

1 **Vertical distribution of buoyant *Microcystis* blooms in a**
2 **Lagrangian particle tracking model for short-term forecasts**
3 **in Lake Erie**

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17 **Key Points**

18 *Microcystis* vertical distribution is a dynamic balance between turbulence and buoyancy

19 Appropriate time step and numerical scheme avoid artifacts in random walk models

20 Vertical mixing with buoyancy improved simulation of bloom spatial distribution

21 **Abstract**

22 Cyanobacterial harmful algal blooms (CHABs) are a problem in western Lake Erie, and in
23 eutrophic fresh waters worldwide. Western Lake Erie is a large (3000 km²), shallow (8 m mean
24 depth), freshwater system. CHABs occur from July to October, when stratification is intermittent
25 in response to wind and surface heating or cooling (polymictic). Existing forecast models give
26 the present location and extent of CHABs from satellite imagery, then predict two-dimensional
27 (surface) CHAB movement in response to meteorology. In this study, we simulated vertical
28 distribution of buoyant *Microcystis* colonies, and 3D advection, using a Lagrangian particle
29 model forced by currents and turbulent diffusivity from the Finite Volume Community Ocean
30 Model (FVCOM). We estimated the frequency distribution of *Microcystis* colony buoyant
31 velocity from measured size distributions and buoyant velocities. We evaluated several random-
32 walk numerical schemes to efficiently minimize particle accumulation artifacts. We selected the
33 Milstein scheme, with linear interpolation of the diffusivity profile in place of cubic splines, and
34 varied the time step at each particle and step based on the curvature of the local diffusivity profile
35 to ensure that the Visser time step criterion was satisfied. Inclusion of vertical mixing with
36 buoyancy significantly improved model skill statistics compared to an advection-only model, and
37 showed greater skill than a persistence forecast through simulation day 6, in a series of 26
38 hindcast simulations from 2011. The simulations and in-situ observations show the importance of

39 subtle thermal structure, typical of a polymictic lake, along with buoyancy in determining vertical
40 and horizontal distribution of *Microcystis*.

41 **Index Terms**

42 4239 Limnology, 4842 Modeling, 4855 Plankton

43 **Keywords**

44 Numerical modeling, Ecological forecasting, Lake Erie, *Microcystis*, Harmful algal bloom,
45 cyanobacteria

46

47 **1 Introduction**

48 Harmful algal blooms (HABs) are a global problem that is linked to anthropogenic eutrophication
49 of inland and coastal waters, and may be exacerbated by climate change [O'Neil, J. et al., 2012].

50 Lake Erie has experienced recurring blooms of toxin-producing cyanobacteria since the mid

51 1990s [Brittain, S. M. et al., 2000; Michalak, A. M. et al., 2013; Wynne, T. T. and R. P. Stumpf,

52 2015]. Lake Erie is the most productive, warm, and shallow of the Laurentian Great Lakes of

53 North America. In the open waters of Lake Erie, cyanobacterial harmful algal blooms (CHABs)

54 generally occur from July to October, and are dominated by the species *Microcystis aeruginosa*,

55 which produces the group of hepatotoxin compounds known as microcystins [Rinta-Kanto, J. M.

56 et al., 2009]. CHABs occur primarily in the shallow western basin, which receives the main

57 hydraulic load from the Detroit River in the north and the main nutrient load from the Maumee

58 River [Kane, D. D. et al., 2014] in the southwest (Fig. 1a). Occasionally, CHABs are transported

59 through the islands into the deeper central basin, while the eastern basin is largely free of CHABs
60 (Fig. 1a)[*Wynne, T. T. and R. P. Stumpf, 2015*]. A bloom of record-breaking intensity and extent
61 occurred in 2011. Analysis by Michalak et al. [2013] indicated that the conditions of meteorology
62 and agricultural land use that caused the 2011 record bloom are likely to recur, and the 2015
63 bloom subsequently exceeded the severity of the 2011 record bloom [*Stumpf, R. P. and T. T.*
64 *Wynne, 2015*].

65 Forecasts of CHAB abundance and spatial distribution are useful to water treatment plant
66 operators, anglers, recreational boaters, and beach users. Lake Erie is a source of drinking water
67 to 11 million people [*French, R. et al., 2011*]. In 2014, the City of Toledo issued a do-not-drink
68 order that affected a half million residents for two days as a result of contamination of treated
69 water by microcystins [*Henry, T., 2014*]. In addition to spatial forecasts, forecasts of *Microcystis*
70 vertical distribution are of interest to water treatment plant operators because intake structures are
71 usually located sub-surface, so the risk of toxins in their raw water may be greater during mixing
72 events than when *Microcystis* colonies are concentrated on the surface. In addition to providing
73 drinking water, Lake Erie supports economically valuable sport and commercial fisheries as well
74 as recreation and tourism.

75 Short-term and seasonal CHAB forecasts have been developed for Lake Erie. Seasonal forecasts
76 predict the annual bloom severity using statistical models based on the cumulative March to July
77 phosphorus load from the Maumee River, which are able to explain > 90% of the interannual
78 variance in bloom severity [*Obenour, D. R. et al., 2014*]. Bloom severity is defined as the lake-
79 wide cyanobacterial biomass averaged over the 30 days containing the maximum biomass

80 [Obenour, D. R. et al., 2014; Stumpf, R. P. et al., 2012], and has been estimated using both
81 satellite [Stumpf, R. P. et al., 2012] and plankton tow data [Bridgeman, T. B. et al., 2013]. The
82 seasonal forecast is used by water treatment plant managers for seasonal planning, to determine
83 recommended phosphorus load reductions to meet commitments under the Great Lakes Water
84 Quality Agreement [GLWQA Annex 4, 2015], and is distributed to over 1600 subscribers. Short-
85 term forecasts are distinct from the seasonal forecast in the greater importance of physical
86 transport processes over biological mechanisms in explaining short-term bloom variability.
87 Experimental short-term forecasts have been developed for Lake Erie that indicate the present
88 location and extent of CHABs from satellite imagery, then predict future movement of the CHAB
89 using forecast meteorology, a hydrodynamic model, and a Lagrangian particle tracking model to
90 simulate horizontal advection of neutrally-buoyant particles at the water surface [Wynne, T. T. et
91 al., 2013; Wynne, T. T. et al., 2011].

92 Skill assessment of short-term CHAB forecasts is needed so that forecast users may have an
93 appropriate level of confidence in forecast data, and for development of improved models.
94 However, CHAB forecast skill assessment can be challenging, and methods are not well
95 established. Wynne et al. [2011] showed that a forecast model more accurately predicted
96 horizontal movement of the bloom centroid (center of mass) than a persistence forecast over a
97 ten-day simulation for an August 2003 event. A persistence forecast is a benchmark used for
98 model skill assessment in which no change is assumed from the initial observed location. The
99 same quantitative skill assessment method could not be applied in the case of an August 2008
100 event because cloud cover and vertical mixing obscured parts of the bloom in subsequent images

101 and prevented an accurate estimate of the centroid location [Wynne, T. T. et al., 2011]. In a
102 detailed analysis of the August 2008 bloom in Lake Erie, Wynne et al. [2010] found that wind
103 speed was a significant predictor of apparent bloom intensity in satellite imagery, and presented
104 evidence to support the hypothesis that variation in the mixing depth of the buoyant
105 cyanobacterial colonies was the underlying mechanism causing changes in surface concentration.
106 In a qualitative analysis of the first three years of the experimental Lake Erie CHAB forecast
107 (2008-2010), Wynne et al. [2013] concluded that the forecast provided useful information, but
108 could be improved by a means to fill in cloud-covered areas in the satellite images, and by
109 simulation of vertical mixing of the buoyant cyanobacteria.

110 To simulate concentration profiles of buoyant particles produced by vertical mixing in stratified
111 turbulence, the partial differential equations governing advection and diffusion may be solved
112 from an Eulerian or a Lagrangian perspective; each approach has strengths and weaknesses. Real
113 phytoplankton communities have properties that vary among individuals within a population,
114 such as size, specific gravity, light exposure history, and nutrient quotas. A strength of
115 Lagrangian particle models is that nearly continuous distributions of properties can be
116 represented by allowing properties to vary by particle [e.g., Ross, O. N. and J. Sharples, 2008],
117 while in Eulerian models property distributions are usually represented by a limited number of
118 discrete property classes [e.g., Medrano, E. A. et al., 2013].

119 A weakness of the Lagrangian approach is that artificial accumulation of particles can occur in
120 low diffusivity areas in random-walk turbulence simulations for the case of spatially (vertically)
121 non-uniform diffusivity if an inappropriate numerical scheme or time step is used [Visser, A.,

122 1997]. These artifacts can be easy to misinterpret as features of interest. Several studies have
123 applied Lagrangian particle models to the 1D column case with steady, idealized diffusivity
124 profiles [Gräwe, U. et al., 2012; Ross, O. N. and J. Sharples, 2008; Visser, A., 1997]. In 3D
125 ocean and lake models, use of random-walk vertical mixing schemes can be challenging because
126 the required time step varies spatially and temporally over the model domain. A fixed time step
127 may be selected that is adequate in most places and times [Huret, M. et al., 2007], or in some
128 cases a well-mixed profile was imposed in shallow regions where a small time step would be
129 required to avoid formation of artifacts [Gilbert, C. et al., 2010]. Our application required
130 realistic rather than idealized diffusivity profiles for a polymictic lake in which conditions are
131 alternately turbulent and stratified, and to simulate concentration profiles rather than specify a
132 well-mixed condition, even in shallow areas, therefore some development of the random-walk
133 approach was required.

134 We describe a short-term forecast system for CHAB abundance and distribution in Lake Erie that
135 takes a similar approach to that of Wynne et al. [2011], but uses updated hydrodynamic and
136 Lagrangian particle tracking models, and includes a means of filling in cloud-covered areas of
137 satellite images using model data from a previous run. In addition, we describe a method to
138 simulate vertical distribution of buoyant cyanobacteria in stratified turbulence. We evaluated the
139 performance of several random-walk turbulence numerical schemes in terms of computational
140 efficiency and their ability to minimize artifacts in simulations with vertically-varying diffusivity
141 typical of a large polymictic lake. We compared simulated vertical distributions of cyanobacteria
142 to observed profiles in Lake Erie. Finally, we show that model skill statistics were improved by

143 including vertical mixing with buoyancy in hindcast simulations of the spatial distribution of the
144 2011 bloom in Lake Erie.

145

146 **2 Methods**

147 **2.1 Hydrodynamic model**

148 The Finite Volume Community Ocean Model (FVCOM, v. 3.2) is an unstructured grid, finite-
149 volume, free surface, three-dimensional primitive equation ocean model that solves the
150 momentum, continuity, temperature, salinity, and density equations [Chen, C. *et al.*, 2003].
151 Turbulence closure is implemented through the MY-2.5 scheme for vertical mixing [Galperin, B.
152 *et al.*, 1988], and the Smagorinsky scheme for horizontal mixing [Smagorinsky, J., 1963].
153 FVCOM has been adapted and implemented for the Great Lakes in several recent studies
154 [Anderson, E. J. *et al.*, 2015; Anderson, E. J. and D. J. Schwab, 2013; Anderson, E. J. *et al.*,
155 2010; Bai, X. *et al.*, 2013], yielding accurate predictions of temperature, water levels, and
156 currents. In particular, FVCOM has been applied to Lake Erie for extreme storm prediction
157 [Anderson, E. J. *et al.*, 2015] and serves as the oceanographic model underlying NOAA's next-
158 generation Lake Erie Operational Forecast System (LEOFS), a real-time short-term
159 hydrodynamic forecast model (<http://tidesandcurrents.noaa.gov/ofs/leofs/leofs.html>).

160 The FVCOM-based LEOFS model was applied for this study with bathymetry interpolated from
161 the NOAA National Geophysical Data Center.

162 (www.ngdc.noaa.gov/mgg/greatlakes/greatlakes.html). The unstructured grid consisted of 6,106

163 nodes and 11,509 elements (Fig. 1). Spatial resolution was 2 km in the central basin, 1.5 km in
164 the western basin and 0.5 km in Maumee Bay and the islands (Fig. 1b) to improve representation
165 of currents in these key areas for CHAB transport, with 20 uniform vertical sigma layers.

166 Dynamic water levels (6-minute) were prescribed at open boundaries at the Detroit River and the
167 Niagara River, taken from adjustments to the NOAA/NOS gauges at Gibraltar (9044020) and
168 Buffalo (9063020), to drive the primary inflow and outflow, respectively (Fig. 1a). Atmospheric
169 forcing conditions were generated using station-based interpolation methods as in the NOAA
170 Great Lakes Coastal Forecasting System (Beletsky et al., 2003; Schwab and Beletsky, 1998).

171 Hourly meteorological forcing variables of wind speed, wind direction, air temperature, dewpoint
172 temperature, and cloud cover were interpolated over Lake Erie from several land-based
173 meteorological stations and offshore NOAA/NDBC buoys (45004, 45132, and 45142), when
174 available. Wind speeds were adjusted to 10-m height and empirical relationships were used to
175 adjust land-based meteorological variables for over-lake modification (Beletsky et al., 2003;
176 Schwab and Beletsky, 1998).

177 Hydrodynamic model simulations were based on the real-time LEOFS nowcast, which was
178 initialized on January 1, 2004 with uniform temperature of 2 °C. For the 2011 scenario presented
179 here, the hydrodynamic model simulation was initialized on January 1, 2011 with initial
180 conditions provided by the simulation from the previous year, and produced hourly output of
181 three-dimensional currents, water temperature, turbulent diffusivity, and 2D water level
182 fluctuations.

183 **2.2 Lagrangian particle tracking model**

184 Lagrangian particle tracking was accomplished using a Fortran program developed previously to
185 study transport of larval cod in the Gulf of Maine [*Churchill, J. H. et al., 2011; Huret, M. et al.,*
186 2007], which is distributed as part of the FVCOM code package
187 (<http://fvcom.smast.umassd.edu/>), and has previously been applied in the Great Lakes [*Anderson,*
188 *E. J. and M. S. Phanikumar, 2011*]. Advection of particles was determined by

$$\frac{d}{dt}\mathbf{X}(t) = \mathbf{V}(\mathbf{X}(t), t) \quad (1)$$

189 where $\mathbf{X}(t)$ is the three-dimensional particle position at time t , and $\mathbf{V}(\mathbf{X}(t), t)$ is the three-
190 dimensional, time varying velocity field. Linear interpolation in space and time was used to
191 obtain $\mathbf{V}(\mathbf{X}(t), t)$ from hourly-archived FVCOM output. The contribution of advection to the
192 particle position was updated by integrating Eq. (1) using an explicit fourth-order Runge-Kutta
193 scheme with a time step, $\Delta t = 600\text{s}$. Vertical mixing due to turbulent eddy diffusivity was
194 optionally simulated using one of several random-walk schemes, described below. We used
195 reflected boundary conditions at vertical and horizontal boundaries in all simulations.

196 **2.3 Well-mixed condition test**

197 We compared the performance of several random walk vertical mixing numerical schemes using
198 a well-mixed condition test case [*Visser, A., 1997*] that can be used to evaluate whether a given
199 numerical scheme and time step will produce artifacts in simulated concentration profiles. In our
200 case, the well-mixed condition simulation was performed by initiating 1000 particles, uniformly
201 distributed through the water column, then simulating 1D vertical mixing with neutral buoyancy

202 using time series of Lake Erie diffusivity profiles output from FVCOM for the month of August,
203 2011 (Table 1). Random noise in the simulated concentration profiles decreases with increasing
204 number of particles, and 1000 particles were found to be adequate in 1D simulations [Ross, O. N.
205 and J. Sharples, 2004]. These simulations are expected to give uniform concentration at all time
206 regardless of the diffusivity profile, consistent with the Eulerian solution to the 1D diffusion
207 equation with initial uniform concentration [Visser, A., 1997].

208 To evaluate performance in the well-mixed condition test in a way that is directly relevant to our
209 application, we defined a signal-to-noise ratio, SN , based on simulated surface concentration,

$$SN(t) = C(t)/|1.0 - C_{wm}(t)| \quad (2)$$

210 where $C_{wm}(t)$ is the time-dependent concentration in the 1-m thick surface layer, normalized to
211 the column-mean concentration, in a well-mixed condition simulation with neutral buoyancy. A
212 constant value of 1.0 is expected for $C_{wm}(t)$, therefore $|1.0 - C_{wm}(t)|$ represents the magnitude
213 of “noise” due to artifacts and random fluctuations due to calculating concentration by counting
214 discrete particles in a control volume. The “signal” is provided by the analogous surface
215 concentration, $C(t)$, from an identical simulation *with* buoyancy. A large value of SN indicates
216 that artifacts are small compared to the effect of interest, which is the fluctuation in surface
217 concentration due to buoyancy. SN can be made arbitrarily large by using a small time step and a
218 large number of particles, but at the expense of computational time. We selected a value of 5 as a
219 goal minimum value for SN , based on the “Rose criterion” for the detection limit of the human
220 eye for image features [Rose, A., 1948]. Accordingly, we used the frequency of occurrence of SN

221 < 5 over the hourly records in a simulation as the performance criterion by which to evaluate the
222 numerical schemes and time step criteria.

223 **2.4 Random walk vertical mixing schemes**

224 We evaluated the Visser [1997] scheme, as-implemented by Huret et al. [2007]

$$z(t + \delta t) = z(t) + w_b \delta t + K'(z(t)) \delta t + R \sqrt{\frac{2K(\tilde{z}) \delta t}{\sigma^2}} \quad (3)$$

225 where $z(t)$ is the vertical position of the particle at time t , δt is the vertical random walk time
226 step, w_b is a floating/sinking/swimming vertical velocity component, K is the vertical turbulent
227 diffusivity, $K' = dK/dz$, R is a random variable sampled from a uniform distribution with zero
228 mean and standard deviation σ , and $\tilde{z} = z(t) + 0.5K'(z(t))\delta t$ is a vertical position displaced
229 from the particle position as a function of the diffusivity gradient. Following Ross and Sharples
230 [2004], a cubic spline interpolation was used to obtain a continuous, differentiable diffusivity
231 profile.

232 In addition to the Visser scheme, described above, we evaluated random walk vertical mixing
233 schemes with higher order accuracy including the Milstein, Strong 1.5 (S1.5), and Platen two-
234 step (PC2) schemes that were implemented in Fortran for the General Ocean Turbulence Model
235 by Gräwe [2011]. After evaluating the performance of ten random walk schemes [Gräwe, U.,
236 2011], Grawe et al. [2012] recommended the use of either the Milstein scheme or higher-order
237 schemes such as S1.5 or PC2. The order of accuracy (rate of convergence) of numerical
238 approximations to stochastic differential equations is separated into weak and strong cases, where

239 the weak case relates to the accuracy of the ensemble particle distribution, while the strong case
240 relates to the accuracy of particle trajectories [Gräwe, U., 2011]. The weak order of accuracy of
241 the schemes that we evaluated was 1 for Visser and Milstein, and 2 for S1.5 and PC2. The strong
242 order of accuracy was 1 for Milstein, 1.5 for S1.5, and was not defined for Visser or PC2 [Gräwe,
243 U., 2011]. The Milstein scheme is

$$z(t + \delta t) = z(t) + w_b \delta t + 0.5K'(z(t))[\Delta W^2 + \delta t] + \Delta W \sqrt{2K(z(t))} \quad (4)$$

244 where ΔW is a random variable drawn from a Gaussian distribution with zero mean and standard
245 deviation $\sqrt{\delta t}$. Because the Milstein scheme is first order, linear interpolation of the diffusivity
246 profile to obtain $K(z(t))$ and $K'(z(t))$ are sufficient, and the added computational expense
247 compared to the Visser scheme (Eq. 3) is minimal. The second order schemes S1.5 and PC2
248 retain additional terms from the stochastic Taylor expansion, including higher order derivatives
249 that require the application of cubic splines to the diffusivity profile, and S1.5 requires an
250 additional random variable, therefore the second order schemes have greater computational
251 expense. We refer to Gräwe [2011] for a full explanation of the S1.5 and PC2 schemes.

252 **2.5 Random walk time step**

253 Visser [1997] introduced a time step criterion, $\delta t \ll \min(1/|K''|)$, where K'' is the second
254 derivative of the diffusivity profile, so that the diffusivity profile is reasonably approximated by a
255 first-order Taylor series expansion over the range of particle displacement. Ross and Sharples
256 [2004] found that

$$\delta t = \frac{1}{100|K''|} \quad (5)$$

257 is acceptable in most applications. To ensure the use of an appropriate time step throughout a 3D
258 model domain, and to avoid the need to specify an appropriate δt in advance, we modified the
259 code of Huret et al. [2007] to allow an appropriate value of δt to be calculated and applied for
260 each particle at each δt . To evaluate Eq. (5) independently of spline interpolation, we calculated
261 K'' directly at the FVCOM sigma levels using a centered finite difference approximation, and
262 extended the profile beyond the surface and bottom by reflection. We further limited $0.01 \text{ s} \leq$
263 $\delta t \leq \Delta t$. We tracked the minimum δt during simulations to confirm that the lower limit of 0.01
264 seconds was rarely applied.

265 **2.6 Measurement of *Microcystis* colony size distribution**

266 We assigned buoyant velocity in the model based on measured size distributions and regressions
267 of buoyant velocity versus colony diameter. We measured *Microcystis* colony diameter of Lugol
268 preserved samples collected from western Lake Erie in the summers of 2012, 2013, and 2014. In
269 2012 and 2013 colony diameters were measured by microscopy (Table 1). In 2014 we used the
270 FlowCam (Fluid Imaging Technologies). The FlowCam captures images of individual colonies
271 and estimates their equivalent spherical diameter by image analysis. Wang et al. [2015] showed
272 that counts and colony diameters of *Microcystis* given by FlowCAM and microscopic image
273 analysis diameters were nearly identical. Colonies in Lake Erie are typically $> 50 \mu\text{m}$
274 [Vanderploeg, H. A. et al., 2001] and buoyant. Samples were preserved with 1% formalin upon
275 collection, immediately refrigerated, and analyzed within 24 hours. FlowCam analyses were

276 performed with a 2× objective and 1 × 3 mm field of view flow cell with samples diluted as per
277 manufacturer recommendations to avoid capturing more than one image per trigger event.
278 Fluorescent triggering mode was used to avoid imaging detrital material that might be confused
279 with *Microcystis*. Samples were diluted in 0.2 µm filtered algal culture media [e.g., *Vanderploeg,*
280 *H. A. et al.*, 2001] and injected into the FlowCAM with a 60-mL syringe, which was constantly
281 turned over so as to prevent the buoyant colonies from aggregating in the syringe. The image
282 analysis algorithm was calibrated to identify the colony outline including the mucilage.

283 **2.7 Measurement of *Microcystis* colony buoyant velocity**

284 *Microcystis* colony buoyant velocity was measured using microscopic videography [*Bundy, M.*
285 *H. et al.*, 1998; *Strickler, J. R.*, 1985], a method in principle similar to that of Nakamura et al.
286 [1993]. Surface water samples were collected on 15 and 21 July 2015 (Table 1) at station WE15
287 (Fig. 1b, -83.0, 41.6) during the early afternoon. Water was placed in 1-L glass bottles in an
288 incubator outdoors with a neutral density filter to cut light intensity to 50% of surface irradiance.
289 Measurements of colony velocities were made throughout the morning and afternoon of the next
290 day. Digital video clips were captured of individual colonies rising through a 2-cm × 2-cm cross
291 section × 30-cm tall glass cuvette filled with ambient lake water inside of a water jacket in a
292 temperature controlled environmental room maintained at the lake temperature. Image capture
293 and analysis software (Templo Motus, Vicon Motus, Contemplas, GmbH, Germany) was used to
294 measure the velocity of the rising colonies. Diameters of the colonies were determined from
295 image analysis of video images using Image-Pro software (Media Cybernetics, Rockville, MD).

296 Water samples were diluted as needed with 0.2- μm filtered lake water to avoid turbulence
297 induced by multiple rising colonies.

298 **2.8 Measured vertical profiles of temperature and cyanobacterial concentration**

299 Vertical profiles of temperature and cyanobacterial concentration (reported in μg chlorophyll a L^{-1})
300 were measured using the FluoroProbe (bbe Moldaenke, GmbH), which uses spectral
301 fluorometry to partition total chlorophyll into multiple phytoplankton classes on the basis of their
302 characteristic pigments (green algae, cyanobacteria, diatoms/dinoflagellates, cryptophytes), with
303 correction for possible interference by colored dissolved organic matter [Catherine, A. et al.,
304 2012; Kring, S. A. et al., 2014]. Standard factory calibration settings for representative algal
305 classes were used. Profiles were measured at 11 stations (Fig. 1) weekly from June through
306 September, 2015 (Table 1). Profiles were selected for model skill assessment in which the
307 cyanobacterial chlorophyll was greater than chlorophyll from other algal classes, which occurred
308 13 July to 28 September.

309 **2.9 Satellite remote sensing data**

310 To initialize model simulations, and for model skill assessment, we used a series of images of
311 cyanobacterial blooms in Lake Erie from July to October 2011 (Table 1) that were derived from
312 the Medium Resolution Imaging Spectrometer (MERIS) [Wynne, T. T. and R. P. Stumpf, 2015].
313 MERIS standard level 2 data sets (in units of sr^{-1}) were used with a spectral shape algorithm
314 based around 680 nm [Wynne, T. et al., 2008] to obtain the cyanobacterial index (CI). CI varies
315 linearly with biomass, with a value of 10^{-3} sr^{-1} corresponding to approximately $10^5 \text{ cells mL}^{-1}$

316 [Stumpf, R. P. *et al.*, 2012], which is the World Health Organization's threshold of significantly
317 increased risk for human health effects [Chorus, I. and J. Bartram, 1999]. For our analysis, we
318 converted CI to chlorophyll concentration using an empirical relationship derived from field
319 radiometry and grab sample extracted chlorophyll from eutrophic lakes in Florida (R^2 0.95, std.
320 error $7.7 \mu\text{g L}^{-1}$, range 16 to $115 \mu\text{g L}^{-1}$). The relationship was also verified for satellite-derived
321 CI, and gave a relative root-mean square error of 27% [Tomlinson, M. C. *et al.*, 2016].

$$\text{Chl} = 12570 \text{ CI} + 10 \quad (6)$$

322 A value of $\text{CI} = 10^{-3} \text{ sr}^{-1}$ is approximately equivalent to $23 \mu\text{g L}^{-1}$ chlorophyll, which we used as a
323 threshold to define the presence of a CHAB.

324 **2.10 Hindcast simulations**

325 Daily satellite images for the period July to October 2011 were evaluated, and 26 images were
326 selected that had > 50% cloud-free views of western Lake Erie. A ten-day model simulation was
327 initialized from each image by assigning surface chlorophyll concentration values to FVCOM
328 nodes by nearest neighbor interpolation. Concentration was converted to Lagrangian particles by
329 specifying a chlorophyll mass per particle ($10^{10} \mu\text{g Chl particle}^{-1}$) and placing the specified
330 number of particles within a control volume. The FVCOM node-centered tracer control elements
331 were used as control volumes (Fig. 1). Vertical layers were specified as constant-thickness (1 m)
332 z-layers.

333 Preliminary hindcast skill assessment indicated a need to censor satellite data within a buffer
334 region of shorelines due to frequent false positives in these areas, which was likely caused by

335 contamination of the water signal from relatively bright surrounding land or by bottom
336 reflectance. The buffer width was set to 1 km south of Stony Point (41.94 °Lat.) and east of
337 Catawba Island (-82.85 °Lon.) and to 1.5 km elsewhere. The buffer was not applied in Maumee
338 Bay because CHABs were often present [Wynne, T. T. and R. P. Stumpf, 2015], reducing the
339 likelihood of false positives. Buffered or missing data areas were assigned values by nearest
340 neighbor if pixels containing valid data were available within 2 km. If no valid pixels were
341 available within 2 km of a node one of two approaches was used: 1) it was assumed that no
342 CHAB was present ($Chl = 0$), or 2) model output was carried forward to fill in the no-data area if
343 a previous model run was available covering the time period.

344 Two types of simulations were run, 2D and 3D. In 2D simulations, the surface chlorophyll
345 concentration was applied to the surface 1 m, with $Chl = 0$ below, and random walk vertical
346 mixing was turned off. In 3D simulations, surface chlorophyll concentration was applied over the
347 surface mixed layer (SML) depth (see below), and random walk vertical mixing was simulated.
348 Both 2D and 3D simulations included 3D advection.

349 **2.11 Estimation of the surface mixed layer depth**

350 It was necessary to estimate the surface mixed layer (SML) depth for buoyant *Microcystis*
351 colonies for the purpose of initializing the vertical distribution of particles (chlorophyll
352 concentration) in 3D simulations from satellite-derived surface chlorophyll concentration. The
353 vertical distribution of buoyant particles in the water column depends on buoyancy in addition to
354 turbulent diffusivity (e.g., Fig. 2d,e); therefore, we used the Lagrangian particle model to estimate
355 the initial vertical distribution of *Microcystis* colonies rather than using diffusivity or temperature

356 profiles from FVCOM directly. Initial vertical profiles were simulated at a subset of FVCOM
357 nodes (stations) because a large number of particles is needed to obtain a well-resolved profile,
358 which would be computationally intensive to simulate at all 6,106 nodes. The SML depth was
359 estimated by running 1D column vertical random walk simulations at 11 station locations (Fig.
360 1b) that were selected to provide representative coverage of the western basin where CHABs are
361 most common, with additional stations added at representative deeper locations. 1D simulations
362 were initialized with 1000 particles uniformly distributed over the column 36 hours prior to the
363 initialization time of the 3D model (satellite image time) to allow the particle distribution
364 sufficient time to adapt to the varying diffusivity. Random walk vertical mixing was forced by
365 hourly diffusivity output from FVCOM. The SML depth for *Microcystis* colonies was estimated
366 as the depth at which the 1D concentration profile decreased to half the surface concentration,
367 and the satellite-derived surface concentration was applied from the surface to this depth; this
368 approach provides an unbiased estimate of the total column biomass for the case of a uniform
369 concentration profile or a profile that can be approximated by a linear decrease. SML depth was
370 interpolated spatially to the FVCOM nodes by the nearest neighbor method.

371 **2.12 Model skill statistics**

372 Comparison of model results to in-situ profile data was conducted using column-integrated
373 quantities to minimize the effect of noise in the profile data on the statistics. Turbulent diffusivity
374 is strongly influenced by the static stability of the water column, which can be quantified using
375 the potential energy anomaly, ϕ ,

$$\phi = \frac{1}{h} \int_0^h (\hat{\rho} - \rho)gzdz; \hat{\rho} = \frac{1}{h} \int_0^h \rho dz \quad (7)$$

376 where ρ is the local density, h is the water column depth, and g is acceleration due to gravity
 377 [Simpson, J. and D. Bowers, 1981; Wiles, P. et al., 2006]. Vertical distribution of concentration
 378 was characterized by calculating the center of mass of the normalized concentration profile, σ_m

$$\sigma_m = \frac{1}{C} \sum_{k=1}^{k_b} c_k \sigma_k; C = \sum_{k=1}^{k_b} c_k \quad (8)$$

379 where $\sigma = z/h$ is the normalized vertical coordinate, c is the concentration at layer k normalized
 380 to the column-mean concentration, $k_b = 20$ is the number of uniformly-spaced σ layers in the
 381 model grid. The observed concentration profile was averaged over the σ layers of the model grid
 382 for the purpose of comparison to the model profiles.

383 Skill assessment in hindcast simulations was conducted by comparing model results to remote
 384 sensing images that were within the model simulation period. Each hindcast simulation was
 385 initialized from a satellite image, and two to four subsequent images were typically available
 386 within the simulation period for skill assessment. Skill assessment was conducted using a binary
 387 categorical variable (CHAB, no CHAB), and pixel-by-pixel comparisons of model to remote
 388 sensing observations were conducted. FVCOM tracer control elements (Fig. 1) were used as the
 389 spatial segmentation (pixels).

390 Our approach to skill assessment statistics followed Hogan and Mason [2012]. Two statistics
 391 were calculated from the elements of the contingency table, which are the number of a , correctly

392 predicted events (hits); b , false events (false alarms); c , false negatives (misses); and d , correct
393 non-events. The frequency bias (B) gives the ratio of the number of forecasts of occurrence to the
394 number of observed occurrences,

$$B = \frac{a + b}{a + c} \quad (9)$$

395 and the Pierce Skill Score (PSS) gives the hit rate minus the false alarm rate.

$$PSS = \frac{ad - bc}{(b + d)(a + c)} \quad (10)$$

396 An unbiased forecast has a frequency bias $B = 1.0$. PSS values range from -1.0 to 1.0, with
397 positive values indicating that the hit rate was greater than the false positive rate, and therefore
398 the model had greater skill than a random forecast or constant CHAB or no-CHAB prediction
399 [*Hogan, R. J. and I. B. Mason, 2012*].

400 To provide a reference forecast for skill comparison, we defined a “persistence” forecast as the
401 assumption of no change from the satellite image that was used to initialize the model, which
402 represents the best available information to a forecast user in the absence of a useful model. We
403 took the further steps of filling in missing data in the persistence forecast with the most recent
404 satellite data for each spatial segment, and applying the same shoreline buffering procedure that
405 was used to initialize the model.

406 To test whether the model had significantly greater skill than the persistence forecast, we used the
407 bootstrap method described by Hogan and Mason [2012] to estimate the confidence interval
408 around the difference in skill score of the model compared to the persistence forecast. Starting

409 with a series of n triplets of observations, model predictions, and persistence predictions, we
410 created 1000 different bootstrap samples, each of length n , by taking random samples with
411 replacement from the series. We then calculated the difference in PSS , ΔPSS , for each bootstrap
412 sample. Finally, we estimated the 95% confidence interval as the 0.025 to 0.975 quantiles of the
413 ensemble of 1000 values of ΔPSS . While analytical formulas are available to estimate the
414 uncertainty in PSS , the bootstrap method accounts for effects of spatial and temporal
415 autocorrelation in environmental data, which effectively reduce the number of independent
416 observations to be $< n$ [Hogan, R. J. and I. B. Mason, 2012].

417

418 **3 Results and Discussion**

419 Western Lake Erie is polymictic, meaning that it does not continuously stratify during the
420 summer owing to shallow bathymetry and exposure to wind. Temperature profiles simulated by
421 FVCOM show periods of temporary stratification that are strongest during calm afternoons when
422 the surface is warmed by the sun and warm summer air (Figure 2a, 19-20 Aug.). At night,
423 cooling of the surface often causes deepening of the surface mixed layer by convection. This diel
424 cycle can be overpowered by shear-induced mixing during windy periods (Fig. 2a, 21-22 Aug.).
425 The temperature difference over the water column during periods of stratification is small (Fig.
426 2a), but the static stability is sufficient to cause turbulent diffusivity to vary by orders of
427 magnitude over a depth range of a few meters (Fig. 2b). Random walk turbulence schemes are
428 susceptible to formation of artificial accumulations of particles in the presence of strong gradients
429 in diffusivity [Visser, A., 1997].

430 **3.1 Vertical random walk schemes**

431 We tested the random walk schemes using a 1D well-mixed condition simulation (see Methods)
432 that was run using hourly time series of diffusivity profiles output from FVCOM for the month of
433 August 2011 at stations representative of the range of conditions that occur during the summer
434 CHAB season in western Lake Erie (Fig. 1). The range of conditions represented by the
435 diffusivity profiles can be characterized by defining a Peclet number that represents the ratio of
436 mixing time scale to floating/sinking time scale of the water column, $Pe = w_b h / \bar{K}$, where h is
437 the water column depth and \bar{K} is the column mean eddy diffusivity [Ross, O. N. and J. Sharples,
438 2004]. Values of $Pe \gg 1$ indicate that w_b has a strong influence on particle concentration
439 profiles. Combining the time series of diffusivity profiles that was used in the well-mixed
440 condition simulations with $w_b = 70 \mu\text{m s}^{-1}$ (see below), values ranged on the order of $0.01 < Pe <$
441 100 . A small time step is usually required for $Pe \ll 1$ because small h and large \bar{K} will produce
442 strong gradients in diffusivity for realistic diffusivity profiles (i.e., $K \sim 0$ at the bottom or at the
443 thermocline), and therefore, small values of the Visser time step criterion (Eq. 3). At the 3-m
444 deep station (WE6) the variable time step occasionally was limited by the specified minimum and
445 maximum values of 0.01 and 600 s, with typical hourly means of 0.2-3 s. At the 13-m deep
446 station (NDBC45005) in the central basin the water column was continuously stratified, and
447 longer time steps could be used; hourly minimum values of the variable time step were typically
448 0.2-2 s, mean values were 3-30 s and maximum hourly values were constrained by the upper
449 limit of 600 s.

450 Example time series of concentration profiles output from a well-mixed condition simulation are
451 shown in Figure 2. A typical particle accumulation artifact is visible in Fig. 2c, where the
452 normalized concentration deviated from the expected constant value of unity. The artifact formed
453 when high diffusivity in the surface mixed layer on 24 August (Fig. 2b) caused particles to jump
454 across the sharp diffusivity gradient into the area of low diffusivity in the lower half of the water
455 column without the opportunity for the gradient term (Eq. 3-4) to push the particles back toward
456 the high diffusivity SML. Improved performance can be seen in Fig. 2d, where the variable time
457 step was reduced during the high diffusivity event on 24 August, thereby limiting the maximum
458 particle displacements and reducing the magnitude of the artifact.

459 The Gräwe Milstein scheme [Gräwe, U., 2011] with the variable time step provided the best
460 combination of computational efficiency and accuracy of the random-walk numerical schemes
461 tested. The shortest run time was achieved by the Gräwe Milstein scheme (Fig. 3a), which was
462 unique in the use of linear interpolation of diffusivity to the particle position, while the other
463 schemes used cubic splines at a greater computational cost. The second order schemes, PC2 and
464 S1.5, required greater computational effort to calculate additional terms and had the longest run
465 times. The run time of the variable time step simulation was similar to that of the fixed 1 s time
466 step for the Visser and Milstein schemes because the average of the variable time step was
467 similar to the value of the fixed 1 s time step (Fig. 3a). The Milstein schemes showed improved
468 accuracy compared to the Visser scheme with the fixed 1 s time step (Fig. 3b), consistent with the
469 finding of Gräwe [2011]. All schemes were more accurate with the variable time step than with
470 the fixed 1 s time step (Fig. 3b) because the Visser time step criterion (Eq. 5) was always

471 satisfied in the case of the variable time step. The second order schemes did not offer sufficiently
472 improved accuracy to compensate for their greater computational effort (Fig. 3a,b).

473 In contrast to our result, Gräwe (2011) found that the second order schemes did offer improved
474 accuracy that justified the additional computational effort, but for the case of idealized diffusivity
475 profiles specified at high vertical resolution and for a realistic test case of a tidally-mixed bay
476 with model diffusivity output at 200 levels. In our case of a shallow polymictic lake, diffusivity
477 profiles were highly irregular with sharp gradients (Fig. 2b), and diffusivity was output at only 20
478 levels. We found that spline fits often had spurious features between the levels at which
479 diffusivity was specified by FVCOM that were not representative of physically realistic
480 diffusivity profiles. Higher-order random walk schemes depend on higher-order derivative terms
481 from the spline fits to the diffusivity profiles [Gräwe, U., 2011], which may not be accurate in the
482 case of a non-representative spline fit. The higher order schemes might produce better results if
483 we were to output diffusivity at a large number of levels, but that would come at the expense of
484 greater computational effort in the hydrodynamic model. For our application, the Gräwe Milstein
485 scheme produced satisfactory results and did not require a spline fit, so it was selected for further
486 work.

487 **3.2 *Microcystis* colony size distribution and buoyant velocity**

488 The parameter w_b represents the *Microcystis* colony terminal velocity resulting from the balance
489 of forces between buoyancy and fluid drag. Our approach was to specify a *Microcystis* colony
490 size distribution, then apply an empirical relationship between w_b and colony diameter to obtain a
491 frequency distribution of w_b for use in the model. According to Stoke's law for the terminal

492 velocity of a floating/sinking spherical particle in a fluid, one might expect the relationship
493 between w_b and colony diameter to give a straight line on a log-log plot with a slope of 2.
494 However, Nakamura et al. [1993] showed that *Microcystis* colony specific gravity approaches
495 that of the surrounding fluid as colony diameter increases owing to the fractal geometry of the
496 colonies and the increasing volume of void spaces filled with the surrounding water; the result is
497 that the slope of the log-log plot is < 2 .

498 The *Microcystis* colony diameter frequency distribution measured by FlowCam was unimodal
499 with a median of 117 μm and a maximum of 740 μm (Fig. 4a, Station WE12, 4 Aug 2014). The
500 size frequency distribution measured by microscopy on samples collected at stations WE 2,4,6,
501 and 8 in July – October 2013 and June – July 2014 gave a similar size distribution to that of the 4
502 August 2014 sample (Fig. 4a). It is likely that the colony size distribution varies to some extent
503 spatially and temporally [e.g., Lin, L. et al., 2014], and our estimate could be refined through
504 additional measurements. Even so, the consistency between our two estimated size distributions
505 gives some indication of representativeness.

506 Our measured values of Lake Erie *Microcystis* colony buoyant velocity, w_b , were similar to those
507 of Nakamura et al. [1993] for colonies larger than 200 μm in their sample collected from a lake in
508 Japan on 3 August 1990, and generally less than their 18 September sample (Fig. 4b). We were
509 not able to resolve colonies smaller than 200 μm by our method; however, large colonies account
510 for the majority of biomass and toxin concentration. For example, colonies $> 112 \mu\text{m}$ accounted
511 for 93% of *Microcystis* cells (biomass) in Lake Erie samples [Chaffin, J. D. et al., 2011], and
512 colonies $> 100 \mu\text{m}$ showed the highest proportion of microcystin-producing genotypes, highest

513 microcystin cell quotas, and highest microcystin production rate, compared to smaller colony size
514 classes in Lake Wannsee, Germany [Kurmayer, R. et al., 2003]. In addition to colony size,
515 *Microcystis* buoyancy is a function of *Microcystis* strain and light exposure history, as it affects
516 gas vacuole and carbohydrate content of the cells [Ibelings, B. W. et al., 1991; Xiao, Y. et al.,
517 2012]. Further research is necessary to define buoyant velocities over a wide size range of Lake
518 Erie *Microcystis* under a variety of environmental conditions. Our results, while limited in size
519 range, do show similarity between the > 200 μm values of Nakamura et al. [1993] and samples
520 from two different dates in Lake Erie, and support using the lower estimate of buoyancy (Fig. 4b,
521 N93 3 Aug.) from Nakamura et al. [1993].

522 For the model simulations, we assigned buoyant velocities to Lagrangian particles by random
523 sampling with replacement from the frequency distributions shown in Figure 4c, which were
524 obtained by applying the regression lines (Fig. 4b) from the data of Nakamura et al. [1993] to the
525 diameter frequency distribution from the 8 August 2014 sample from Lake Erie (Fig. 4a). We
526 tested the sensitivity of 1D model simulations to the two buoyant velocity frequency distributions
527 shown in Figure 2c, and refer to these hereafter as “N93 3 Aug” and “N93 18 Sep”. Example
528 time series of concentration profiles simulated with the low estimate of buoyancy (N93 3 Aug)
529 are shown in Figure 2e.

530 **3.3 Vertical profiles of cyanobacterial concentration and temperature**

531 We tested the ability of the random-walk model with buoyancy to simulate realistic *Microcystis*
532 concentration profiles by comparing measured profiles of cyanobacterial chlorophyll

533 concentration from Lake Erie (predominantly *Microcystis*) to corresponding 1D simulations. On
534 20 July, the concentration profile showed strong accumulation within the surface two meters
535 (Fig. 5a), which corresponded to a surface mixed layer defined by a thermocline at 2-m depth
536 (Fig. 5b). A second profile was measured on 9 September, which showed concentration
537 enrichment within a 3-m thick surface mixed layer (Fig. 5c), which was similarly defined by a
538 thermocline at 3-m depth (Fig. 5d).

539 The temperature difference across the thermocline in both cases was only about 1 °C (Fig. 5b,d),
540 but the FVCOM simulations indicated that such subtle stratification features can have a strong
541 influence on diffusivity (e.g., Fig. 2a,b). The accuracy of temperature simulations in
542 hydrodynamic models is often only within a few degrees, which brings into question whether the
543 subtle stratification features that are influencing the *Microcystis* vertical distribution can be
544 reasonably simulated by a hydrodynamic model. Even though the simulated temperature profiles
545 have a warm bias of 1-2 °C, at these locations and times, they show thermoclines at multiple
546 levels that are similar to the observed profiles (Fig. 5b,d). The deeper thermocline may have
547 formed due to convective deepening of the SML overnight, followed by surface warming during
548 the day that produced the shallower thermocline; the profiles were captured in the afternoon. It is
549 the static stability of the profile rather than the absolute temperature that is important in
550 simulation of the diffusivity, and the static stability of the simulated and observed profiles is in
551 reasonable agreement (Fig. 5b,d). Over the full set of 69 profiles, the frequency distribution of
552 static stability simulated by FVCOM was in reasonable agreement with the observed frequency

553 distribution, although the model was biased slightly less stable than the observations (Fig. 6b,
554 Table 2).

555 Simulated normalized concentration profiles of buoyant particles showed enrichment within the
556 surface mixed layer, similar to the observed profiles (Fig. 5a,c). We calculated the center of mass,
557 σ_m , of the concentration profile as a column-integrated indicator of the vertical distribution of
558 concentration (horizontal lines in Fig. 5a,c). Concentration was weighted toward the surface (σ_m
559 > -0.5) in $> 80\%$ of the observed profiles (Fig. 6a), which is consistent with the assumption to
560 treat *Microcystis* colonies as buoyant particles in the model.

561 We selected the lower estimate of buoyancy (N93 3 Aug) for use in the hindcast simulations. The
562 simulated frequency distribution of σ_m was in reasonable agreement with the observed
563 distribution for both the low and high estimates of w_b , although the low estimate was closer to the
564 observations (Fig. 6a, Table 2). Similarly, the direct measurements of w_b also indicated better
565 agreement with the lower estimate of buoyancy (Fig. 4b).

566 **3.4 Hindcast simulations of CHAB intensity and distribution**

567 Having shown that 1D random walk simulations reasonably approximated the changing vertical
568 distributions of buoyant *Microcystis* colonies in response to varying turbulence, we went on to
569 test whether inclusion of this mechanism in the forecast model improved model skill. Hindcast
570 simulations were initiated from each of the 26 quality satellite images of CHAB distribution for
571 the 2011 CHAB season.

572 In one example, a hindcast simulation was initialized on 6 August, which was a calm day (wind <
573 5 m s^{-1}) with an intense CHAB event throughout the central western basin (Fig. 7a). On the
574 following day, wind increased ($5\text{-}10 \text{ m s}^{-1}$), and a second satellite image indicated reduced
575 surface CHAB intensity and distribution (Fig. 7d). The 3D simulation captured the reduced
576 surface CHAB intensity on 7 August, while the 2D model did not, which can be seen
577 qualitatively by comparing Figures 7e and 7f, and was indicated quantitatively by reduced
578 frequency bias (B) of the 3D simulation compared to the 2D simulation (3D B = 1.10; 2D B =
579 1.34). On day 9 (15 Aug.), the simulated CHAB distribution was distinctly different between the
580 2D and 3D models (Fig. 7h,i). In comparison to the 2D model, the 3D model CHAB distribution
581 was more similar to the observed distribution (3D PSS = 0.56; 2D PSS = 0.41), having less
582 CHAB coverage in the central basin east of Sandusky and more continuous coverage along the
583 coast from Monroe to Toledo. Both 2D and 3D models simulated the advection of CHAB to Port
584 Clinton (Fig 7b,c and h,i), which was minimally affected by CHAB on 6 August and fully
585 covered on 15 August (Fig. 7a,g).

586 In a second example, a hindcast simulation was initialized on 29 August, which was a date with
587 only partial coverage by the satellite image, leaving no data over much of the western basin (Fig.
588 8a). Output from a previous model run was used to initialize the CHAB distribution in the
589 western basin (Fig. 8b,c). On simulation day four (2 Sept.) a second partial satellite image
590 indicated extensive CHAB coverage in the western basin (Fig. 8d), consistent with both models.
591 Both 2D and 3D models underestimated the CHAB coverage, although the 3D model better
592 matched the observed coverage (2D B = 0.81, PSS = 0.76; 3D B = 0.90, PSS = 0.84; Fig. 8d,e,f).

593 The partial image on 2 September did not show the extensive CHAB outbreak into the central
594 basin east of Leamington, Ontario, although it was simulated by both 2D and 3D models (Fig.
595 8e,f), and was revealed the following day in the 3 September satellite image (Fig. 8g). The 3D
596 model better simulated the CHAB distribution on simulation day 5 (3 Sept.) than the 2D model
597 (2D B = 0.79, PSS = 0.68; 3D B = 0.99, PSS = 0.80; Fig. 8g,h,i).

598 The examples in Figures 7 and 8 show, that both 2D and 3D models capture some observed
599 events that may be attributed to advection, but the 3D model performed better in several cases.
600 The 3D model is initialized with a better estimate of total biomass than the 2D model because an
601 estimate of the surface mixed layer depth for buoyant *Microcystis* colonies is used to assign the
602 depth over which the satellite-derived surface concentration is applied. In addition, the 3D model
603 is able to simulate changing surface concentration in response to changing mixed layer depth.
604 Finally, the 3D model produced different final CHAB spatial distribution than the 2D model,
605 which likely results from the more accurate vertical distribution within a complex 3D flow field.

606 **3.5 Summary of hindcast skill statistics**

607 Skill statistics were summarized by simulation day to evaluate how long the model can be run
608 from initialization before skill begins to decline. The Pierce skill score (PSS) gave positive values
609 for the 2D model, 3D model, and the persistence forecast on simulation days 1-10 (Fig. 9).
610 Positive values of PSS indicate that the hit rate was greater than the false positive rate, and
611 therefore the model had greater skill than a random forecast or constant CHAB or no-CHAB
612 prediction [Hogan, R. J. and I. B. Mason, 2012]. The frequency bias was less than 1.0 on 8 of 10
613 forecast days, indicating that both models had an overall bias toward under-prediction, although

614 not consistently so (Fig. 9a,b). The 95% confidence intervals on the difference in PSS indicated
615 that the 3D model displayed significantly greater skill in the hindcast simulations than the
616 persistence forecast through simulation day 6, and was not significantly worse than the
617 persistence forecast through day 10 (Fig. 10). The 3D model had significantly greater skill than
618 the 2D model over the full simulation period. The 2D model had significantly less skill than the
619 persistence forecast on all simulation days (Fig. 10).

620 It may be surprising that the persistence forecast displayed a reasonable level of skill. This can be
621 explained in that the spatial distribution of CHABs in Lake Erie has a number of persistent
622 features. For example CHABs often persist in the southern and western portions of the western
623 basin, while they are rarely present in the Detroit River plume and in the central basin east of the
624 islands, as indicated by 13 years of Lake Erie CHAB spatial patterns compiled by Wynne and
625 Stumpf [2015]. The model does not necessarily preserve these persistent features. For example
626 CHABs may be erroneously flushed from Maumee Bay in long simulations, although this
627 happened to a lesser extent in the 3D model than in the 2D model (Fig. 7g,h,i). The skill of the
628 persistence forecast indicates that the most recent satellite image is a reasonable indication of the
629 CHAB distribution for several days after.

630 Skill statistics based on pixel-by-pixel comparisons, and use of a persistence forecast as a
631 benchmark, provide a useful point of comparison among models, but do not capture all aspects of
632 model performance. For example, the large simulated plume that extended into the central basin
633 on 3 September (Fig. 8g,h,i) does not exactly match the observed plume in terms of shape and
634 position. This pattern mismatch detracted from pixel-by-pixel skill statistics, but both models

635 provided information regarding the existence of this transport event before it could be seen in
636 satellite imagery; information that would be useful to forecast users even if the shape of the
637 plume is not entirely accurate. A persistence forecast can score reasonably well in skill statistics
638 that compare spatial patterns, but cannot provide any information on likely transport trajectories.
639 Therefore, even though the 2D model had less skill than the persistence forecast (Fig. 10), this
640 does not indicate that the 2D model has no value because it may provide useful information on
641 likely transport trajectories. It is a challenge to formulate skill statistics that test for accuracy in
642 simulation of transport events, largely because it is difficult to identify and quantify transport
643 events by comparing among subsequent satellite images. Wynne et al. [2011] attempted to
644 calculate skill statistics based on movement of the bloom centroid; however, this approach is
645 likely to work only for special cases because accurate calculation of the bloom centroid is
646 sensitive to missing data (cloud cover) and the bloom often consists of multiple patches that may
647 move in different directions rather than one distinct patch. Formulation of appropriate skill
648 statistics for CHAB forecasts is an area for further work.

649 **3.6 Ecological significance**

650 Aside from the specific application of CHAB forecasting, the observations and simulations
651 shown here provide interesting insights on the physical processes that influence phytoplankton
652 ecology in a polymictic lake. Previous studies of Lake Erie circulation and thermal structure
653 considered the western basin to be largely unstratified [e.g., *Beletsky, D. et al.*, 2013], but our
654 study highlighted the importance of fine-scale thermal structure in the western basin in a
655 biological context. Surface mixed layer depth varies hour by hour due to subtle features in the

656 temperature profile caused by the diel cycle of surface heating and cooling, further modified by
657 varying wind stress. Colony buoyancy is sufficient to keep *Microcystis* concentrated within the
658 constantly changing surface mixed layer depth. The thermal structure is subtle in comparison to
659 the continuous seasonal stratification that occurs in deeper lakes, and in the central and eastern
660 basins of Lake Erie, but important nonetheless to the *Microcystis* vertical distribution. The
661 position of *Microcystis* colonies in the water column is critical to their light exposure, nutrient
662 acquisition, and ultimately to their ability to dominate the phytoplankton community, and
663 produce toxic blooms.

664

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676

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824

825 Table 1. Dates of measurements and model simulations.
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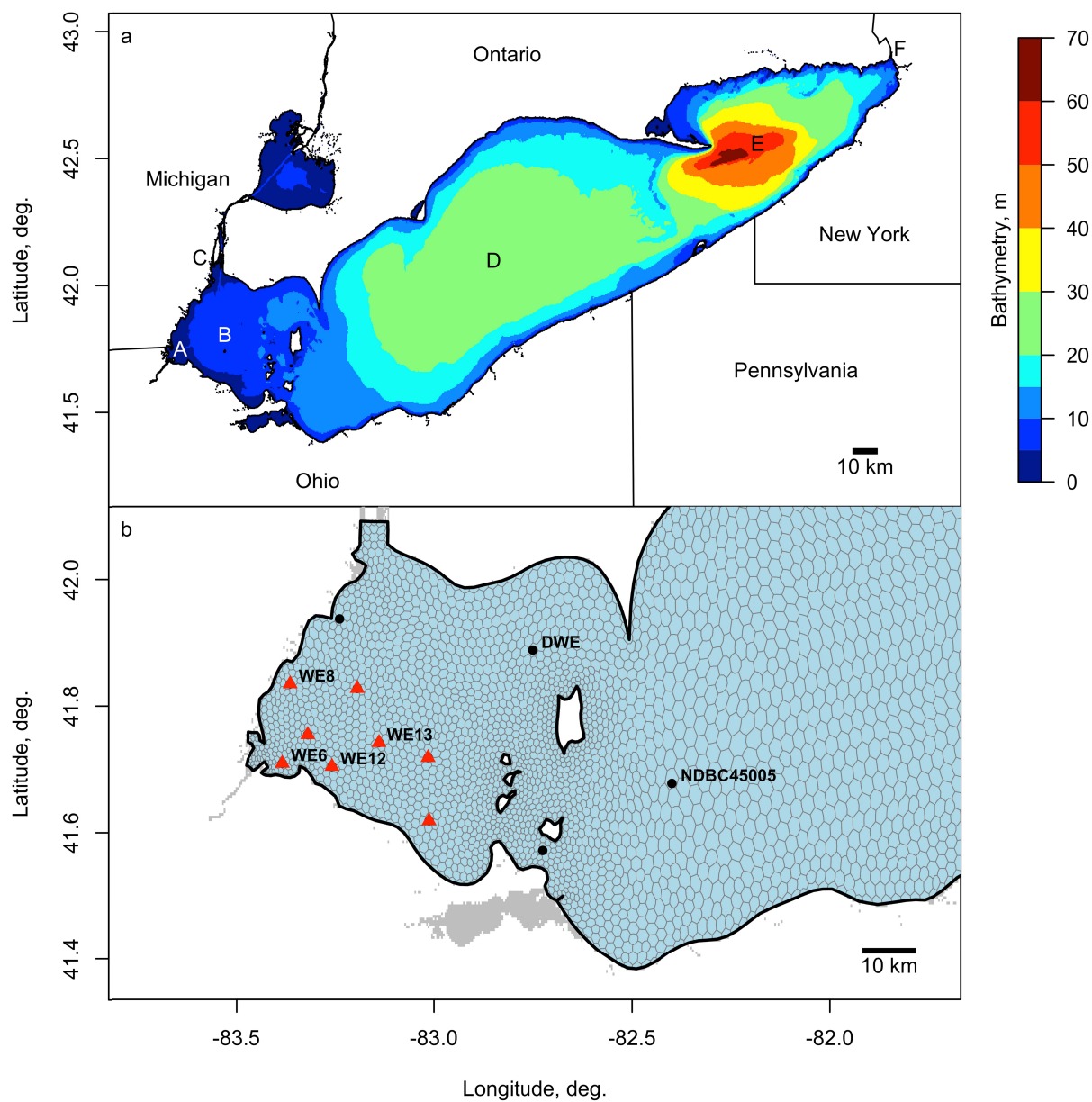
Measurement or simulation	Date
Well-mixed condition simulations	Aug., 2011
Satellite images, 2D, 3D simulations	26 dates, July to Oct., 2011
Colony size distribution (FlowCam)	4 Aug., 2014
Colony size distribution (microscopy)	Weekly sampling July to Sep., 2012 and 2013
Buoyant velocity measurements	15 and 21 July, 2015
Vertical profiles (FluoroProbe)	Weekly sampling July to Sep., 2015

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830 Table 2. Statistics evaluating the skill of the Lagrangian particle model in simulating the vertical
831 distribution of cyanobacterial chlorophyll concentration (center of mass of the normalized
832 concentration profile) and of FVCOM in simulating temperature profiles (potential energy
833 anomaly). The statistics are the mean bias, root mean square deviation (RMSD), and Pearson
834 correlation coefficient (r).

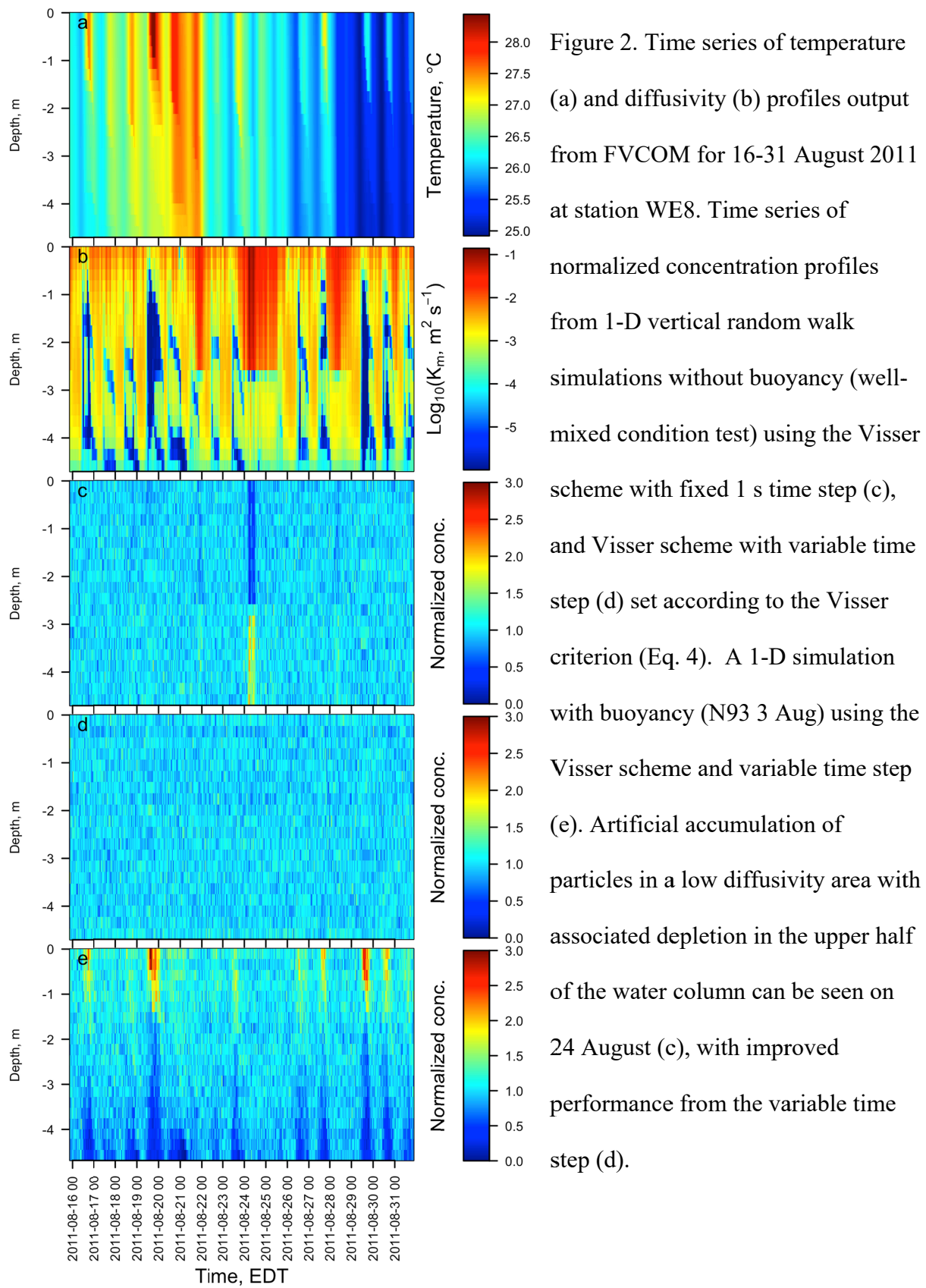
	Bias	RMSD	r
Center of mass, N93 18 Sep, σ	0.04	0.09	0.56
Center of mass, N93 3 Aug, σ	-0.01	0.08	0.53
Potential energy anomaly, J m^{-3}	-0.20	0.72	0.83

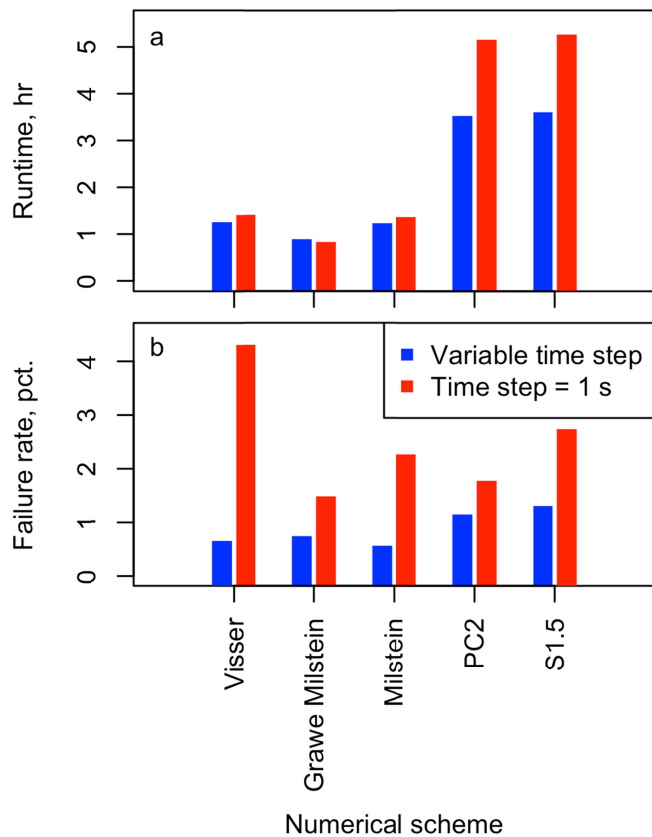
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 837 Figure 1. a) Geographic location and bathymetry of Lake Erie, showing bordering U.S. states and
 838 the Canadian province of Ontario. Features of interest are identified, including A) Maumee Bay,
 839 B) Western Basin, C) Gibraltar on the Detroit River, D) Central Basin, E) Eastern Basin, and F)

840 Buffalo on the Niagara River. b) An enlarged view of the western portion of the FVCOM model
841 domain. FVCOM domain boundaries are indicated with a heavy black line, and node-centered
842 tracer control elements with gray lines. Surface mixed layer depth was estimated in hindcast
843 simulations at the stations identified with symbols. Well-mixed condition simulations were
844 conducted at the named stations. Profiles of temperature and cyanobacterial chlorophyll
845 concentration were measured at the stations indicated by red triangles.
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 850 Figure 3. Performance comparison of the five vertical random walk numerical schemes and the
 851 variable time step scheme. One-dimensional vertical mixing simulations with 1000 particles were
 852 conducted for the month of August, 2011, at the six stations indicated in Figure 1. a) Total run
 853 time for the six simulations. b) percent occurrence of failure to meet the quality criterion out of
 854 4464 hourly records. The quality criterion was a signal to noise ratio > 5 where the signal was
 855 simulated normalized surface concentration with buoyancy (N93 3 Aug) and the noise was
 856 absolute deviation of the normalized surface concentration from 1.0 in a well-mixed condition
 857 test (without buoyancy). Concentration was normalized to the mean column concentration.

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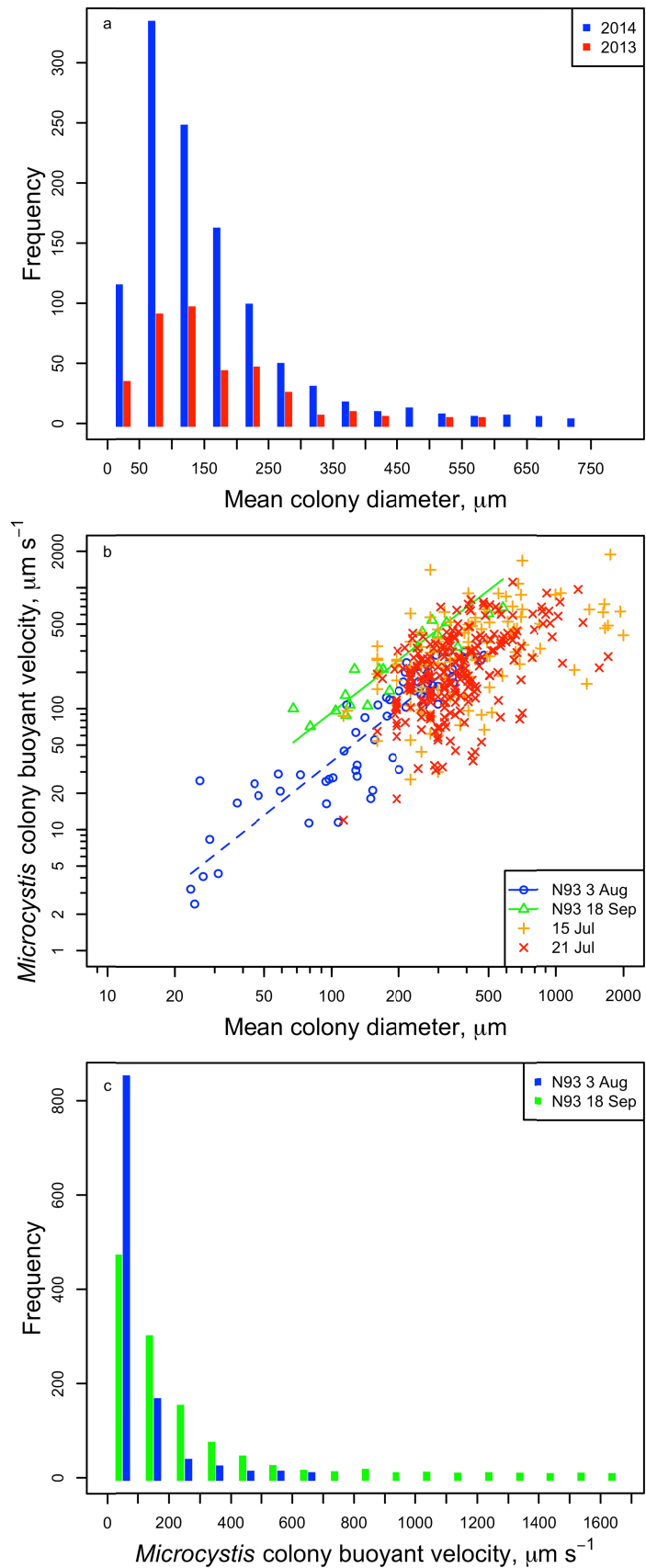
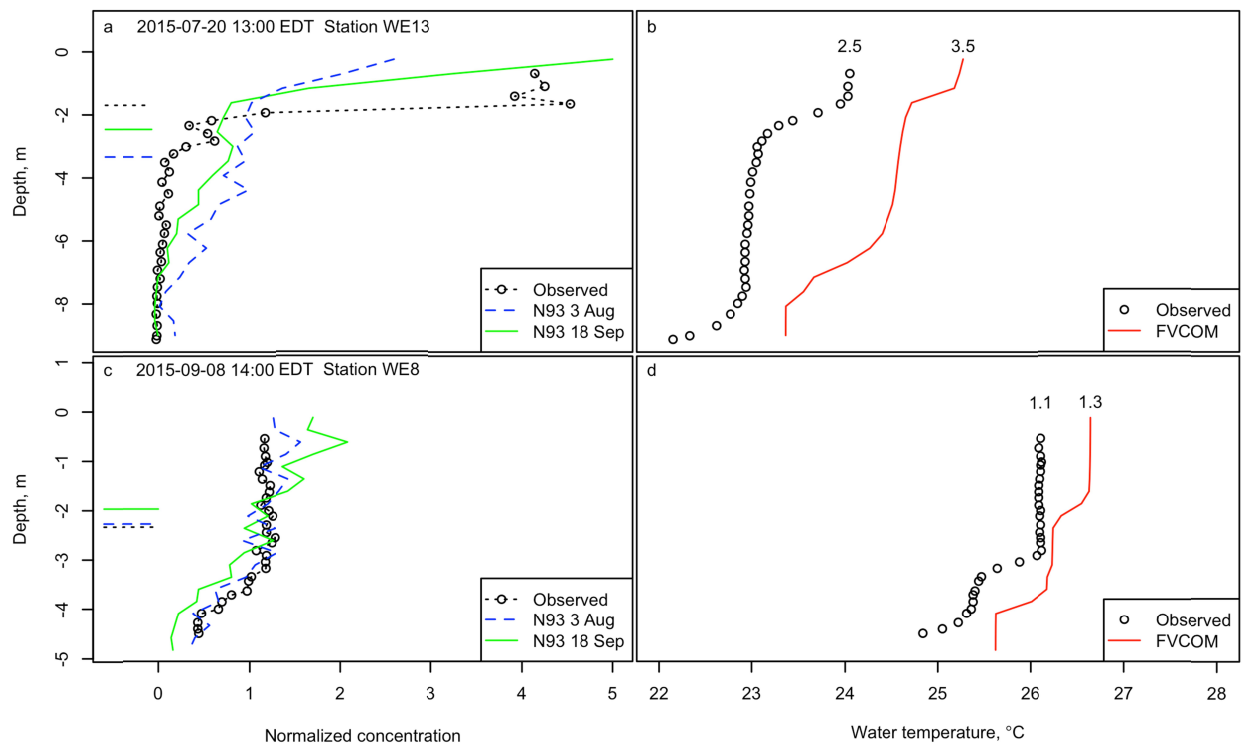


Figure 4. a) Histogram of *Microcystis* colony size distribution for samples collected in Lake Erie in 2013 and 2014. b) *Microcystis* colony buoyant velocity for samples collected in Lake Erie on 15 and 21 July, 2015, and data digitized from Nakamura et al. (1993, their Fig. 3). c) Histogram of *Microcystis* colony buoyant velocity resulting from application of the regression lines in Fig. 2b to the 2015 size distribution in Fig. 2a.



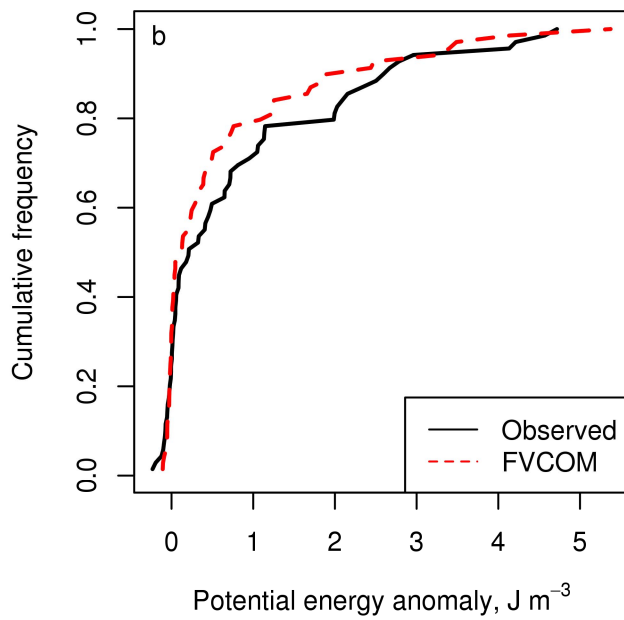
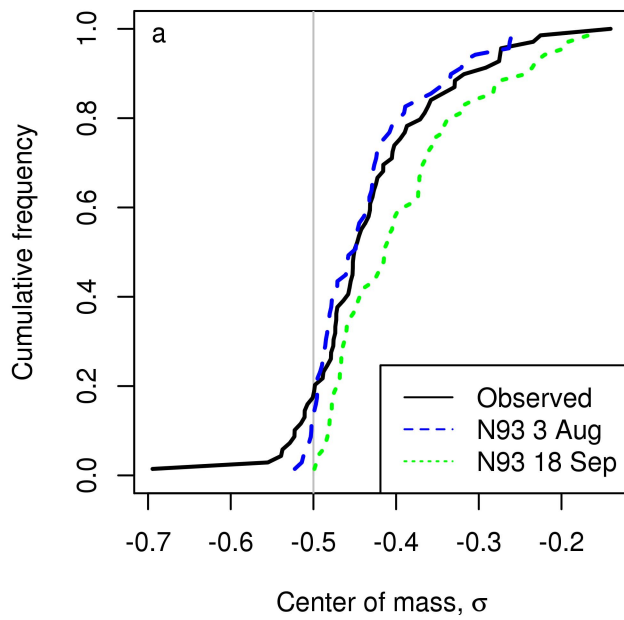
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863 Figure 5. Two examples of observed vertical profiles of normalized concentration of
 864 cyanobacterial chlorophyll and temperature with simulated values plotted for comparison.

865 Horizontal lines indicate the center of mass, σ_m , of the normalized concentration profiles, and the
 866 potential energy anomaly, ϕ , J m^{-3} , is given above the temperature profiles.

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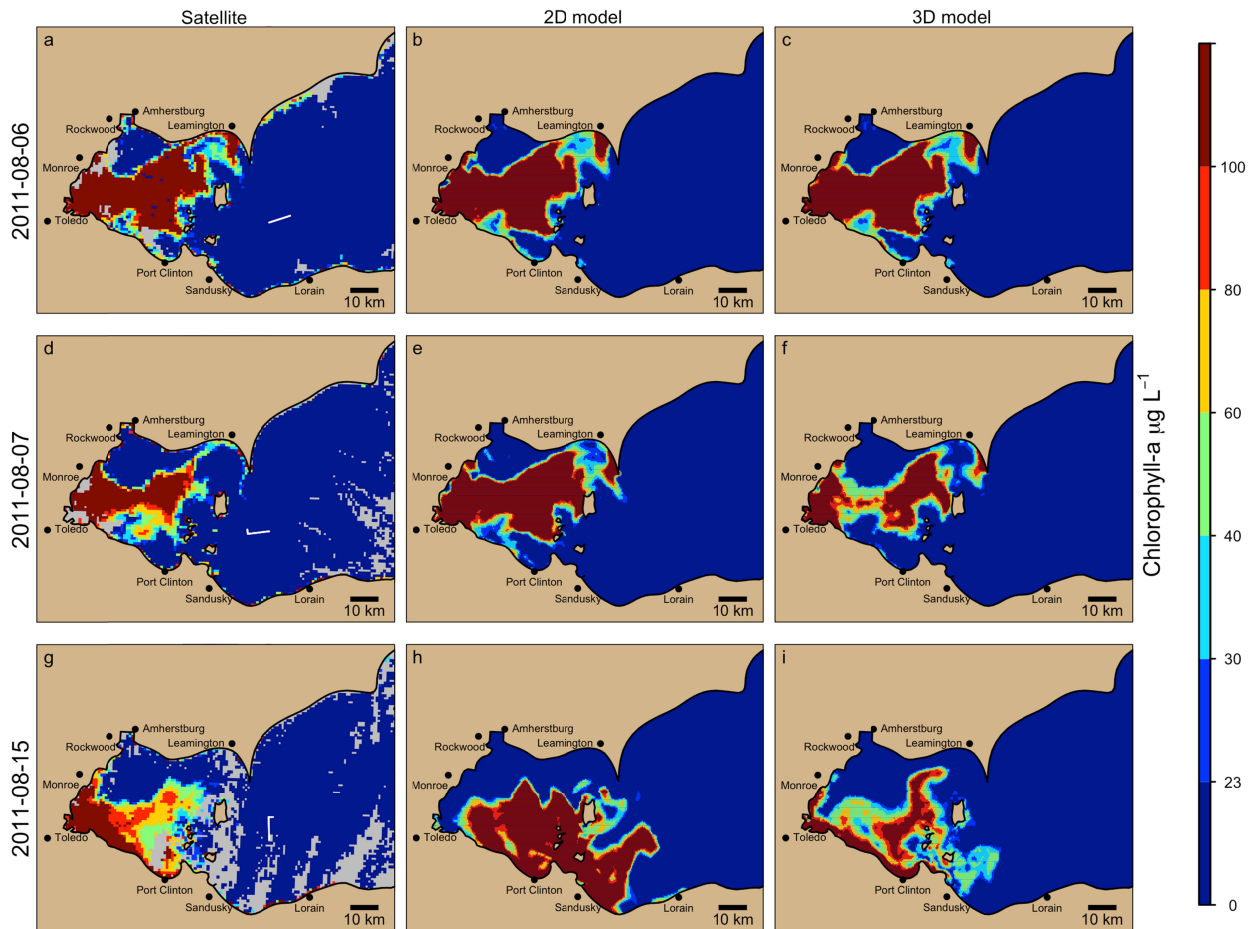


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870 Figure 6. Cumulative frequency distributions of simulated and observed a) center of mass of the
 871 normalized cyanobacteria concentration profile, and b) potential energy anomaly of the

872 temperature profile for 69 profiles collected in July - September of 2015 at stations indicated in
873 Figure 1.

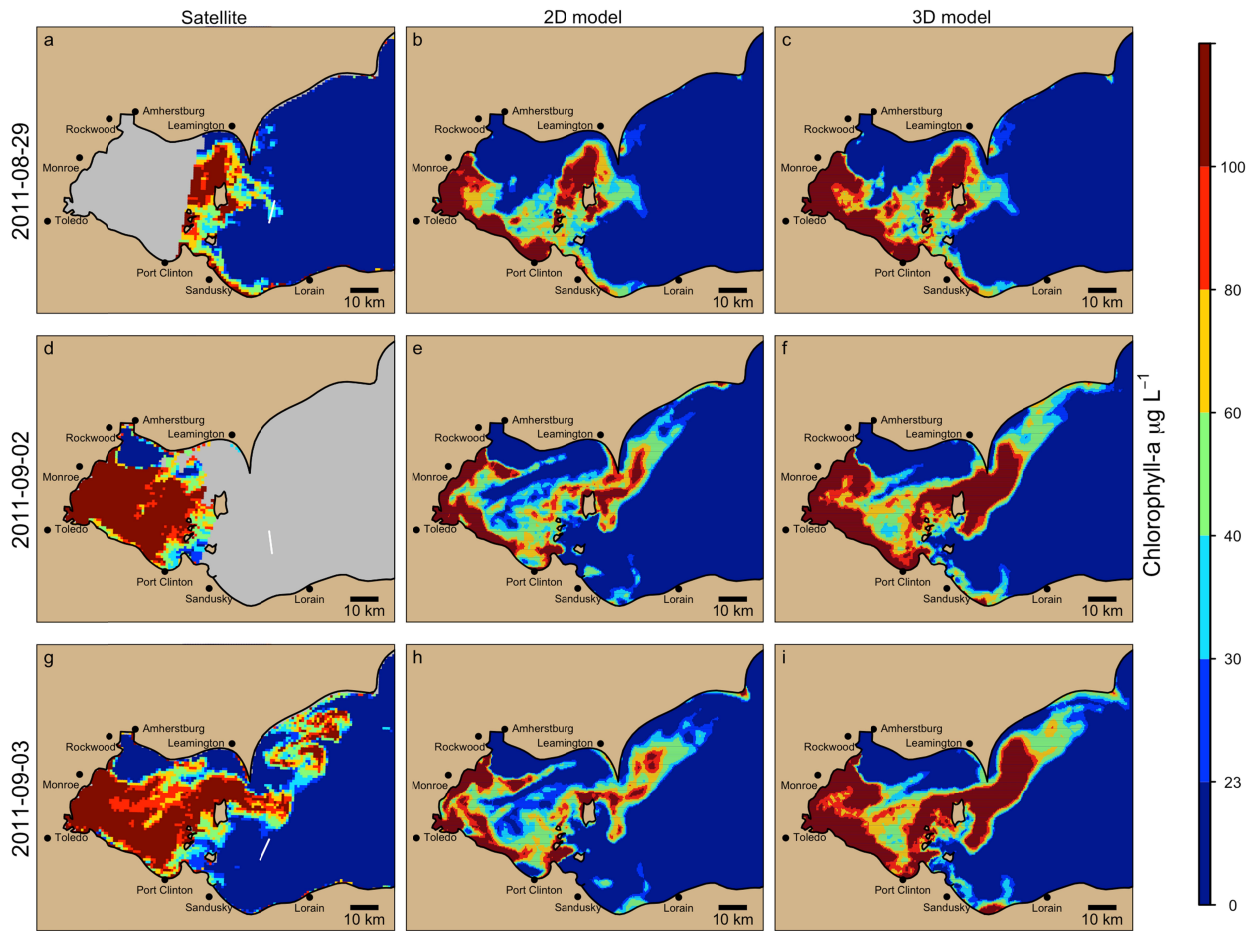
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876 Figure 7. Comparison of 2D and 3D hindcast simulations initialized from satellite-derived
 877 cyanobacterial chlorophyll concentration (a) on 6 August 2011. Subsequent satellite images (d,g)
 878 were used for model skill assessment. Gray color indicates missing data. Wind speed, averaged
 879 over the preceding 12 hours, was $< 5 \text{ m s}^{-1}$ on 6 August and increased to $5\text{-}10 \text{ m s}^{-1}$ on 7 and 15
 880 August (wind barbs a,d,g). In the 2D simulation, particles were initiated in the surface 1m and
 881 vertical mixing was not simulated (advection only). In the 3D simulation, particles were
 882 initialized over the surface mixed layer, as determined by preliminary simulations of 1D mixing
 883 with buoyancy, and the model was run with 3D advection in addition to vertical mixing with

884 buoyancy. While both 2D and 3D models simulated CHAB advection toward Port Clinton
885 observed on 15 August, the 3D model better simulated reduced intensity and coverage observed
886 on 7 and 15 August due to higher winds, and continued CHAB coverage near Toledo and
887 Monroe on 15 August.
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889
 890 Figure 8. Comparison of 2D and 3D hindcast simulations initialized from satellite-derived
 891 cyanobacterial chlorophyll concentration (a) on 29 August 2011. Symbols and model setup are
 892 explained in Figure 8. While both 2D and 3D models simulated CHAB advection east of
 893 Leamington into the central basin observed on 3 September, the 3D model better simulated
 894 CHAB intensity and extent in the western basin observed on 3 September. Wind speed was < 5 m
 895 s^{-1} during the simulation period, but was north at 10 $m s^{-1}$ until 12 hours prior to the initial image.
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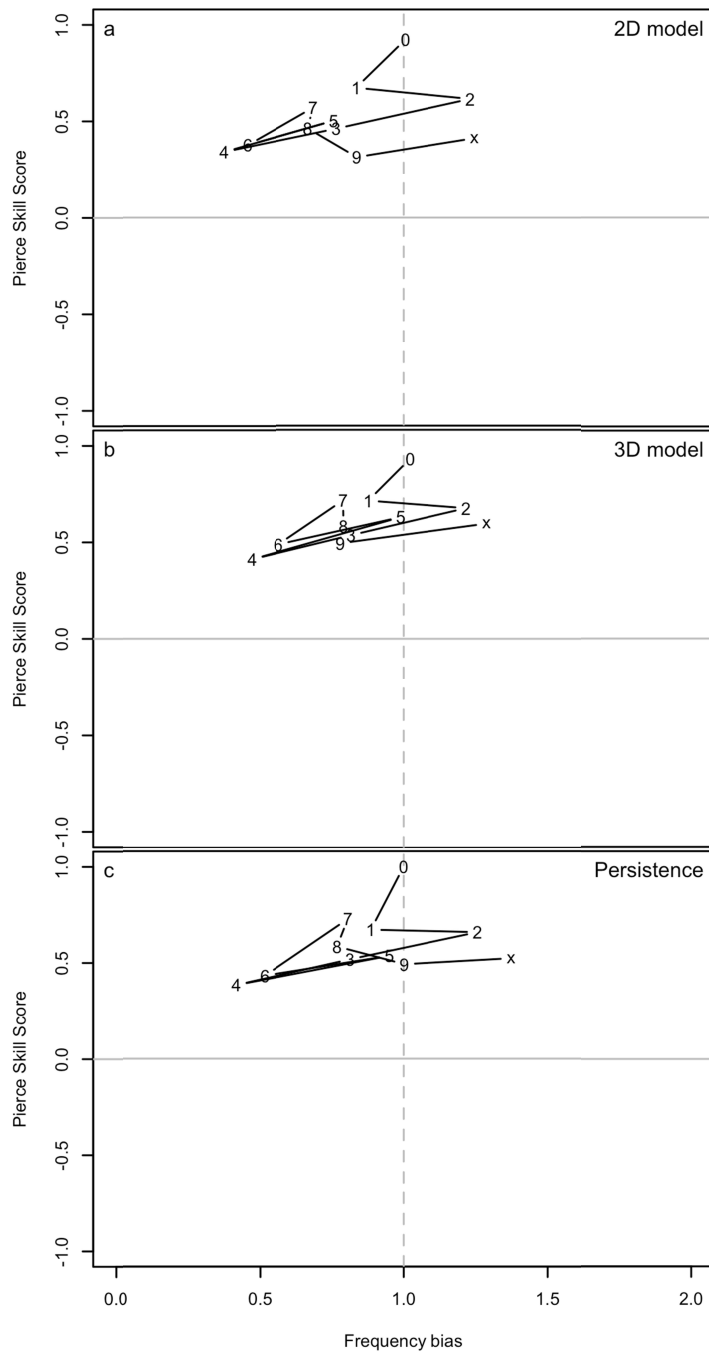
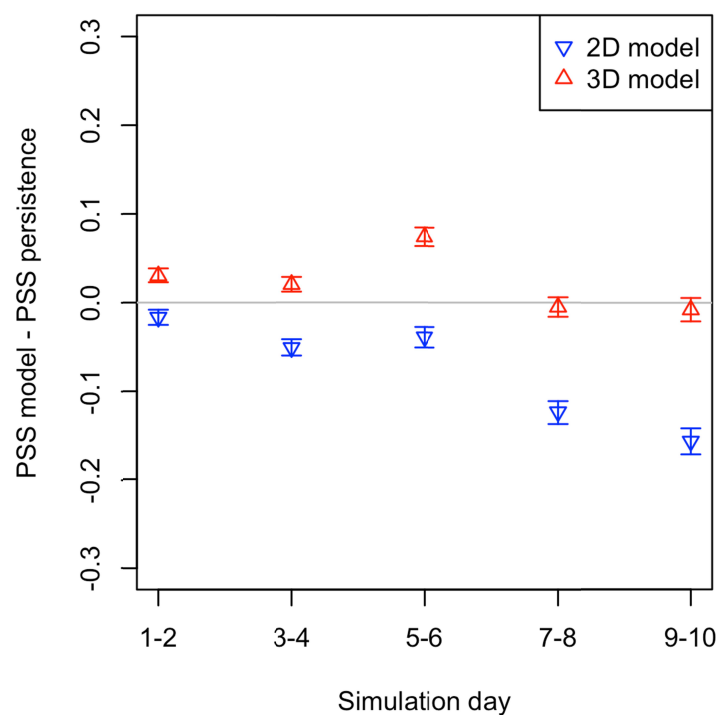


Figure 9. Skill-bias plots for the 2D model (advection only), 3D model (advection, vertical mixing, and buoyancy), and persistence forecasts for the 2011 hindcast simulations. The plot symbol indicates the simulation day (0 = initial, x = day 10). Pierce Skill Score (PSS) is the hit rate minus the false detection rate, and frequency bias (B) is the ratio of forecast hits to observed hits. Positive PSS indicates greater skill than a random forecast. Frequency bias of 1.0 indicates the same number of CHAB pixels were predicted as observed.

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 900 Figure 10. Pierce skill score (PSS) of the model minus PSS of the persistence forecast. Positive
 901 values indicate greater skill for the model than for the persistence forecast. Error bars indicate the
 902 95% bootstrap confidence interval on the difference in PSS for the 26 hindcast simulations from
 903 the 2011 CHAB season, grouped into two-day intervals. The 3D model (including vertical
 904 mixing with buoyancy) had greater skill than the 2D model (advection only) and greater skill
 905 than the persistence forecast through day 6 and comparable skill out to day 10.

Figure 1. Figure

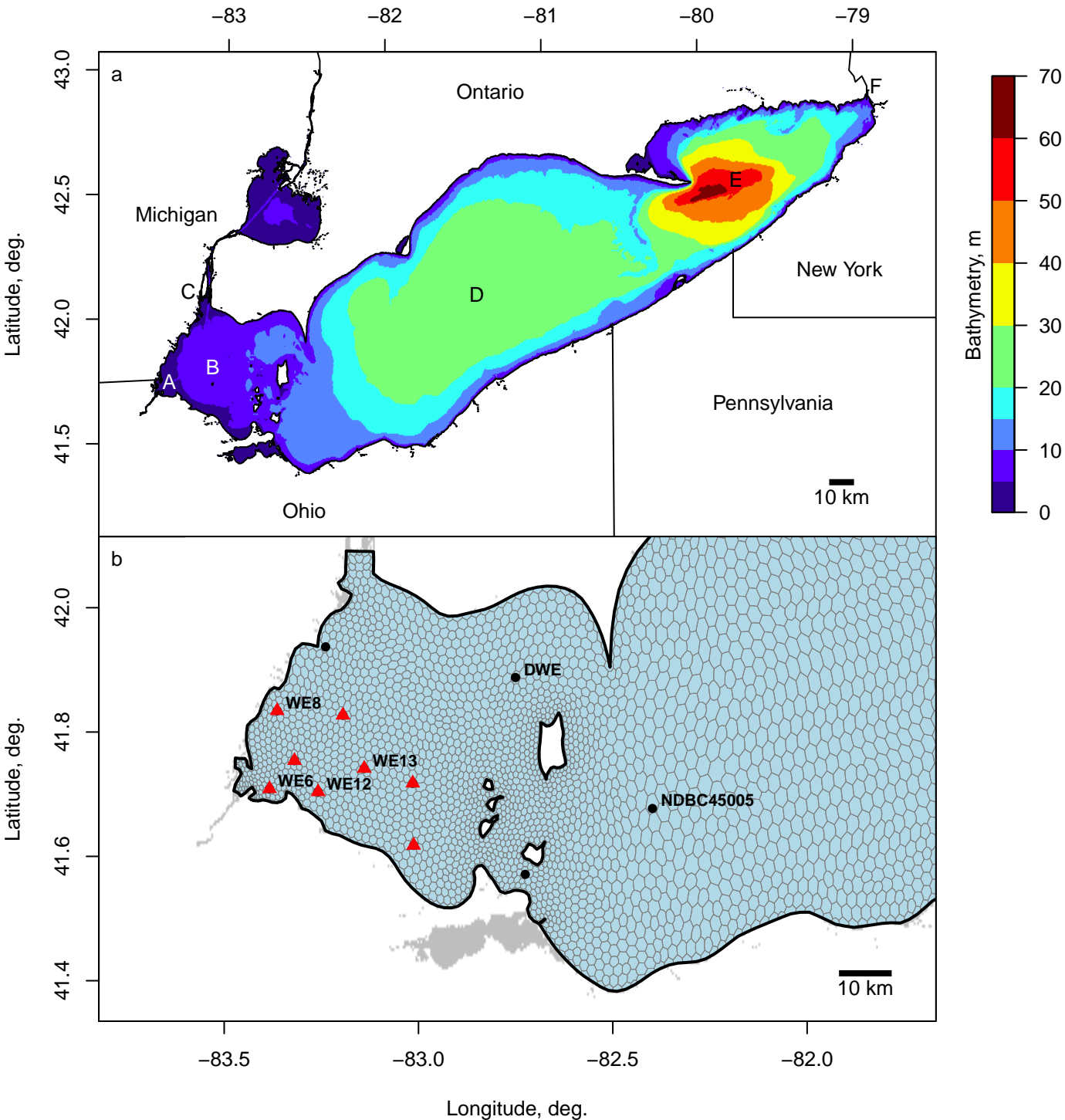


Figure 2. Figure

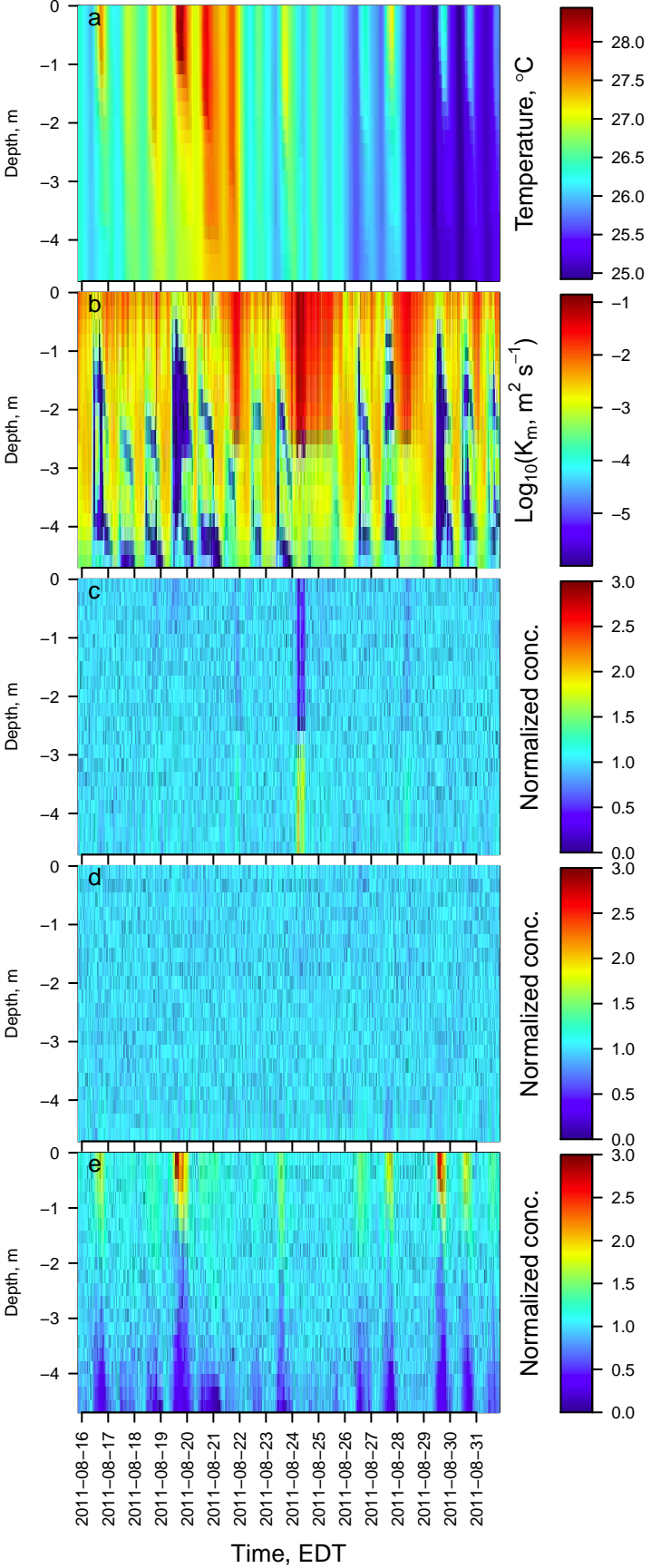


Figure 3. Figure

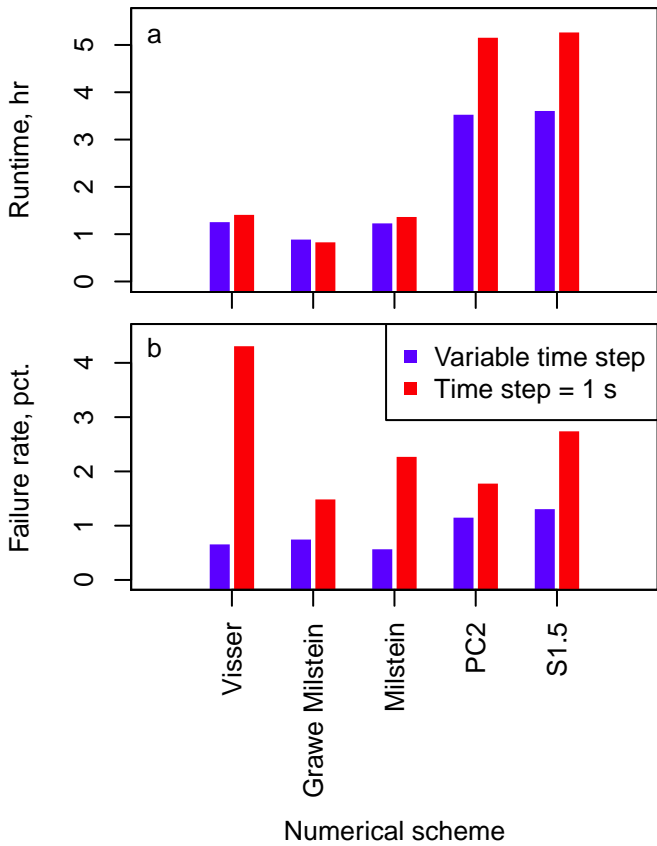


Figure 4. Figure

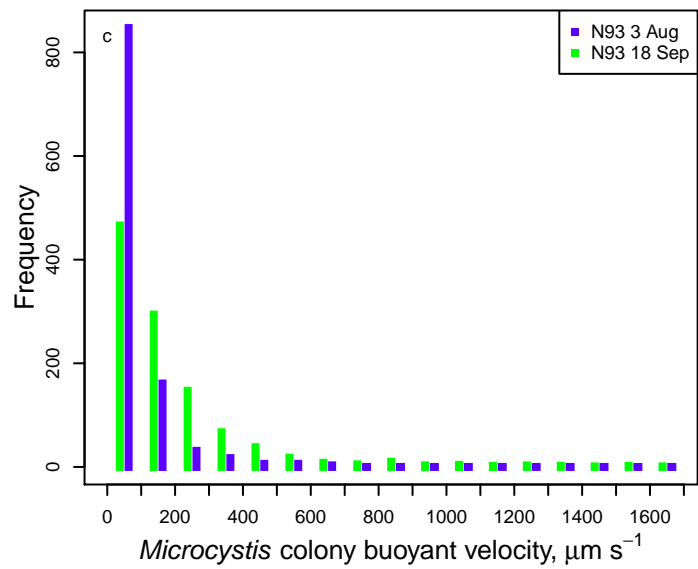
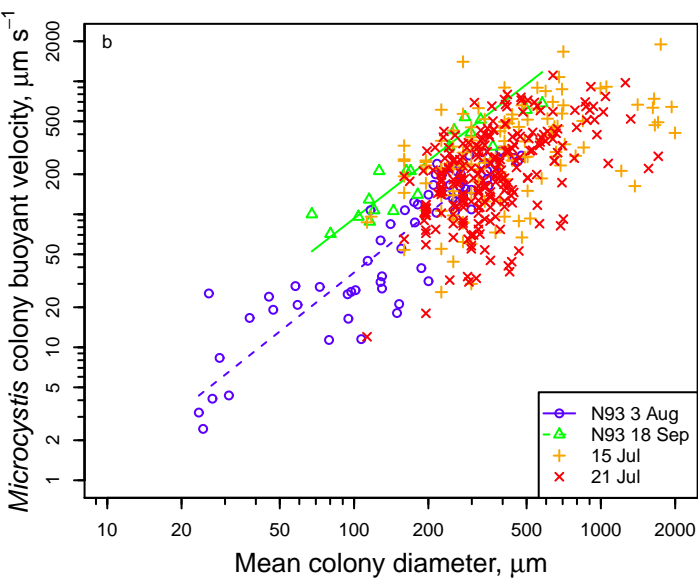
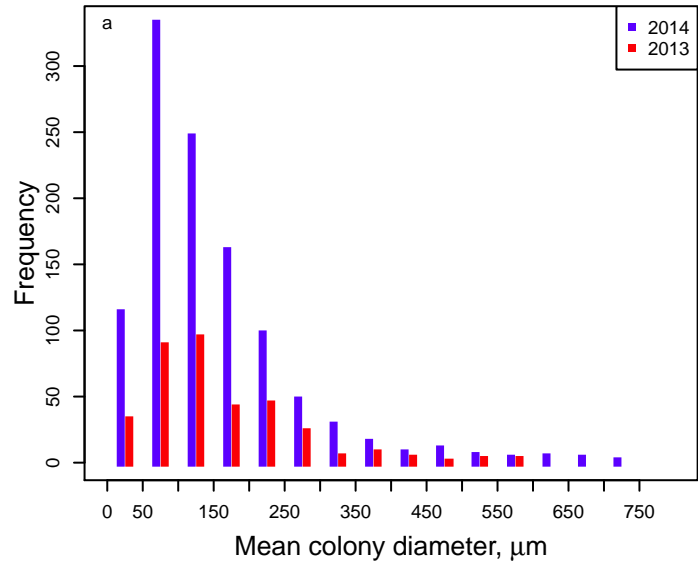


Figure 5. Figure

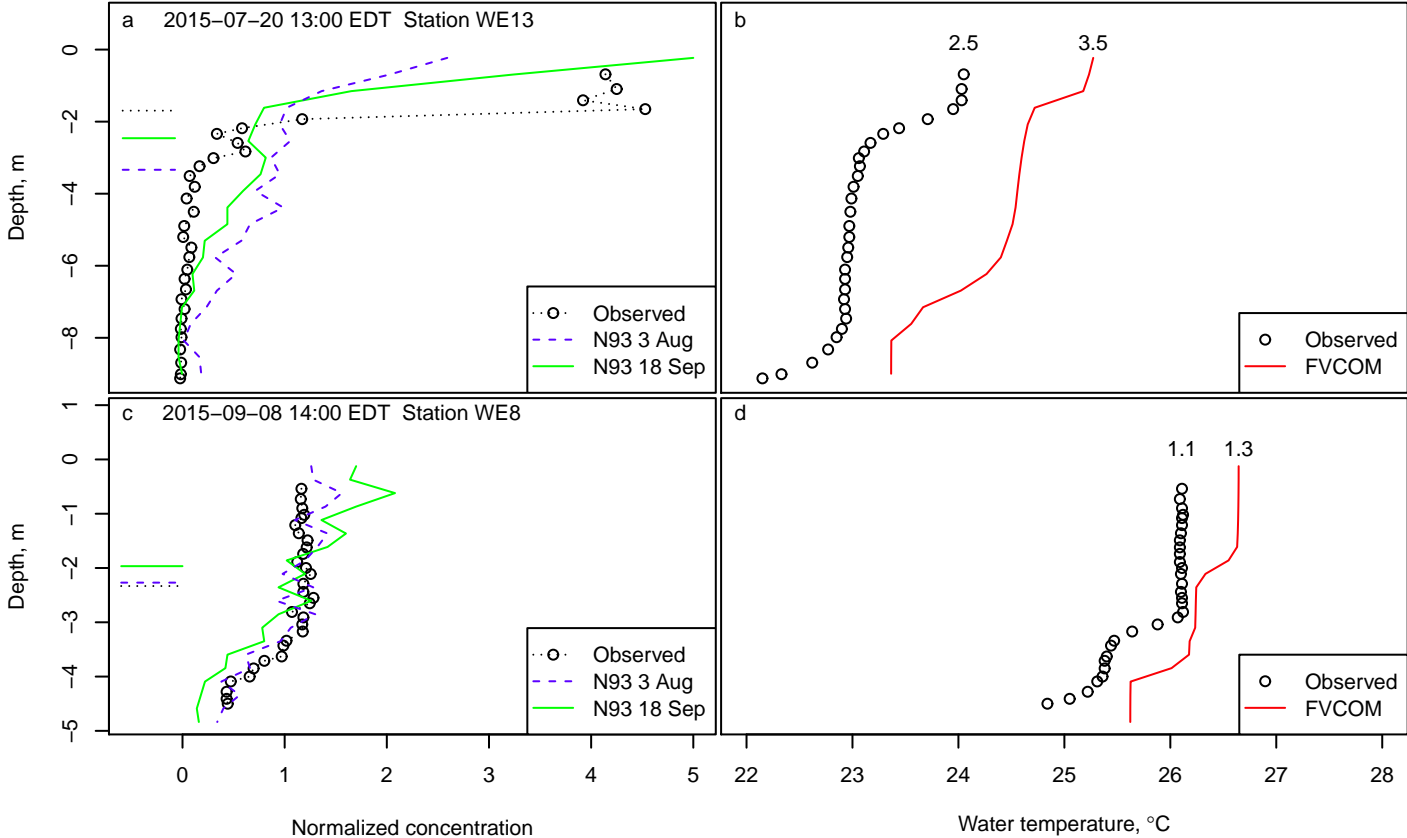


Figure 6. Figure

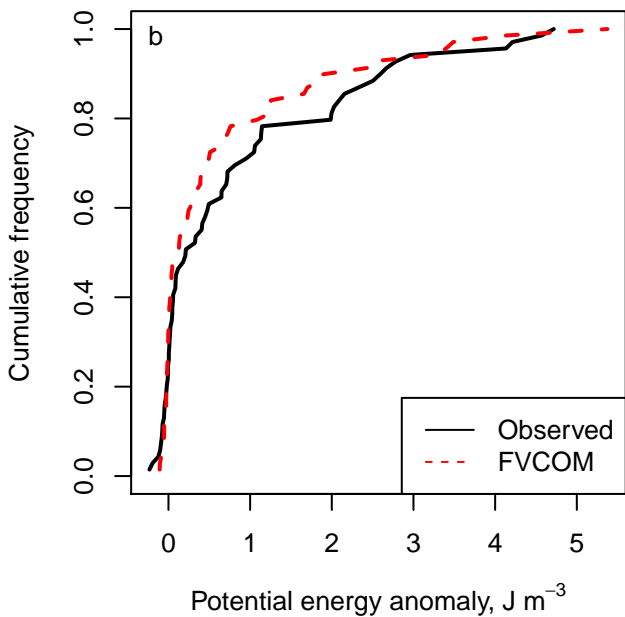
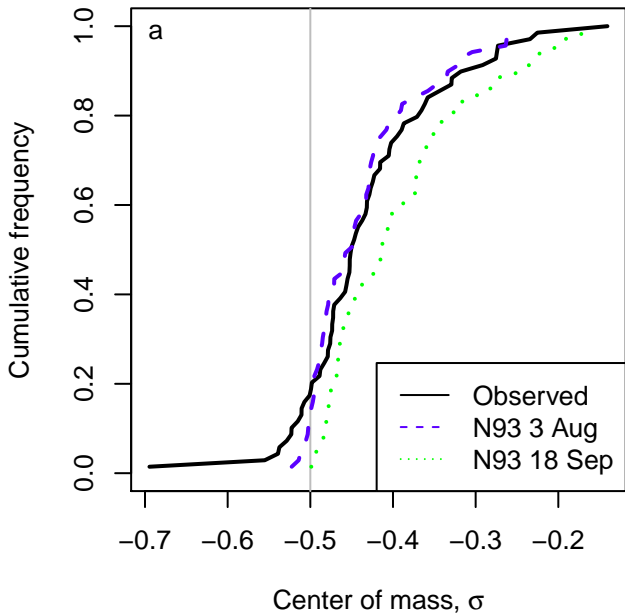


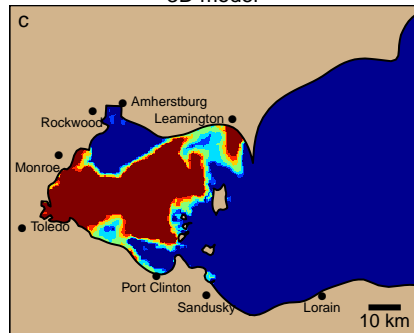
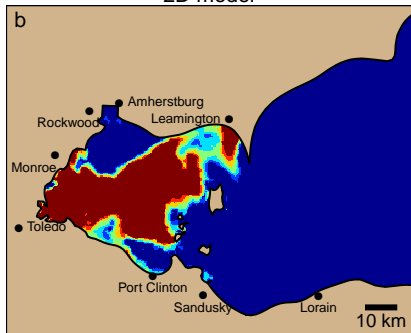
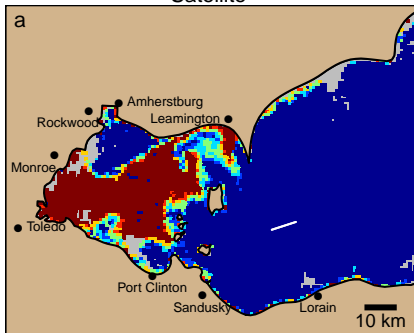
Figure 7. Figure

Satellite

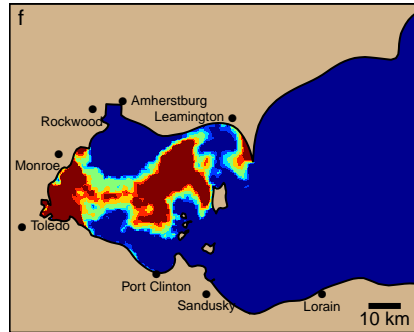
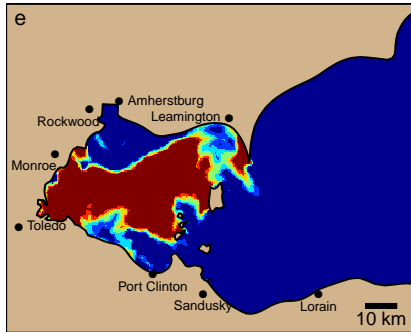
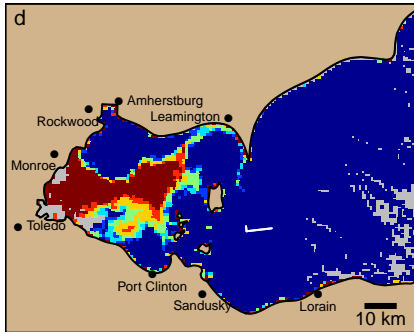
2D model

3D model

2011-08-06



2011-08-07



2011-08-15

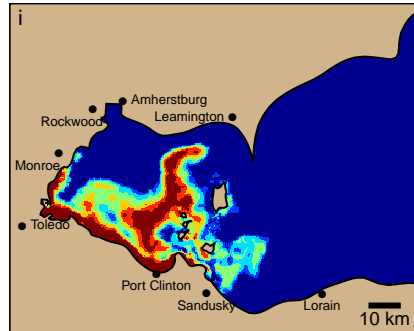
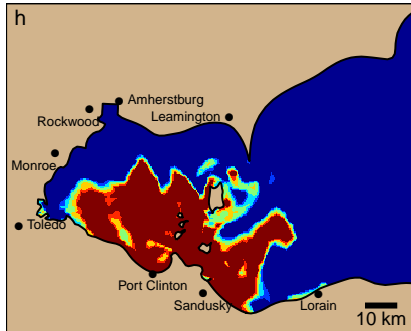
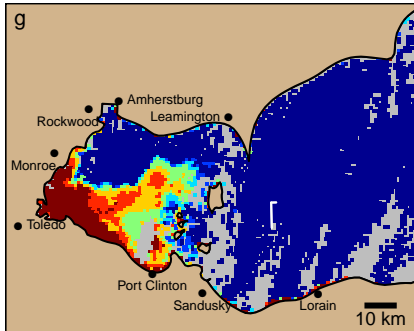
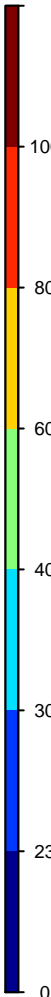
Chlorophyll-a $\mu\text{g L}^{-1}$ 

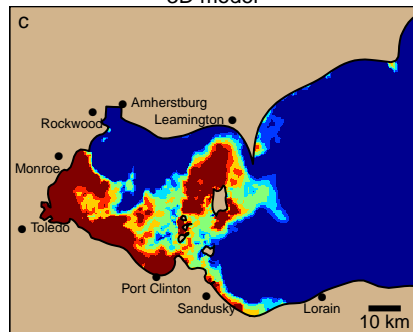
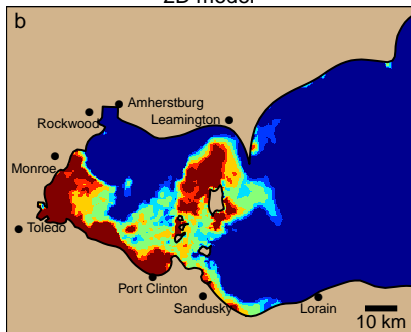
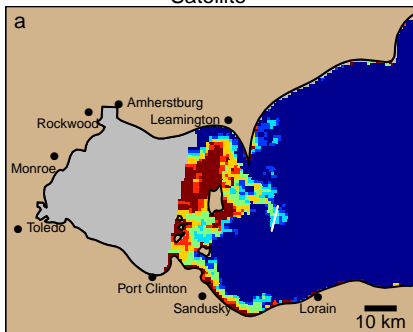
Figure 8. Figure

Satellite

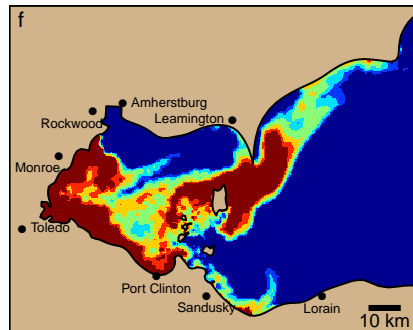
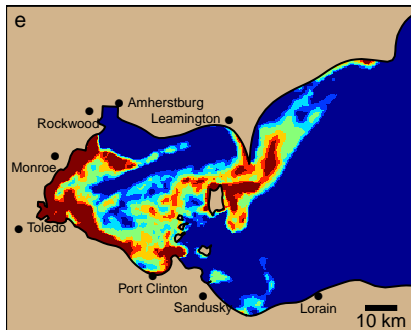
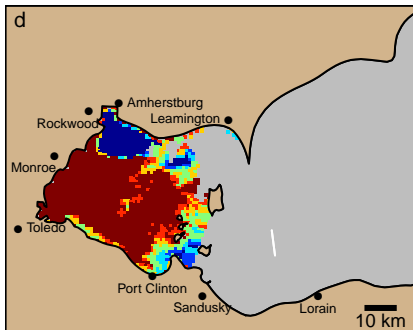
2D model

3D model

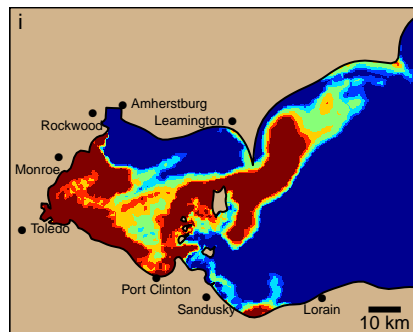
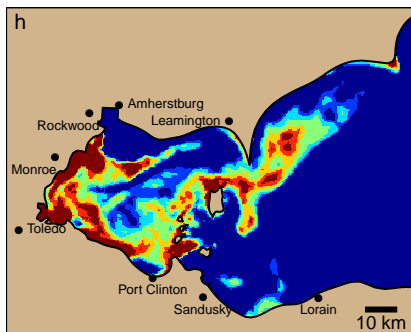
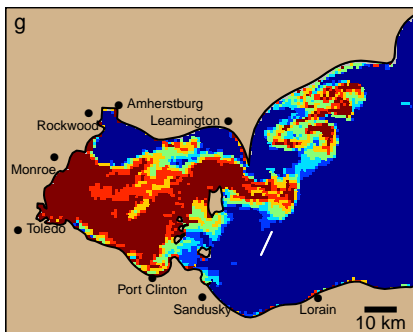
2011-08-29



2011-09-02



2011-09-03

Chlorophyll-a $\mu\text{g L}^{-1}$

100

80

60

40

30

23

0

Figure 9. Figure

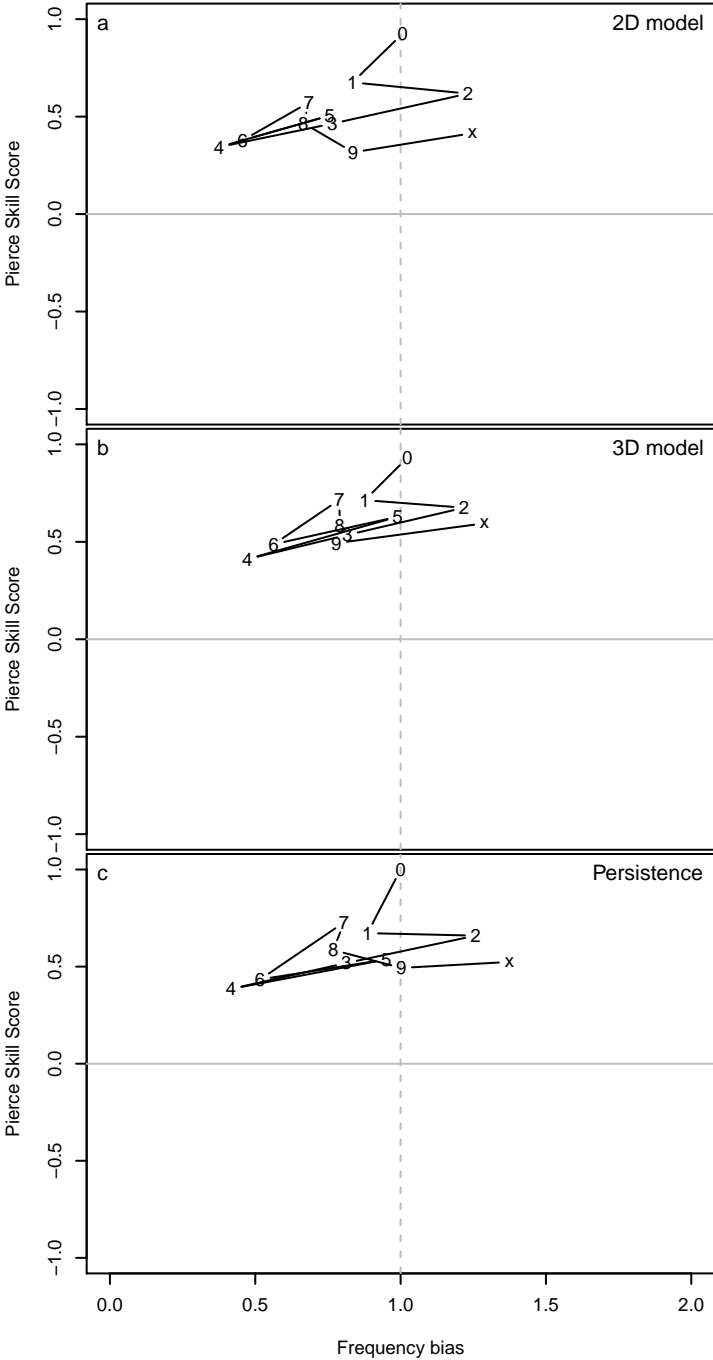


Figure 10. Figure

PSS model - PSS persistence

