5. Toxin measurements were performed by (a) receptor-binding assay and (b) HPLC. Toxin contained in sample extracts is expressed as pmol saxitoxin equivalents  $\text{g}^{-1}$  wet wt of biomass extracted to permit direct comparison between HPLC and receptor-assay values, as well as within and between cruises. Other than high levels of toxin found in smallest size fraction (20 to 64  $\mu\text{m}$ ), which contains the *Alexandrium* spp. cells, most toxin was contained in larger zooplankton size fractions. ST: station

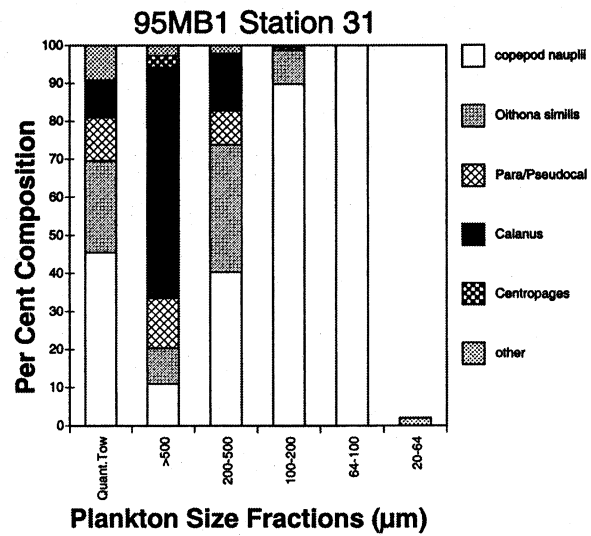


Fig. 5. Percent composition of dominant taxa in quantitative tows and zooplankton size fractions in qualitative tows from Cruise 95MB1, Stn 31. Para/Pseudocal: *Paracalanus parvus* and *Pseudocalanus* spp., Calanus: *Calanus finmarchicus*; Centropages: *Centropages typicus*

instance, on Cruise 95MB5, the contribution of *C. typicus* to the >500  $\mu\text{m}$  size fraction comprised 74.63% at Stn 10 (Fig. 12) and 47.6% at Stn 15 (Fig. 14), whereas respective contributions of *C. typicus* to total community abundance were only 5.56 and 4.80%. The combination of *C. typicus* plus *C. finmarchicus* at

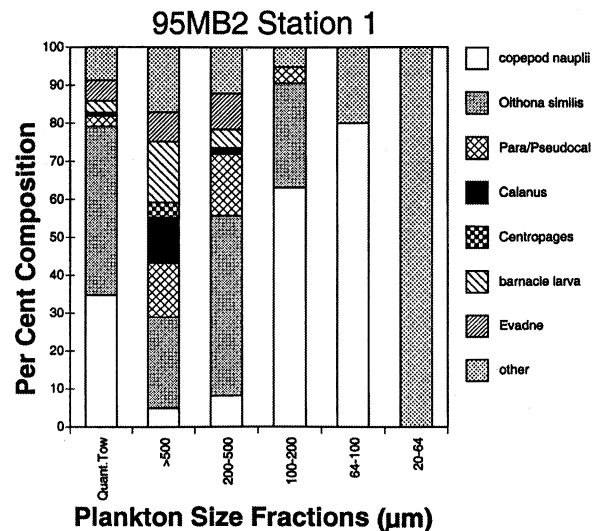


Fig. 6. Percent composition of dominant taxa in quantitative tows and zooplankton size fractions in qualitative tows from Cruise 95MB2, Stn 1. Abbreviations as in Fig. 5

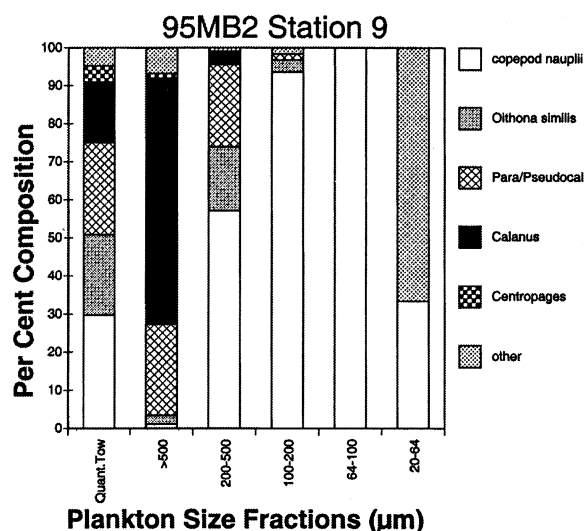


Fig. 7. Percent composition of dominant taxa in quantitative tows and zooplankton size fractions in qualitative tows from Cruise 95MB2, Stn 9. Abbreviations as in Fig. 5

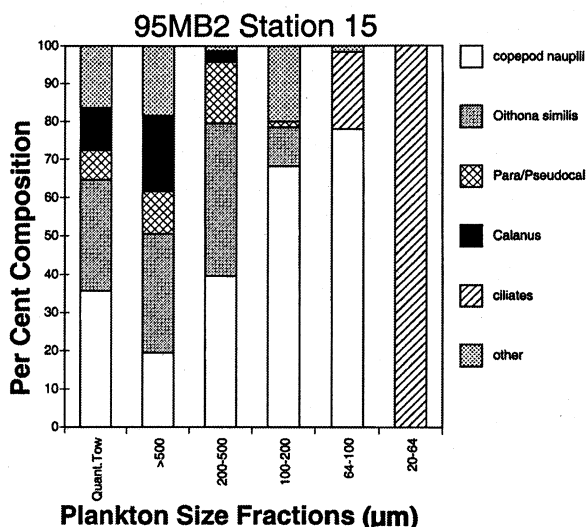


Fig. 9. Percent composition of dominant taxa in quantitative tows and zooplankton size fractions in qualitative tows from Cruise 95MB2, Stn 15. Abbreviations as in Fig. 5

Stn 11 (Fig. 13) comprised 75.7 % of the >500 µm fraction compared to only 20% of the total zooplankton assemblages represented by the quantitative tows. Toxin levels in the >500 µm fraction were also high relative to the smaller size classes at these stations. Similarly, at other stations with high toxicity, the copepods *T. longicornis* and *C. typicus* were more predominant (19.16 and 3.41%, respectively) in >500 µm fractions than in quantitative tow assemblages (3.19 and 0.23%,

respectively) (Cruise 95MB4, Stn 2, Fig. 11). There were, however, other stations with high toxicity where the relative proportions of large and small copepods in quantitative tows and >500 µm fractions were more similar to each other, such as on Cruise 95MB2 at Stns 1 (Fig. 6), 11 (Fig. 8), 15 (Fig. 9), and 19 (Fig. 10). Also, *C. finmarchicus* disproportionately dominated relative abundance in the >500 µm fraction at some stations such as Cruise 95MB1, Stn 31 (Fig. 5) and

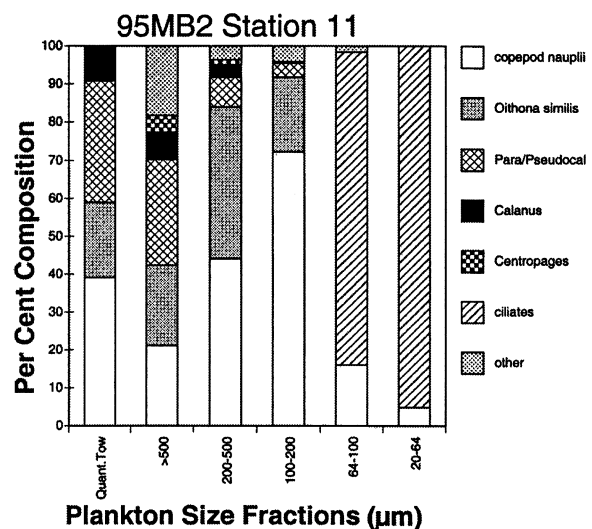


Fig. 8. Percent composition of dominant taxa in quantitative tows and zooplankton size fractions in qualitative tows from Cruise 95MB2, Stn 11. Abbreviations as in Fig. 5

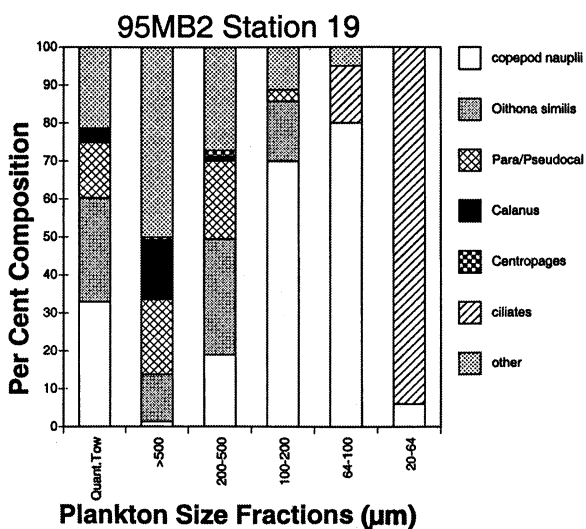


Fig. 10. Percent composition of dominant taxa in quantitative tows and zooplankton size fractions in qualitative tows from Cruise 95MB2, Stn 19. Abbreviations as in Fig. 5

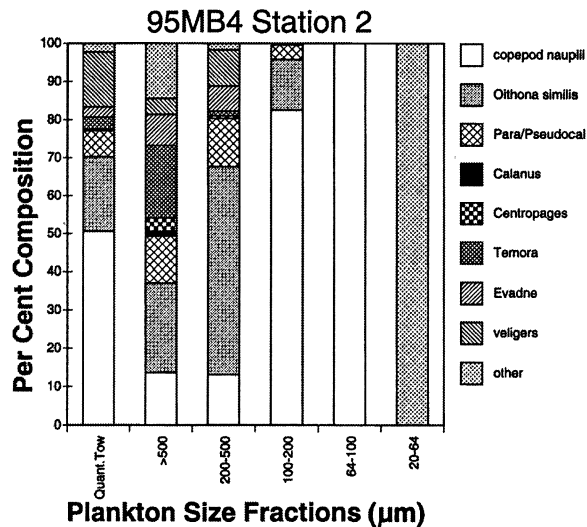


Fig. 11. Percent composition of dominant taxa in quantitative tows and zooplankton size fractions in qualitative tows from Cruise 95MB4, Stn 2. *Temora*: *Temora longicornis*; other abbreviations as in Fig. 5

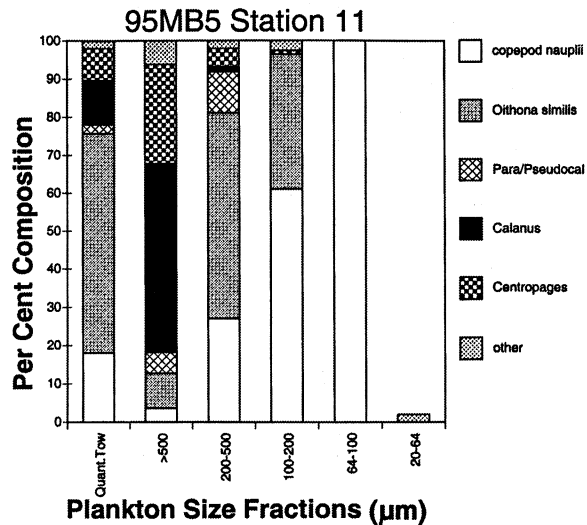


Fig. 13. Percent composition of dominant taxa in quantitative tows and zooplankton size fractions in qualitative tows from Cruise 95MB5, Stn 11. Abbreviations as in Fig. 5

Cruise 95MB2, Stn 9 (Fig. 7) (60.69 and 64.77%, respectively in the >500 µm fraction, compared to 9.59 and 15.95%, respectively in quantitative tows), even when toxicity in this fraction was low.

DISCUSSION

The discovery that the highest 'non-algal' PSP toxin levels occurred primarily in the larger, less-abundant

copepods of the >500 µm size fraction is an important finding. Our HPLC and receptor-assay data provide compelling evidence for the transfer of phytoplankton toxins to zooplankton size fractions dominated by large grazers, which are both direct consumers of *Alexandrium* cells, as well as prey of animals at higher trophic levels. Such a trend was unexpected because these larger copepods accounted for only a minor portion of total mesozooplankton abundance. Absolute confirmation of this route for PSP trophic toxin trans-

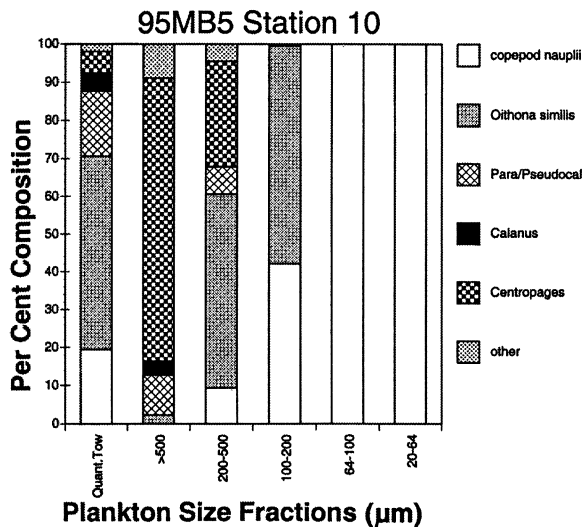


Fig. 12. Percent composition of dominant taxa in quantitative tows and zooplankton size fractions in qualitative tows from Cruise 95MB5, Stn 10. Abbreviations as in Fig. 5

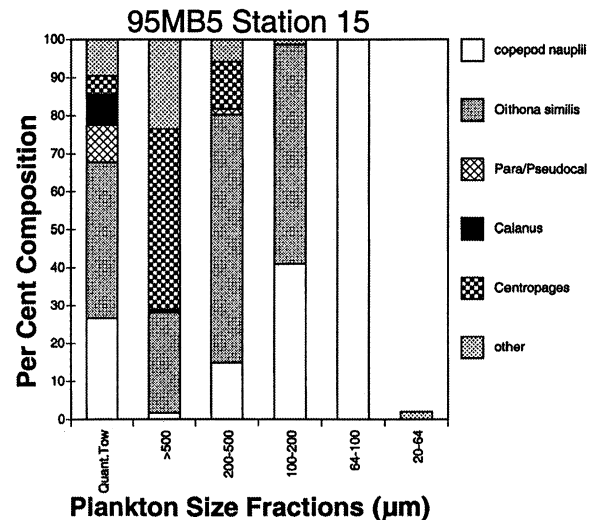


Fig. 14. Percent composition of dominant taxa in quantitative tows and zooplankton size fractions in qualitative tows from Cruise 95MB5, Stn 15. Abbreviations as in Fig. 5

fer by tandem mass spectrometry is currently being pursued.

The overwhelming numerical dominance of meso-zooplankton assemblages in Massachusetts Bay by copepod nauplii and small copepods such as *Oithona similis* has been noted previously (Turner 1994), but it appeared that these small metazoans did not play a direct role in the trophic transfer of PSP toxins to upper levels of the food web during our study. Although the nauplii- and *Oithona*-dominated 200 to 500  $\mu\text{m}$  size fraction exhibited measurable toxin levels in 6 of 10 samples, these values were generally far lower than for the  $>500 \mu\text{m}$  fraction. Similarly, the abundant protistan microzooplankton in most of the 64 to 100  $\mu\text{m}$  size fractions appear to be comparatively unimportant linkages in toxin trophic transfer.

The HPLC is currently the standard method employed for estimating concentrations of individual PSP toxin derivatives. In this comparative examination of field material encompassing both algal and zooplankton size fractions, we observed good agreement between HPLC and saxitoxin receptor-binding assay values. Our findings demonstrate that the PSP toxin activity detected by the receptor assay corresponds closely to HPLC-based estimates of these toxins. Moreover, the receptor assay's rapid, high-throughput format makes it especially useful for screening large numbers of samples and identifying those for which a more detailed, yet time consuming, HPLC analysis might be appropriate.

The receptor-binding assay shows great potential for monitoring toxicity in the water column. Receptor assay measurements of toxins extracted from the plankton were sensitive enough to detect toxic activity in the particulate fractions that were retained on sieves, even when no shellfish toxicity was recorded in shellfish at nearshore PSP monitoring stations in Massachusetts Bay. Reasons for this might include: (1) *Alexandrium* populations were generally offshore and not available to inshore shellfish, even during the downwelling period, and (2) based upon our experience in this region, *Alexandrium* abundances of 50 cells  $\text{l}^{-1}$  are not sufficient to cause shellfish toxicity measurable by mouse bioassay, the method used in shellfish-toxicity monitoring programs. Thus, for offshore waters where benthic shellfish may not be readily harvestable, or even in contact with near-surface blooms, receptor-assay measurements on plankton samples from net tows could serve to monitor for toxicity, even when *Alexandrium* cells are present at low abundance.

Although uptake and depuration rates of saxitoxins from zooplankters such as copepods are not well understood, it is likely that the toxins were transferred from *Alexandrium* cells to the larger-sized copepods

during the southerly transport of water within the Western Maine Coastal Current. Shellfish toxicity was much higher along the southern Maine coast during this time, suggesting that the possibility that *Alexandrium* populations might have been extensively grazed down by the time the water mass reached Massachusetts Bay (approx. 1 to 2 wk transit time depending on wind conditions). Evidence consistent with this scenario was particularly apparent during Cruise 95MB-2 (10 May), when maximum *Alexandrium* cell abundances in Massachusetts Bay (50 cells  $\text{l}^{-1}$ ) were still so low as to be near detection limits, but toxin levels at Stns 11, 15 and 19 were the highest recorded (Fig. 4).

Once toxins are sequestered in larger zooplankton, they may become less available to shellfish, since these zooplankton generally occur more offshore, away from nearshore shellfish beds. Even if these larger zooplankters are transported inshore during downwelling conditions, suspension-feeding bivalves generally do not eat them.

Since molecular evidence indicates that populations of the copepod *Calanus finmarchicus* are genetically homogeneous throughout the Gulf of St. Lawrence, Gulf of Maine, and Georges Bank regions (Bucklin et al. 1996), many of the copepods in Massachusetts Bay are also likely to be advected from the north by the Western Maine Coastal Current. If larger copepods substantially graze upon *Alexandrium* during this journey, this may prevent shellfish toxicity in the southern domain of the Western Maine Coastal Current in some years. Furthermore, since *C. finmarchicus* (Turriff et al. 1995) and *Centropages hamatus* (Turner et al. 1998a) have been shown to accumulate *Alexandrium* toxins through feeding, uptake of such toxins by large copepods may make those toxins more available to pelagic zooplanktivorous consumers such as baleen whales. More data on PSP toxin-depuration rates for zooplankton are needed to more critically evaluate this possibility.

Our finding of disproportionate levels of PSP toxins in larger zooplankton size fractions is of particular interest, since right whales *Eubalaena glacialis* observed in these waters in early spring are known to feed primarily on large copepods such as species of *Calanus* and *Centropages* (Mayo & Marx 1990). Also, humpback whales *Megaptera novaeangliae* occur in these waters in late spring and summer, and mortalities of humpback whales in the Cape Cod region (Geraci et al. 1989) were attributed to vectorial intoxication with *Alexandrium* toxins from ingested mackerel *Scomber scombrus*, which are known to feed primarily upon *Calanus finmarchicus* and other large copepods (Bigelow & Schroeder 1953, p. 320). Further, Nisbet (1983) recorded a kill of  $>70$  common terns

*Sterna hirundo* in the Cape Cod region following ingestion of zooplanktivorous sand lance *Ammodytes americanus* that were shown to contain PSP toxins from *Alexandrium* spp. Vectorial intoxication of other upper-trophic-level marine vertebrates with several other types of phycotoxins have been reviewed by Turner & Tester (1997) and Turner et al. (1998b).

The low toxin levels in microzooplankton size fractions are the inverse of the pattern reported by White (1979) during an *Alexandrium* bloom in the Bay of Fundy. There, toxin content progressively declined from smaller to larger zooplankton size fractions. These inconsistencies are likely to reflect real variability in patterns of toxin movement through zooplankton communities during different blooms. However, part of the explanation may also lie in the greater sensitivity for measurement of PSP toxins by HPLC and receptor-binding assay methods (~2 to 3 orders of magnitude), compared to the mouse bioassay methods that were used by White. White (1981) noted, for example, that in order to obtain reliable PSP measurements in zooplankton with the mouse bioassay, each sample required approx.  $10^9$  copepods. Our methods would have permitted detection of far lower PSP toxin levels contained in larger zooplankton size fractions comprising a small percentage of the total zooplankton numerical abundance.

In summary, we have presented compelling evidence that PSP toxins from *Alexandrium* can accumulate in larger-sized copepods in Massachusetts Bay. This occurred at a time when *Alexandrium* cell abundance was minimal, and there was no recorded shellfish toxicity in the sampling area. Our findings suggest that toxin accumulation in copepods may have occurred through grazing during southward transport of *Alexandrium* cells in the Western Maine Coastal Current. The ability to detect low levels of PSP toxins in zooplankton was facilitated by sensitive HPLC analyses and receptor-binding assays. Disproportionate accumulation of toxins in larger, less-abundant copepods may pose a threat of vectorial intoxication to endangered whales that inhabit these waters, and feed upon such copepods, during the spring-summer red tide season.

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