A red tide of *Alexandrium fundyense* in the Gulf of Maine

D.J. McGillicuddy Jr.[a,*], M.L. Brosnahan[b], D.A. Couture[c], R. He[d], B.A. Keafer[b], J.P. Manning[e], J.L. Martin[f], C.H. Pilskaln[g], D.W. Townsend[h], D.M. Anderson[b]

[a] Department of Applied Ocean Physics and Engineering, Woods Hole Oceanographic Institution, Woods Hole, MA 02543, USA
[b] Biology Department, Woods Hole Oceanographic Institution, Woods Hole, MA 02543, USA
[c] Resource Access International, Brunswick, ME 04011, USA
[d] Department of Maine, Earth, and Atmospheric Sciences, North Carolina State University, Raleigh, NC 27695, USA
[e] National Oceanic Atmospheric Administration, Northeast Fisheries Science Center, Woods Hole, MA 02543, USA
[f] St. Andrews Biological Station, Fisheries and Oceans Canada, St. Andrews, NB, Canada E5B 2L9
[g] School of Marine Sciences, University of Massachusetts Dartmouth, North Dartmouth, MA 02747, USA
[h] School of Marine Sciences, University of Maine, Orono, ME 04469, USA

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ABSTRACT

In early July 2009, an unusually high concentration of the toxic dinoflagellate *Alexandrium fundyense* occurred in the western Gulf of Maine, causing surface waters to appear reddish brown to the human eye. The discolored water appeared to be the southern terminus of a large-scale event that caused shellfish toxicity along the entire coast of Maine to the Canadian border. Rapid-response shipboard sampling efforts together with satellite data suggest the water discoloration in the western Gulf of Maine was a highly ephemeral feature of less than two weeks in duration. Flow cytometric analysis of surface samples from the red water indicated the population was undergoing sexual reproduction. Cyst fluxes downstream of the discolored water were the highest ever measured in the Gulf of Maine, and a large deposit of new cysts was observed that fall. Although the mechanisms causing this event remain unknown, its timing coincided with an anomalous period of downwelling-favorable winds that could have played a role in aggregating upward-swimming cells. Regardless of the underlying causes, this event highlights the importance of short-term episodic phenomena on regional population dynamics of *A. fundyense*.

1. Introduction

Although the term "red tide" is frequently used in reference to harmful algal bloom events, its use to describe blooms of *Alexandrium fundyense* in the Gulf of Maine is largely a misnomer. Concentrations of *A. fundyense* seldom reach levels sufficient to discolor the water, and this species typically constitutes a small fraction of the total phytoplankton biomass. However, there have been some exceptions, including the historic bloom of 1972 during which *A. fundyense* (formerly *Gonyaulax tamarensis*) discolored the water offshore of the northern Massachusetts and New Hampshire coastlines (Hartwell, 1975; Mulligan, 1973; Sasner et al., 1974). *A. fundyense* also discolored water in the Bay of Fundy in 1980, 2003, and 2004 (Martin et al., 2008; Martin and White, 1988).

An unusual "red tide" of *A. fundyense* occurred in the Gulf of Maine in 2009. Initial discovery of the anomaly was serendipitous, taking place on a mooring turnaround cruise prompted by an increase in Paralytic Shellfish Poisoning (PSP) toxicity along the

Maine coast in late June–early July. Visual observations of discolored water prompted surface sampling to and from the mooring site, revealing *A. fundyense* concentrations ranging from hundreds of thousands of cells l^{-1} to in excess of one million cells l^{-1}. This triggered a rapid-response sampling effort, both at sea and via aerial survey. Herein we characterize the phenomenology of this event utilizing these observations together with satellite imagery, shellfish toxicity measurements, flow cytometric analysis of samples from the discolored water, as well as cyst fluxes from a nearby sediment trap and a spatial survey of cysts in coastal sediments the following October. Through synthesis of this diverse set of observations, it is clear that this was a significant event not only in terms of coastal shellfish toxicity but also the regional population dynamics of *A. fundyense*.

2. Methods

2.1. Hydrography

Hydrographic profiles and water samples were collected with a standard CTD-rosette system with Niskin bottles. *A. fundyense*

* Corresponding author. Tel.: +1 508 289 2683; fax: +1 508 457 2194.
 E-mail address: dmcgillicuddy@whoi.edu (D.J. McGillicuddy Jr.).

samples were collected by sieving 2 l through a 20 μm Nitex screen that was washed into a 15 ml centrifuge tube and fixed in 5% formalin for < 24 h. The samples were then centrifuged, the supernatant removed, and ice-cold methanol added. Samples were stored at −20 °C for later enumeration in the laboratory (Section 2.2 below). An additional 10 l sample of surface water was sieved in the same manner for the purposes of an on-board "live" count. Nutrient samples were filtered through Millipore HA filters, placed immediately in a sea water-ice bath for 5–10 min, and frozen at −18 °C. Concentrations of NO_3+NO_2, NH_4, $Si(OH)_4$ and PO_4 were measured ashore following each cruise with a Bran Luebbe AA3 AutoAnalyzer using standard techniques.

2.2. Moored observations

A McLane Laboratories Inc. autonomous Phytoplankton Sampler (PPS) obtained measurements at 5 m depth from April to September in two contiguous deployments. The mooring was located in the vicinity of the Northeastern Regional Association of Coastal and Ocean Observing Systems (NERACOOS) buoy B at 43° 11′N, 70° 26′W (Fig. 1B). A total of 24 samples were taken per deployment, at a frequency of one every 2–3 days. For each sample, the instrument filters 2 l of seawater onto a 15 μm Nitex screen. The instrument was prepared with 10% formalin dissolved in artificial seawater that was adjusted to be lighter (specific density=1.018) than the ambient sample seawater. During the automated filtering process, the "light" 10% formalin solution dispensed into the bottom of the filter reservoirs is diluted and displaced upward onto the filter by the heavier ambient sample seawater. This process resulted in a 1/2 reduction in the concentration of preservative to about 5% final. After the instrument was recovered, the preserved > 15 μm samples were backwashed off the Nitex screen into a 50 ml centrifuge tube for further concentration and A. fundyense cell counting as described in Section 2.3.

2.3. Enumeration of A. fundyense cells and cysts

A. fundyense cells were enumerated from water samples using a species-specific oligonucleotide probe and methods described in Anderson et al. (2005c). Both A. tamarense and A. fundyense occur in the Gulf of Maine, and these are considered to be varieties of the same species (Anderson et al., 1994; Brosnahan et al., 2010; Scholin et al., 1995). Available molecular probes cannot distinguish between them, and only detailed analysis of the thecal plates on individual cells can provide this resolution—which is not practical for large numbers of field samples. Accordingly, for the purpose of this study, the name A. fundyense is used to refer to both forms.

Surface "live" counts consisted of two transects across a Sedgewick-Rafter counting slide using an on-board light microscope at 200x magnification. The slide was loaded with 1 ml of concentrated sieved material (see Section 2.1 above) resuspended to 14 ml. This provided a lower limit of detection of 14 cells l^{-1}. Light microscopy of this kind cannot distinguish A. fundyense from the morphologically similar non-toxic species A. ostenfeldii, and therefore the live counts can sometimes overestimate A. fundyense concentration.

Cysts of A. fundyense were collected and enumerated from sediment samples using primulin-staining methods described in Anderson et al. (2005d). Samples were obtained with a Craib corer in dedicated surveys in fall 2008, 2009, and 2010. The sampling pattern consisted of 14 cross-shore transects in the coastal Gulf of Maine and three transects across Georges Bank, for a total of approximately 120 stations. A. fundyense cysts from the upper 1 cm of oxygenated sediment are viable for germination (Anderson et al., 2005d) and thus only that vertical fraction of the sediment samples is presented herein.
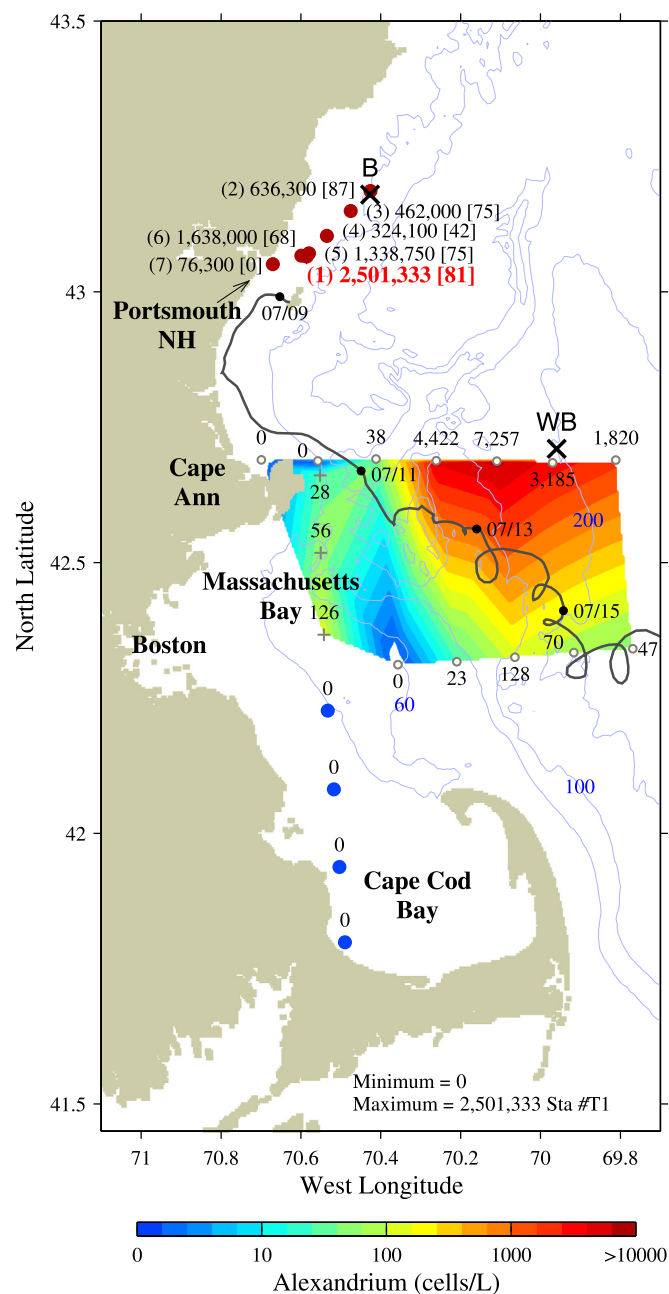

Fig. 1. Dots north of 43°N: surface A. fundyense concentrations (cells l^{-1}) observed on July 10, 2009 (whole cell counts). Station numbers (in parentheses) precede the cell counts, and the percentage of planozygotes follows in brackets. Map, dots, and plus signs south of 43°N: surface A. fundyense concentrations observed on R/V Tioga 383 July 12, 2009 (live counts). Black crosses indicate the locations of NERACOOS mooring "B" where the McLane PPS sampler was located, and the Wilkinson Basin "WB" sediment trap. Trajectory of surface drifter #97201 released on July 9 is plotted as a gray line, with dates provided every two days along track.

Cyst fluxes were measured in time-series sediment traps (Honjo and Doherty, 1988) deployed on subsurface moorings. See Pilskaln et al. (2014) for a complete description of these data. Of particular interest to this study are the traps located in Wilkinson Basin located at 42° 43′N, 69° 58′W (Fig. 1, WB). The traps had a baffled surface collection area of 0.5 m^2, and collected time-series samples in thirteen 250 ml volume cups per deployment. Prior to deployment, trap cups were pre-poisoned with an 8% density-adjusted formalin solution in filtered seawater buffered to a pH of 8.0–8.1. Recovery and redeployment of the trap moorings occurred approximately every 5–9 months, with individual cup collection periods

varying from ∼10–20 days. The traps were programmed to insure that the cups on the upper and lower traps rotated and collected sinking particulate material on the same time interval. Trap samples were processed by removing the formalin solution with sieving through a 20 μm Nitex screen. The material retained on the screen was then treated as a sediment sample using the same primulin-staining protocols noted above (Anderson et al., 2005d). Only primulin-labeled, capsule-shaped cysts with intact internal cellular contents were counted; empty cysts were not quantified.

2.4. Flow cytometric analysis

Subsamples of 10,000 cells from surface samples taken on the July 10 mooring turnaround cruise and from a log-phase laboratory culture of a Gulf of Maine *A. fundyense* isolate (clone 38-3) were stained with propidium iodide (PI) for DNA content analysis by flow cytometry. All of the samples were fixed with 5% formalin and resuspended in ice-cold methanol prior to staining. Methanol was removed by centrifuging the cell suspensions for 10 min at 3000 rcg then aspirating the methanol supernatant. The cell pellets were then washed by resuspension in 1 ml of PBS (40 mM Na_2HPO_4, 22 mM KH_2PO_4, 85 mM NaCl), and centrifuged again for 10 min at 3000 rcg. The overlying PBS solution was then aspirated before the cells were stained in PBS amended with 4 μg ml^{-1} PI and 100 Kunitz ml^{-1} RNAse A for at least 3 h in darkness and at room temperature. At least 2000 particles having PI-associated fluorescence comparable to *A. fundyense* culture cells were recorded from each field sample with a FACSCalibur flow cytometer (BD Biosciences). These observations were used to compute relative frequency distributions of FL2-H band (PI-associated) fluorescence, which provides a quantitative measure of DNA content. All measurements were made with linear signal amplification.

The DNA content of *A. fundyense* cells changes in a quantized fashion as they progress through the phases of the cell division cycle (mitosis) and also the gamete and zygote stages of its sexual life cycle. Haploid vegetative *A. fundyense* oscillate between 1c and 2c DNA content as they undergo mitosis; 1c G1 phase cells become 2c G2 phase cells when they replicate their DNA, then return to the 1c G1 phase when they divide. The G1 phase is long relative to the G2 phase in *A. fundyense* and divisions are often phased by the diel light/dark cycle, such that most occur at daybreak (Rubin, 1981; Taroncher-Oldenburg et al., 1997). As a consequence, growing populations of *A. fundyense* tend to be dominated by 1c G1 phase cells, especially in the late-morning and afternoon (Brosnahan et al., 2014).

In contrast, sexual-phase cells do not oscillate between DNA content levels, but instead increase in DNA content as they progress from one stage to the next—first as 1c gametes, then as 2c planozygotes—until eventually transforming to resting cysts and undergoing pre-meiotic replication to become 4c (or greater) in DNA content. The last transition to 4c DNA content has not been observed directly but is inferred from the resumption of mitosis by haploid, vegetative cells after cyst germination (Pfiester and Anderson, 1987). Because gametes have the same DNA content (1c) as G1 phase vegetative cells and planozygotes have the same content as G2 phase cells, DNA content alone cannot be used to determine the life cycle stage of an *A. fundyense* cell (Cetta and Anderson, 1990). However, the presence of a large fraction of 2c cells within a population can signal a shift from vegetative cell division to the formation of planozygotes, especially when high concentrations of 2c cells are observed during afternoon hours (Brosnahan et al., 2014).

2.5. Satellite observations

A variety of satellite data products were examined to determine their suitability for assessing the spatial and temporal scales of the red tide event. Of all that were considered, the Medium Resolution Imaging Spectrometer (MERIS) case 2 water algorithm for chlorophyll (Doerffer and Schiller, 2007) showed closest correspondence with the discolored water observed on July 10. Of course this algorithm is not specific to *A. fundyense*, and its relevance for this purpose is restricted only to those rare occasions when *A. fundyense* constitutes a significant fraction of the total phytoplankton biomass. There are many instances throughout the MERIS data record when such imagery indicates the presence of high biomass during times when *A. fundyense* concentrations were known to be low (data not shown).

2.6. Drifter tracks

Surface drifters consisted of a 1.3 m long, 0.8 cm diameter PVC cylinder (conventional pipe material) that supports two pairs of fiberglass rods. The rods were mounted radially and orthogonally to hold a set of four vinyl-cloth sails. The cylinder was ballasted so that only the GPS antenna and a small portion were exposed to the wind. While the design is essentially the same as the commercially available Davis-style ("CODE") surface drifters (Davis, 1985), the electronics were replaced with new technology used for tracking vehicles on highways. The units were set to report every 0.5–2 h and communicate via the GLOBALSTAR satellite system. Drifter position data were processed to eliminate obvious bad points according to the methods described by Hansen and Poulain (1996). Data are available via the Open-source Project for a Network Data Access Protocol (OPeNDAP) at (http://www.nefsc.noaa.gov/epd/ocean). For more information see Manning et al. (2009).

2.7. Shellfish toxicity

Shellfish toxicity measurements were based on the blue mussel *Mytilus edulis*, using the standard mouse bioassay (Association of Official Analytical Chemists, 1984; AOAC Official Method 959.08). These data were kindly provided by the Maine Department of Maine Resources (http://www.state.me.us/dmr/). A set of coastal locations are monitored on a weekly basis, and shellfish beds are closed when the mouse bioassay approaches the quarantine level of 80 μg saxitoxin equivalents (STX) per 100 g of shellfish tissue.

3. Results

3.1. Surface distributions of A. fundyense vegetative cells

Surface samples collected in between Portsmouth NH and NERACOOS mooring B on 10 July 2009 documented *A. fundyense* concentrations ranging from 10^4 cells l^{-1} to in excess of 10^5 cells l^{-1} in offshore waters (Fig. 1). *A. fundyense* was considerably less abundant in a near-shore sample, although the observed concentration of 25,690 cells l^{-1} far exceeds the 200–1000 cells l^{-1} that typically leads to toxicity in shellfish sufficient to require regulatory closure in the region. These observations prompted a rapid-response survey cruise on board R/V Tioga on July 12. Sampling consisted of (1) underway surface counts on a south-to-north line in transit to Cape Ann, (2) full hydrographic stations along a transect off Cape Ann, and (3) full hydrographic stations on the eastern half of a transect off Boston (Fig. 1). Cell counts were low in Massachusetts and Cape Cod Bays. The western part of the Cape Ann line was devoid of cells, but offshore the numbers were high, peaking at over 7000 cells l^{-1}. Due to time constraints it was not possible to delimit

the offshore edge of the population, as the easternmost station was near 2000 cellsl^{-1}. Given the southward flow characteristic of this area, it seems logical to infer that this offshore population was connected with the extremely high concentrations in the discolored water observed along the coast to the north. In fact, the trajectory of a surface drifter released on July 9 just south of where the red water was observed on July 10 illustrates oceanographic connectivity between the two sets of observations on precisely the right time scale (Fig. 1). Thus these data suggest the southern extent of the population had not been transported much further south than the Cape Ann line, at least in surface waters.

A regional-scale mapping effort was conducted on voyage #386 of R/V Tioga July 19–23. Cell counts were very low overall, with the 100 cells l^{-1} threshold broken only in a few places along transects off Casco Bay, Isle au Haut, and Southwest Harbor (Fig. 2). Fine-scale sampling of a near-coastal area with high toxicity (Southwest Harbor, Frenchman Bay, Winter Harbor, Schoodic Point, and Prospect Harbor) also failed to detect cell concentrations in excess of 200 cells l^{-1} (Fig. 2 inset). Nutrient concentrations were low throughout the region sampled during the survey, with dissolved nitrate plus nitrite concentrations at or below detection limits in the upper 10 m (ca. 1.0 μM or less); however, ammonium concentrations were relatively high, reaching 2 μM at the surface and 10 m at the inner-most stations of the western gulf (not shown).

3.2. Aerial survey

Shipboard sampling on July 19 was accompanied by an aerial survey executed by spotter pilot Norman St. Pierre and observer Michael Brosnahan. Atmospheric conditions were ideal, and altitude was maintained at 200–500 m on tracks from the coast out to 10 km offshore. The survey spanned coastal areas from Cape Cod Bay to Bar Harbor, and water coloration indicative of high concentrations of *A. fundyense* was not found.

3.3. Satellite observations

A MERIS case 2 chlorophyll image from July 9 (Fig. 3, top) indicates the presence of a chlorophyll anomaly precisely where the discolored water was observed on July 10. The region of enhanced chlorophyll extends along the coast southward to Cape Ann and northeast to the western side of Penobscot Bay. More modest enhancements are evident east of Penobscot Bay along the Maine coast and up into the Bay of Fundy, although they appear as more isolated patches rather than a continuous feature. A time-series of images zoomed in on the region of discolored water illustrates the highly-ephemeral nature of this phenomenon (Fig. 3, bottom). There was no trace of the feature on July 5, and isolated patches appeared on July 6. Concentrations peaked on July 9, and were on the decline by July 11. By July 12–15 the surface expression had completely disappeared.

3.4. Toxicity measurements

As of the week of June 15, PSP toxicity patterns observed along the Maine coast were fairly typical: toxicity was on the decline in western Maine, and had yet to rise in eastern Maine (Fig. 4). By the week of June 22, toxicity began to rise in far eastern Maine,
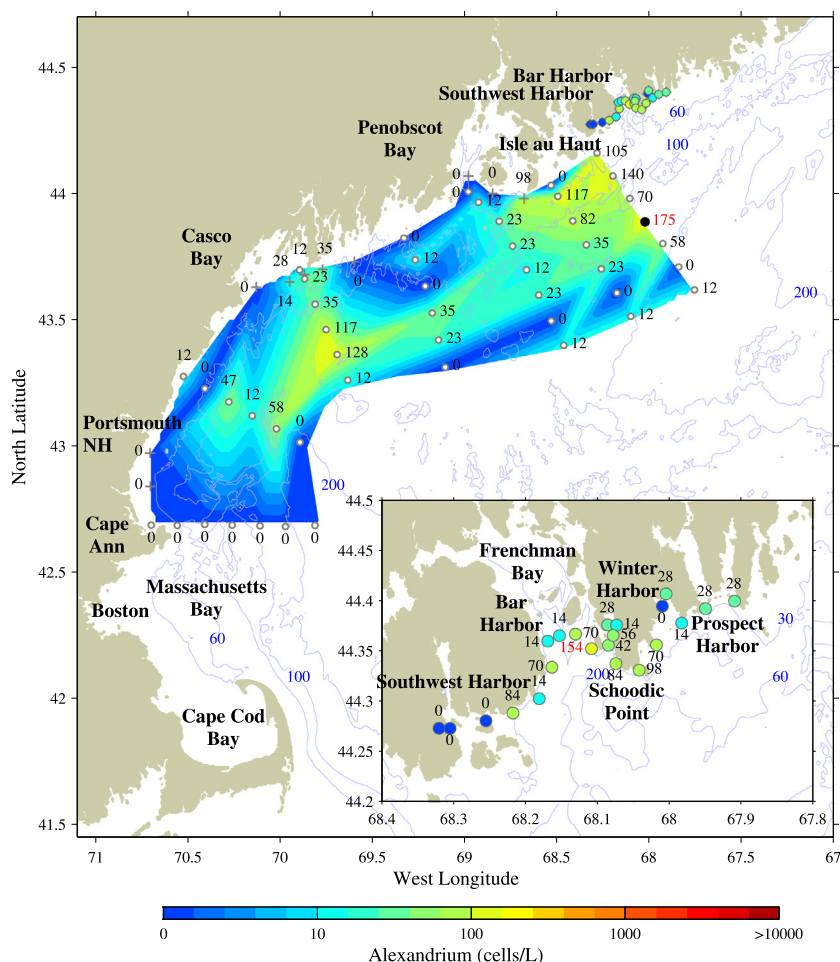

Fig. 2. Surface of *A. fundyense* concentrations observed on R/V Tioga 386 July 19–23, 2009 (live counts). Inset shows a zoom view of the near shore underway data collected at the northern terminus of the survey.
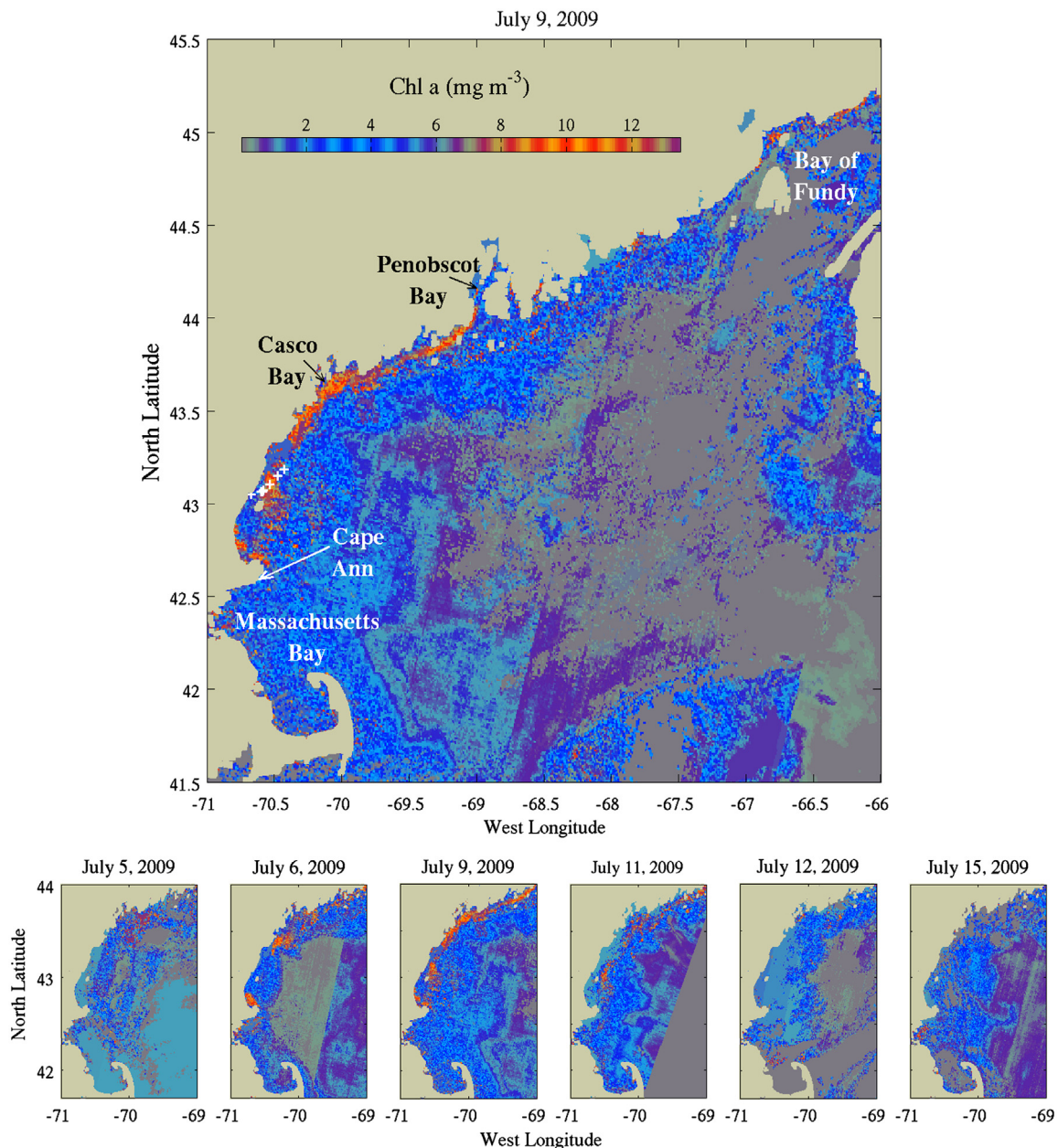
Fig. 3. Top: MERIS "Algal 2" image for July 9, 2009 depicting chlorophyll for case-2 waters. White crosses indicate the positions of *A. fundyense* measurements in the discolored water (Fig. 1). Bottom: time-series of images zoomed into the area of discolored water.

whereas it was still relatively low in western Maine. During the week of June 29, toxicity continued to rise in far eastern Maine, and low-level toxicity was detected nearly coast-wide at the outermost points and islands. A coast-wide onset of toxicity continued during the week of July 6, reaching its peak the week of July 13. By the week of July 20 toxicity was on the decline, with highest values occurring in western Maine where peak toxicities were highest. Toxicity declined further the week of July 27, and by the week of August 3 the episode was essentially over.

Summer 2009 was unusual for *A. fundyense* and shellfish toxicity in the Bay of Fundy, particularly in Passamaquoddy Bay. Although shellfish in most years become toxic in Passamaquoddy Bay, it is usually at low levels—although there has been the occasional year that levels have not gone above the threshold level and shellfish beds have remained opened to harvesting. Shellfish toxicities prior to 2009 and following 2009 have always

been among the lowest anywhere in the Bay of Fundy. 2009 was the first year since sampling began that *Mya arenaria* toxicity values exceeded > 1000 µg STX equiv. 100 g meat. On June 20 *Mya* toxicity was 41 µg STX equiv. 100 g meat; and the next measurement on July 2 was 4120 µg STX equiv. 100 g meat. The following week (July 7) toxicity values had decreased to 441 and on July 14 were 130 µg STX equiv. 100 g meat.

In a time-series dating back to 1988, *A. fundyense* cells have been observed at Brandy Cove in Passamaquoddy Bay each year (Fig. 5). In all years except 2009, concentrations were considerably lower than elsewhere in the Bay of Fundy outside Passamaquoddy Bay. Weekly sampling at Brandy Cove indicated that on June 16, 2009, the concentration of *A. fundyense* was 288 cells l^{-1}. By June 23, it had increased to 5616 cells l^{-1}, and the following week on June 30 concentrations had increased to 2.79×10^5 cells l^{-1}. On July 7, one week later they had decreased to 2480 cells l^{-1}.
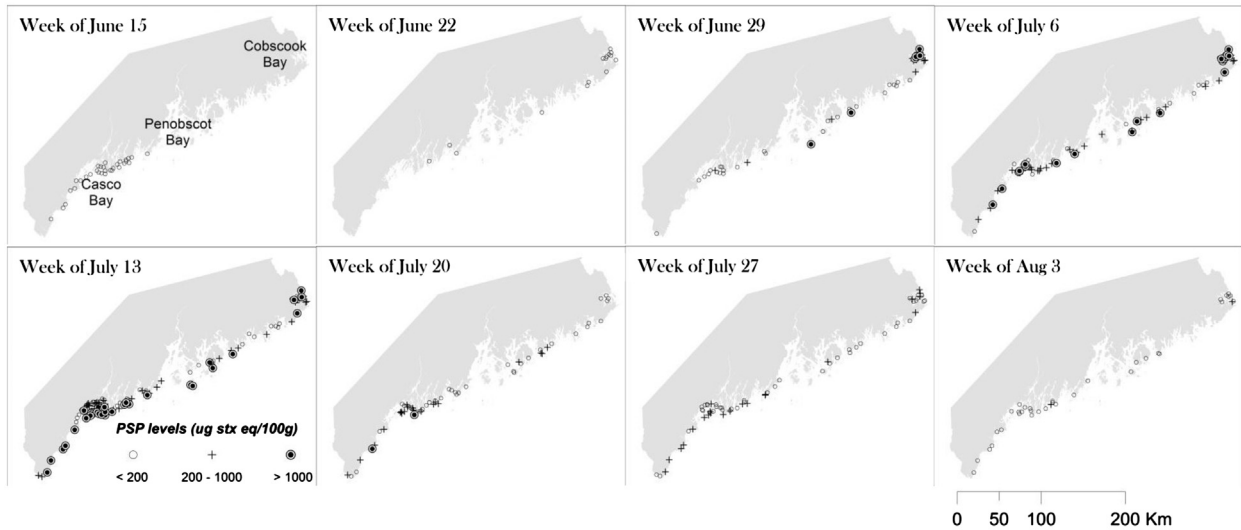
Fig. 4. Weekly toxicity maps for the coast of Maine from mid-June to early August, 2009.
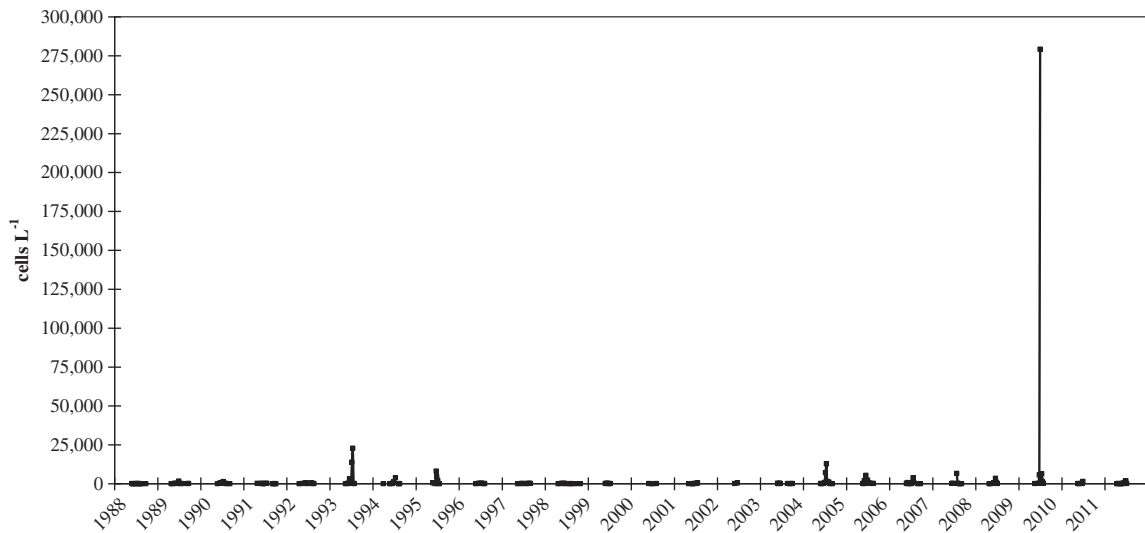

Fig. 5. *A. fundyense* concentrations from weekly sampling at Brandy Cove, Passamaquoddy Bay from 1988 to 2011.

Interestingly, *A. fundyense* concentrations at Brandy Cove and shellfish toxicity at Bar Road in Passamaquoddy Bay were the highest for the whole Bay of Fundy in 2009—a first on record.

3.5. Moored time-series

The McLane PPS time-series of vegetative cells corroborates the highly-ephemeral nature of the red tide event (Fig. 6). *A. fundyense* concentrations began to rise on July 7, peaked at the very next sample on July 11, and were back to background levels by the 17th of July. Note that the peak concentration observed at 5 m depth is two orders of magnitude smaller than that measured from near-surface samples obtained on July 10 from R/V Tioga (Fig. 1). This suggests the cells were highly concentrated near the surface, above the intake port of the PPS sampler. This extreme layering of the population was facilitated by calm conditions during that time, in which wind-driven mixing was minimal. Unfortunately there are no vertical profiles available within the discolored water to characterize the vertical distribution in detail.
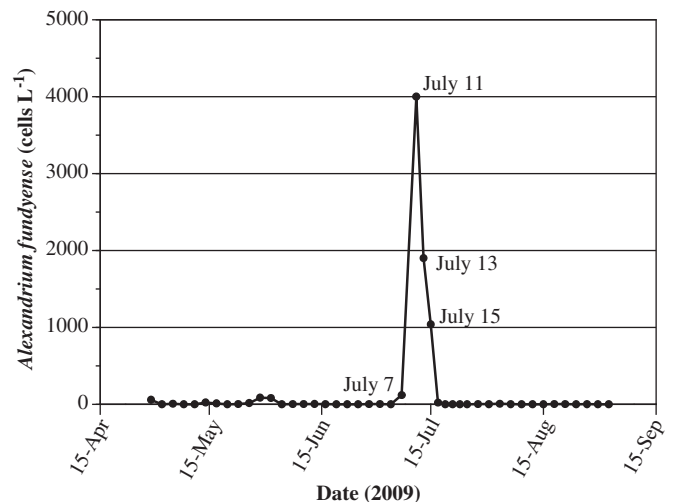

Fig. 6. Time-series of *A. fundyense* cell concentrations at 5 m depth collected from the McLane PPS moored at 43° 11′N, 70° 26′W (Fig. 1, "B").

3.6. Flow cytometric analysis of the red tide population

DNA-associated fluorescence of 1c and 2c *A. fundyense* cells from field samples was assessed by comparison to a log-phase culture sample that contained abundant 1c (G1 phase) and scarcer 2c (G2 phase) vegetative cells (Fig. 7). The correspondence of fluorescence modes between the culture and field samples was quite high: all 1c modes from field samples occurred between 305 and 355 FL2-H units versus 347 in the culture sample, and 2c modes from field samples occurred between 581 and 695 versus 618 in the culture sample. However, in contrast to the culture sample, 2c cells were much more abundant than 1c cells in all of the red water samples taken on July 10 (Fig. 1, stations 1–6 near NERACOOS mooring B).

The proportion of cells that were 2c was estimated by counting the number of cells in each sample that had FL2-H measurements greater and less than 450 units. By this criterion, 2c cells were greater than 95% of those sampled at 4 of the 6 red water stations (1, 2, 5 and 6), and 93% and 80% of those sampled at the other two (3 and 4). Station 7, which was outside the area where discolored water was observed, had a smaller proportion of 2c cells (47%).

This indication that a large fraction of the population in the discolored water was comprised of planozygotes was confirmed with traditional microscopy. The species-specific counting method described in Section 2.3 can be used to distinguish planozygotes from vegetative cells to the trained eye based on size and staining characteristics. That method of enumerating planozygotes, albeit less precise, also 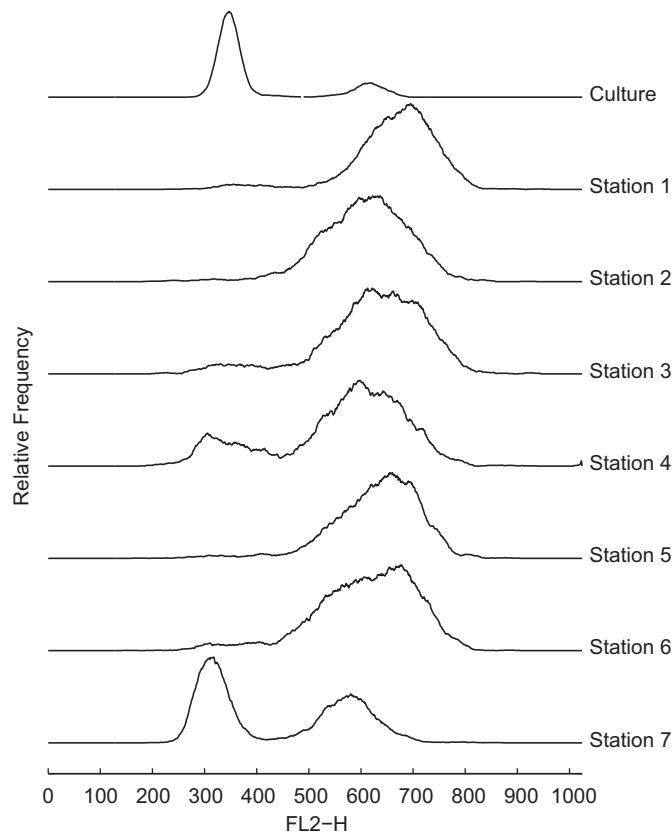indicated a high fraction of planozygotes in the discolored water offshore, and found no evidence of planozygotes at the inshore station (not shown).

3.7. Cyst fluxes

The red-tide event was followed by extremely large fluxes of cysts measured nearby in Wilkinson Basin (Fig. 8). Peak fluxes of 75,236 cysts $m^{-2} d^{-1}$ at 95 m and 292,894 cysts $m^{-2} d^{-1}$ at 180 m were 4000–5000 times larger than the median fluxes recorded in those time series. This cyst flux was the largest measured in all of the traps deployed in the Gulf of Maine in 1995–1997 and 2005–2009 as reported in Pilskaln et al. (2014).

The peak cyst flux at 95 m was a single point corresponding to the sampling interval July 9–19. A peak occurred simultaneously in the 180 m trap, but the flux persisted at nearly the same level for the subsequent sampling interval July 19–30. This is consistent with a longer residence time of cysts in the benthic nepheloid layer where near-bottom turbulence resuspends sedimentary material (Pilskaln et al., 2014).

3.8. Cyst maps

The abundance of cysts in coastal sediments increased dramatically from 2008 to 2009 (Fig. 9). Integrated abundance in the top 1 cm layer of sediment in 2009 was the highest observed in the yearly time-series from 2004 to 2009 (McGillicuddy et al., 2011) and measurements thereafter in 2010 and 2011 (Anderson et al., 2014). In addition to the overall increase in abundance, the western Gulf of Maine cyst bed (also known as the mid-coast Maine cyst bed) spread farther south than in all prior observations. Cyst concentrations in excess of 1000 cysts cm^{-3} extended south and east of Cape Ann, mostly beyond the 200 m isobath.
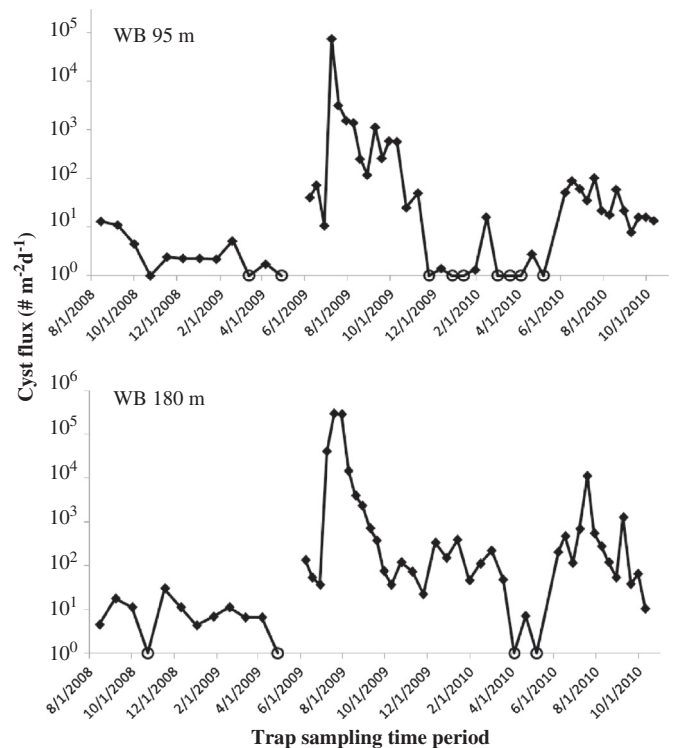

Fig. 7. Relative frequency of FL2-H (DNA-associated) fluorescence from flow cytometry analysis of a log-phase culture of vegetative *A. fundyense* (top curve) and red tide samples taken from stations 1–7 on July 10 (Fig. 1). The frequency distributions have been smoothed and are plotted as deviations from zero (e.g. no cells were observed with FL2-H fluorescence less than 200). Stations 1–6 are the offshore locations where *A. fundyense* concentrations were sufficient to discolor the water (highest *A. fundyense* concentration at Station 1), whereas station 7 was the near-coastal location where the concentration of *A. fundyense* was 25,690 cells l^{-1}.

Fig. 8. Cysts fluxes measured at the northern Wilkinson Basin site (Fig. 1, "WB") at 95 m (top) and 180 m (bottom). Open circles indicate zero cyst flux/no cysts collected in the trap cup during the particular collection period. Peak cyst fluxes in 2009 occurred in consecutive samples 4–7 of the deployment starting in June 2009. The time intervals for samples 4–7 were: July 9–19, July 19–30, July 30–August 9, August 9–19. Note that each dot in the time-series is plotted at the beginning of each sampling interval. Modified from Pilskaln et al. (2014).
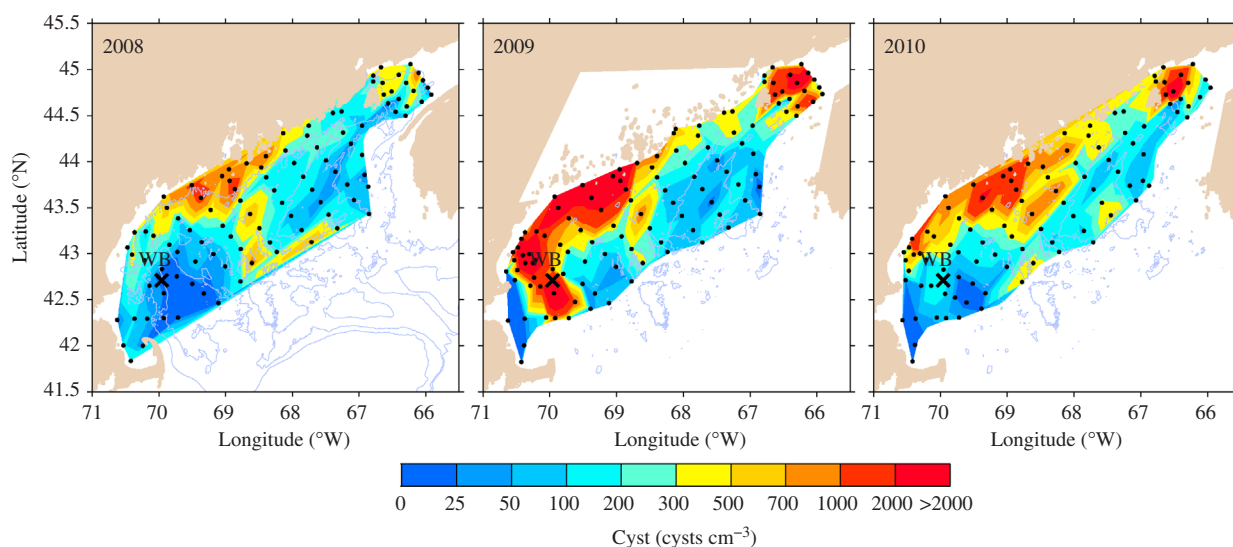
Fig. 9. *A. fundyense* cyst abundance in the upper 1 cm layer of sediment observed in October 2008 (left), 2009 (middle), and 2010 (right). Black dots denote the locations of sediment samples used to construct the maps. Location of the Wilkinson Basin "WB" sediment trap is indicated by a black cross.

The southward tongue of cysts deposited in 2009 had disappeared by the time the same area was sampled again in 2010 (Fig. 9, right panel), as the southern terminus of the western Gulf of Maine cyst bed returned to its more characteristic position north and east of Cape Ann. See Anderson et al. (2014) for more details on cyst dynamics in this region.

4. Discussion

What conditions led to the unusual red-tide event observed in July 2009? We examined the observations described herein, as well as data from the coastal ocean observing system (see Li et al., 2014), and were unable to discern an unequivocal causal factor. However, one aspect did stand out as clearly anomalous: wind forcing. As in Li et al. (2014), we define an upwelling index $UI = \tau_x / \rho f$ following the method of Schwing et al. (1996), where τ_x is the alongshore component of the wind stress calculated using Large and Pond (1981), ρ is the density, and f is the local Coriolis parameter. Positive (negative) UI represents upwelling- (down-welling-) favorable wind conditions. The cumulative UI (CUI) was computed by integrating the resulting UI over time (i.e., $CUI = \int UI dt$) between April 1 and August 1 for each year, 2004–2011 (Fig. 10). The slope of CUI is particularly informative, as upwelling- (downwelling-) favorable wind conditions are represented by a rising (declining) temporal trend in CUI. Wind speeds of zero or winds oriented in the cross-shore direction would cause no change in the CUI, or a slope of zero during such periods. In every year examined except for 2009, the June–July time period is characterized by upwelling-favorable winds driven by the prevailing summertime southwesterlies in this region. In contrast, winds were generally downwelling-favorable from late June through early July 2009. A similar downwelling-favorable trend was evident during the same time period in the eastern Gulf of Maine at NERACOOS Buoy I (Li et al., 2014), suggesting this was a regional pattern.

Wind forcing is a key regulator of *A. fundyense* transport, insofar as upwelling-favorable winds tend to transport near-coastal populations offshore, and downwelling-favorable winds tend to transport offshore populations shoreward (Anderson et al., 2005a; Franks and Anderson, 1992a, b; Hetland et al., 2002; McGillicuddy et al., 2003). As such, coincidence of the anomalous episode of downwelling-favorable winds in late June/early July
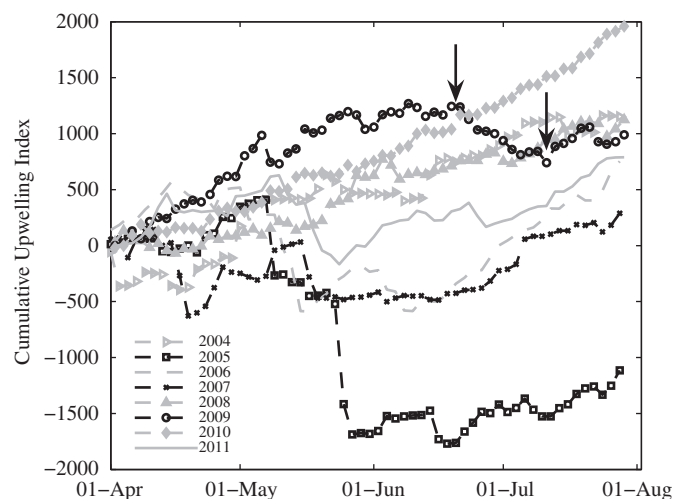

Fig. 10. Time-series of cumulative upwelling index (m³ (100 m coastline)$^{-1}$ d^{-1}) for 2004–2011 at NERACOOS buoy B (see Fig. 1 for buoy location). Vertical arrows bracket the time period of unusual downwelling during the summer of 2009.

2009 with a coast-wide onset of toxicity (Fig. 4) is suggestive of onshore transport of an offshore population with an alongshore extent spanning the entire Maine coast. Blooms of similar scale have been observed in the past (Anderson et al., 2005b; Townsend et al., 2001), and their regional scope is consistent with alongshore advection of coastal populations originating from cyst beds (Fig. 9) in the Bay of Fundy and offshore of mid-coast Maine (Anderson et al., 2005d; McGillicuddy et al., 2005). Leakiness of the Bay of Fundy gyre has been hypothesized to be a factor influencing the magnitude of blooms entering the Maine Coastal Current (Aretxabaleta et al., 2008; Aretxabaleta et al., 2009), but unfortunately there are no measurements to quantify the export of *A. fundyense* cells from the Bay of Fundy in 2009.

Although this event was apparently coast-wide in extent, its visual manifestation appears to have been confined to the western Gulf of Maine. Direct observations do not permit delineation of the spatial scale of the discolored water (Fig. 1), but satellite data reveal a pronounced surface expression extending along the coast from western Penobscot Bay to Cape Ann (Fig. 3). This area corresponds directly to the portion of the coastline in which toxicities were highest (Fig. 4).

Unfortunately, information about the vertical extent of the population in the discolored water is scant. The peak cell concentration observed at the PPS mooring site on July 11 at 5 m (Fig. 6) was two orders of magnitude smaller than surface samples collected by bucket from R/V Tioga on July 10 (Fig. 1) suggests that the population was confined to a thin near-surface layer. However, the non-simultaneity of those measurements constitutes an important caveat to this inference. Nevertheless, the presence of such high surface concentrations is indicative of upward swimming, and vertical swimming speeds for *A. fundyense* and similarly-sized dinoflagellates (diameter $\sim 40\,\mu$) are typically between 5 and 15 m day^{-1} (Anderson and Stolzenbach, 1985; Bauerfeind et al., 1986; Fauchot et al., 2005; Kamykowski et al., 1992). Upward swimming in the presence of near-coastal convergence created by downwelling favorable wind-forcing (Fig. 10) would tend to accumulate *A. fundyense* biomass, as has been demonstrated in a variety of frontal systems (Franks, 1992, 1997).

A potential underlying cause of the apparent surface-seeking behavior could have been the need to aggregate the population to concentrations sufficient to make sexual reproduction practical (Wyatt and Jenkinson, 1997). Indeed, the predominance of 2c DNA content cells in the samples from the discolored water (Fig. 7) is consistent with a conversion to the sexual phase of the *A. fundyense* life cycle. However, this is not the only explanation for the high proportion of 2c cells. Because *A. fundyense* are haplontic (dividing only during its haploid, vegetative phase), *A. fundyense* may have 2c DNA content either during the G2 phase of mitosis or as newly-formed planozygotes (Brosnahan et al., 2014). In 5 of the 6 samples from the discolored water, more than 90% of *A. fundyense* cells were 2c, a ratio that strongly suggests that at least some of the cells had become planozygotes. The alternative – that the population consisted only of vegetative cells – is highly improbable because of the preponderance of 2c cells observed. Furthermore, all of the samples were collected between 1420 and 1702 local time, when most vegetative *A. fundyense* are in the 1c G1 phase of the cell division cycle rather than the 2c G2 phase. If it were assumed that all cells were vegetative and undergoing phased division, the minimum growth rate μ can be calculated as $\mu = 1/t \ln(1 + f_{max})$, where t is time and f_{max} is the fraction of the population with 2c DNA content (Chisholm, 1981). The implied minimum growth rate of 0.64 day^{-1} is at the upper limit of this organism's capability (Stock et al. 2005 and references therein), making that an unlikely explanation. Moreover, phased division by *A. fundyense* more typically occurs at daybreak, not during the late afternoon (Rubin, 1981). For all these reasons, it is much more likely that a substantial proportion of the 2c cells in these samples were planozygotes.

The unusually large flux of *A. fundyense* cysts observed in Wilkinson Basin (Fig. 8) further attests to the association of the discolored water with a massive sexual reproduction event. As indicated by the drifter trajectory (Fig. 1), the transit time between the area of discolored water and the sediment trap is approximately ten days. Given sinking rates of *A. fundyense* cysts on the order of 10 m day^{-1} (Anderson et al., 1985), arrival at the 95 m sediment trap along this advective pathway is certainly plausible. Although the cyst flux of 75,236 cysts m^{-2} d^{-1} observed during July 9–19 at 95 m was the highest of all of the mid-depth deployments described in Pilskaln et al. (2014), the total flux measured during this ten-day interval constitutes less than 10% of what accumulated in the top 1 cm of bottom sediments at this location between October 2008 and October 2009 (Fig. 9). Low-level cyst fluxes throughout the rest of the year are not nearly able to make up the difference. This implies that the vast majority of the cysts deposited on the bottom in that location passed through the 95 m depth horizon elsewhere, perhaps further upstream—and that the cysts passing through 95 m at this location were deposited

further downstream in an area where there was a more modest (ca. 75 cysts cm^{-2}) increase in cyst abundance. The peak flux in the 180 m trap was four times higher than at 95 m, but it is not possible to partition that increase between lateral inputs and resuspension in the benthic nepheloid layer (see Pilskaln et al., 2014). In any case, the dramatic accumulation of ca. 1000 cysts cm^{-2} east of Cape Ann is consistent with production of cysts from an overlying population of 100,000 cells l^{-1} spread over a 1 m-thick layer, assuming 20% success in sexual reproduction (two cells fusing to make one cyst) followed by 100% deposition in the upper 1 cm of sediment.

From an historical perspective, it is interesting to note that the 1972 event was also associated with cyst formation. Although the cyst stage of *A. fundyense* was not fully described until later that decade (Anderson and Wall, 1978; Dale, 1977), Mulligan (1973) reported higher concentrations of cysts in water samples taken toward the end of the bloom. Subsequently, Mulligan (1975) suggested that the abundance of *A. fundyense* cysts in coastal sediments had increased as a result of the 1972 event, hypothesizing that this reservoir could seed blooms in future years. Thus, it appears that both the 1972 and 2009 red tides observed in the western Gulf of Maine involved sexual reproduction and major encystment events.

5. Conclusions

The 2009 *A. fundyense* red tide was extraordinary for a number of reasons. To our knowledge, this is only the second confirmed episode of discolored water in the western Gulf of Maine directly attributable to *A. fundyense*, the first being associated with the historic bloom of 1972 (Hartwell, 1975; Mulligan, 1973; Sasner et al., 1974). The event was clearly associated with sexual reproduction, yielding the highest cyst fluxes that have ever been measured in the region. A large deposit of cysts ensued in coastal sediments, temporarily extending the mid-coast Maine cyst bed farther south than previously observed.

The discolored water appears to have been the southern terminus of a coast-wide phenomenon, leading to widespread toxicity from the Bay of Fundy to western Maine. The event was preceded by a period of anomalous downwelling-favorable winds, which favor outflow from the Bay of Fundy gyre (Aretxabaleta et al., 2008; Aretxabaleta et al., 2009) where a large source population is typically located. Downwelling-favorable winds also tend to accelerate the alongshore current, thereby enhancing transport of *A. fundyense* along the coast into the western Gulf of Maine. A third potential impact of the downwelling-favorable wind arises from the associated convergence along the coast, which would tend to accumulate upward-swimming *A. fundyense*.

Although the precise mechanisms leading to the red-tide event are not known, one aspect is clear: it was an extremely-ephemeral phenomenon. A combination of satellite imagery and rapid-response shipboard surveys constrain the duration of water discoloration to a period of less than two weeks. Similarly, the 1972 red water event ended abruptly, with *A. fundyense* disappearing from the plankton within 5–7 days after peak bloom conditions (Sasner et al., 1974). The short duration of these episodes is particularly humbling from an observational perspective, insofar as the duration of the event is shorter than the typical interval in between "synoptic" surveys of bloom dynamics carried out in regional research programs. This begs the question of how many such events may have been missed in the past due to observing strategies not capable of resolving them. Fortunately, with the deployment of in situ monitoring devices capable of species-specific measurements (Scholin et al., 2009), future prospects are bright for observing these highly-transient processes that obviously play an important role in the regional population dynamics of *A. fundyense* and other harmful algae around the world.

Acknowledgments

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