OC460 Cruise Report Draft 5/11/10

Voyage #460 of R/V *Oceanus* was the first of four cruises in 2010 organized to serve complementary scientific objectives of two different projects. The two projects are:

GOMTOX: Dynamics of *Alexandrium fundyense* distributions in the Gulf of Maine: an observational and modeling study of nearshore and offshore shellfish toxicity, vertical toxin flux, and bloom dynamics in a complex shelf sea – NOAA ECOHAB

Objectives:

Investigate *A. fundyense* bloom dynamics and the pathways that link this organism to toxicity in nearshore and offshore shellfish.

Investigate the vertical structure of *A. fundyense* blooms, vertical toxin flux, and linkage to toxicity in offshore shellfish.

Alexandrium population biology in the Gulf of Maine – Woods Hole Center for Oceans and Human Health – NSF/NIEHS¹

Objectives:

Sample genetic variability of *Alexandrium* subpopulations throughout the Gulf of Maine.

Measure changes in relative abundance of *Alexandrium* genotypes in space and time.

Objectives common to both projects include:

Assess hydrodynamic and hydrographic context for interpretation of *Alexandrium* spp. measurements.

Incorporate field observations into a suite of numerical models for hindcasting and forecasting applications.

The primary domain of interest is Georges Bank, where a large bloom of *A. fundyense* was observed in 2007 and shorter and less intense bloom occurred in 2008. The four cruises in 2010 are designed to (1) resolve the seasonal variation of the Georges Bank bloom, and (2) quantify its interannual variability.

A secondary objective was added to the 2010 cruises when the results of the fall 2009 cyst survey (OC440) revealed that cyst abundance offshore of mid-coast Maine is now higher than in all prior measurements, including those that preceded the severe blooms of 2005 and 2008. This field season thus offers an exceptional opportunity for testing the hypothesis that the magnitude

¹ <u>http://www.whoi.edu/science/cohh/whcohh/projects/habs1_abstract.htm</u>

of the bloom in the western Gulf of Maine and Southern New England is set by the abundance of cysts. We therefore must consider the possibility of redirecting some of this year's observational effort from Georges Bank to the Gulf of Maine. These choices will be informed by a number of factors, including real-time nowcasting and forecasting activities², as well as state agency toxicity monitoring efforts along the coasts of Maine, New Hampshire, and Massachusetts. If widespread toxicity appears along the coast, that would be consistent with the cyst hypothesis. However, if widespread toxicity does not appear, that would not necessarily be inconsistent with the hypothesis, as a large bloom could be present offshore. It is in this latter circumstance that diverting to the western Gulf of Maine would be most advantageous for hypothesis testing, insofar as confirming the absence of a large bloom would provide evidence for rejecting the hypothesis.

OC460 began with a survey of Georges Bank (Figure 1). Surface live counts indicate very low *Alexandrium* concentrations over the entire bank, with most samples coming up zero. The only systematic pattern in the data is on the northern edge of the bank, where concentrations are consistently low, although they are at the limit of detection.

It is interesting to compare this survey with a prior one at the same time of year, in which cell concentrations were significantly higher (Figure 2; OC445, April 28 – May 5, 2008). Surface live counts on OC445 indicated cell concentrations ranging from zero to several hundred cells per liter, with highest concentrations on the northwest part of the bank. Tens to hundreds of cells per liter extend through the western half of the crest to the southern flank.

Water mass analysis suggests interannual variability in hydrographic properties. Temperaturesalinity diagrams reveal both Georges Bank water and warm/salty water characteristic of the continental slope (Figure 3). Focusing on the Georges Bank water (4-8°C, 31.5-33.5 psu), it appears to be nearly 2 degrees warmer and perhaps 0.5 psu fresher in 2010 than 2008. The cause of this interannual variability is unknown at this time.

The OC460 coastal survey consisted of a series of transects spanning the area from Cape Cod Bay to Bar Harbor Maine (Figure 2). Surface live counts revealed low *Alexandrium* abundance south and west of Penobscot Bay, with cell concentrations at or below the limit of detection in that area. From Penobscot Bay to the east, cell concentrations were patchy, but higher overall—with values ranging from zero to 578 cells 1^{-1} .

Just as on Georges Bank, water mass properties in the Gulf of Maine show significant interannual variability. Slope waters in the deep basins are more than a degree warmer in comparison with this same time period in 2008 (Figure 3). Maine intermediate waters are a few tenths of a degree warmer and a few tenths of a psu fresher than in 2008. However, these results must be treated with caution as the salinities have not yet been calibrated with salt bottle data yet.

These findings are in stark contrast with those from 2007, where the western Gulf of Maine was virtually devoid of *Alexandrium* cells in mid-to-late May (even later in the bloom season). Live counts from EN437 (May 17-31 2007) were almost all zeros, with *Alexandrium* detected at only

² <u>http://omglnx3.meas.ncsu.edu/GOMTOX/2010forecast/</u>

a few isolated stations. The peak concentration of 84 cells l⁻¹ was observed at the innermost station of the Saco Bay transect.

Observations in this area during the late April / early May time period during the ECOHAB-GOM era range from zero to approximately 200 cells l^{-1} (Table 1). In comparing the live counts from the present cruise with those published observations, one must bear in mind the methodological differences: the former are not able to distinguish *A. fundyense* from other species that are morphologically similar, whereas the latter are more species specific. Nevertheless, the initial results of OC460 suggest higher than average cell concentrations in the western Gulf at this very early stage of the bloom season. Our findings are thus not inconsistent with the hypothesis that the 2010 bloom will be anomalously large. However, we note that the observed cell concentrations are significantly lower than those predicted by the forecast model (Figure 4).

A total of 3 surface drifters were deployed at the inshore stations of the Casco Bay transect (Figure 5; Appendix A, Table 3)³. Trajectories of all three indicate a significant offshore velocity component, with only a modest alongshore velocity component.

Date	Max surface	Location	Reference			
	concentration					
	(cells l^{-1})					
21-23 April 1998	200	Inshore Casco Bay	Keafer et al. 2005a			
28-29 April 2000	50	Offshore Casco Bay	Keafer et al. 2005a			
April/May 2000	200	Inshore Casco Bay	Townsend et al. 2005			
3-4 May 2000	200	Offshore Casco Bay	Keafer et al. 2005a			
6-11 May 2001	0	N/A	Keafer et al. 2005b			
Table 1. Maximum A. fundyense cell concentrations found in western Gulf of Maine waters						
(Casco Bay and westward) in late April / early May.						

³ Also see <u>http://nefsc.noaa.gov/drifter</u>







2008 (blue) and OC460 in 2010 (red). Left: Georges Bank; right: Gulf of Maine.





Appendix A: Measurements made on OC460

Underway measurements

a. Acoustic Doppler Current Profiler

b. Meteorological sensors

Core hydrographic measurements

a. CTD (pressure, temperature, salinity, oxygen, fluorescence, beam attenuation, PAR)

b. Alexandrium cell counts: 1, 10, 20, 30, 40, 50m plus 250/near bottom

c. Nutrients: standard depths plus 100, 150, 200, 250m

Water budget:

Bottle #	Depth	Live	Spare	Whole Cell	SHA	Nuts/Chl	Pseuds	total
1	1			2	2	1.0	1.0	7.0
2	1	10						10
3	1		10					10
4	10			2	2	1.0	1.0*	7.0
5	20			2	2	1.0	1.0*	7.0
6	30			2	2	1.0	1.0*	7.0
7	40			2	2	1.0	1.0*	7.0
8	50			2	2	1.0	1.0*	7.0
9	100					1.0		1.0
10	150					1.0		1.0
11	200					1.0		1.0
12	250 /					1.0		1.0
	near							
	bottom							

Af water- For Whole Cell (WC) and Sandwich Hybridization (SHA) - 4 liters collected total and 20 μ m sieved and split between the two assays.

4L/depth combined/split x 6 depths=6 WC tubes&6 filters/station (6 hole-manifold #1 loaded once)

Pseuds – At each station: 1) 125 ml whole water will be filtered for *Pseud* SHA onto 0.45 μ m Duropore filters; 2) 125ml whole water will be filtered for ARISA samples onto 0.45 μ m Isopore HA filters (as in 2008); 3) 125 ml whole water will be filtered for Domoic Acid onto 0.45 μ m Isopore HA filters. SHA filters will be frozen in LN2 Dewar

3 filters and cryo-vials/station will be needed. Use a 3-hole manifold—Note that the Pseud SHA filter and the ARISA/DA filters are the same pore size, but not the same material. Do not mix up the filter types.

*A vertical profile of *Pseuds* will be sampled at 4-6 selected stations with high abundance, in different hydrographic regimes as conditions permit. Same procedure as above but repeat for all 6 std depths.

A Domoic Acid "calibration" station will also be done at selected stations-details TBD.

Opportunistic samples– a spare 10L live sample will be available for multiple purposes; e.g., culturing of Pseuds and/or *Alexandrium*, life cycle stage samples, and possibly microsatellite analysis of *Alexandrium* populations. Additional opportunistic samples may be taken in areas of high *Alexandrium* and/or *Pseud* abundance.

Toxin size fractionation – Turner

Pump profiles were carried out at selected locations. Sampling depths were chosen to coincide as closely as possible with hydrographic sampling and sediment trap measurements. Pump deployments are summarized in Table 2.

	Date	Time	Latitude	Longitude	Station	Live Count	
		(local)					
1	May 1		41 34.3 N	68 23.0 W	FDA shellfish time-series site		
					Cultivator Shoal, CTD 55		
2	May 9		42 17.6 N	70 39.3 W	Near Minots Light, Boston,		
					CTD 123		
Table 2. Pump stations.							

Drifters

ID	Mon	Dav	Year	Local Time	Lon	Lat	Drogue depth(m)	Station Number
105430691	5	7	2010	0612	69 52.050W	43 39.750N	1	CB1B
105430692	5	7	2010	0703	69 48.642W	43 33.720N	1	CB1C
105430693	5	7	2010	0750	69 45.102W	43 27.738N	1	CB1D
Table 3: Summary of drifter releases on Casco Bay line. For more information see								
http://nefsc.noaa.gov/drifter.								

Microbial community structure and bacterial abundance – Amaral-Zettler and Murphy

OC460 #1 - CTD55 - Georges Bank - Alexandrium live count = 0 5/5/10 41 34.3 N / 68 23.0 W Sfc (1),(2) 10m (1),(2) 20m (2) [replicate 1 lost] OC460 #2 - CTD79 - Casco Bay - Alexandrium live count = 05/7/10 43 39.8 N / 69 52.0 W [replicate 1 lost] Sfc (2) 10m (1),(2) 20m (1),(2) OC460 #3 - CTD106 - Isle au Haut - Alexandrium live count = 05/8/10 43 53.6 N / 68 25.1 W Sfc(1),(2)10m(1)[replicate 2 lost] 20m(1),(2)OC460 #4 - CTD123 - Minots Light - Alexandrium live count = 5/9/10 42 17.6 N / 70 39.3 W Sfc (1),(2)10m (1),(2) [no filter left for replicate sample] 20m(1)





















Appendix C: Vertical sections.

































































Appendix D: Satellite imagery











Appendix E: Personnel

McGillicuddy	Dennis	WHOI
Keafer	Bruce	WHOI
Payette	Jack	
Conroy	Brandon*	NEU
Adams	Jillian*	NEU
Stone	Mollie*	NEU
Smith	Keston	WHOI
Rebuck	Nathan*	UMe
Thomas	Maura	UMe
Kosnyrev	Olga	WHOI
Turner	Jeff	UMassD
Petitpas	Chrissy*	UMassD
Milligan	Peter*	UMassD
Emde	Grant	NMFS
Stessel	Robert*	UMe
Wilson	Emily*	Memorial University
Zhang	Gordon*	WHOI

*Student/postdoc

Watch schedule

Watch number	1	2	3
4 on / 8 off	8-12	12-4	4-8
1. CTD Operator	Gordon	Keston	Robert
2. Cell Counter	Bruce*	Brandon*	Chrissy*
3. Nutrient sampler	Olga#	Nathan#	Maura#
4. Water sampler	Jack#	Grant#	Jeff
5. Water sampler	Mollie	Jillian	Peter#

* Wetlab chief

CTD slip line handlers