



Are predator–prey model predictions supported by empirical data? Evidence for a storm-driven shift to an alternative stable state in a crab–clam system

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ABSTRACT: A dynamic systems approach can predict steady states in predator–prey interactions, but there are very few examples of predictions from predator–prey models conforming to empirical data. Here, we examined the evidence for the low-density steady state predicted by a Lotka–Volterra model of a crab–clam predator–prey system using data from long-term monitoring, and data from a previously published field survey and field predation experiment. Changepoint analysis of time series data indicate that a shift to low density occurred for the soft-shell clam *Mya arenaria* in 1972, the year of Tropical Storm Agnes. A possible mechanism for the shift is that Agnes altered predator–prey dynamics between *M. arenaria* and the blue crab *Callinectes sapidus*, shifting from a system controlled from the bottom up by prey resources, to a system controlled from the top down by predation pressure on bivalves, which is supported by a correlation analysis of time series data. Predator–prey ordinary differential equation models with these 2 species were analyzed for steady states, and low-density steady states were similar to previously published clam densities and mortality rates, consistent with the idea that *C. sapidus* is a major driver of *M. arenaria* population dynamics. Relatively simple models can predict shifts to alternative stable states, as shown by agreement between model predictions (this study) and published field data in this system. The preponderance of multispecies interactions exhibiting nonlinear dynamics indicates that this may be a general phenomenon.

KEY WORDS: *Mya arenaria* · *Callinectes sapidus* · Blue crab · Soft-shell clam · Alternative stable state · Lotka–Volterra · Nonlinear dynamics · Chesapeake Bay

1. INTRODUCTION

Predators play a key role in ecosystem stability and function by consuming dominant competitors (Lubchenco & Gaines 1981, Boudreau & Worm 2012). Predators can also destabilize ecosystems or collapse food webs if they become too abundant (Estes et al. 2009), or if their prey do not have natural defenses against predation (Johnston et al. 2015). Except in the case of new introductions (e.g. invasive species), populations of predators and prey that have overlapping distributions have generally coevolved adapta-

tions that allow them to coexist (Vermeij 1994). Prey have anti-predator behaviors or morphological adaptations to avoid being eaten (Bibby et al. 2007, Whitlow 2010). Similarly, predators have adaptations or behaviors that help them to forage optimally and take advantage of prey when they are available (Meire & Ervynck 1986, Rindone & Eggleston 2011).

One of the ways the balance between predator and prey adaptations manifests itself in nature is through density-dependent predation. Predators can exhibit a numerical response to prey densities by increasing reproduction rates due to an overabundance of prey

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(demographic response) or by gathering in areas with relatively high densities of prey (aggregative response) (Holling 1959). An individual predator may also adjust its predation rate to prey density through a 'functional response' (changes in a predator's consumption rate in response to prey density). Density-dependent mechanisms tend to stabilize prey population dynamics (Royama 1992, Turchin 2003) and can maintain population viability when a population is reduced to low levels (Cushing 1975).

Certain characteristics of a predator–prey system can help predict which functional response will be observed. A linear relationship between consumption rate and prey density (Type I functional response) is expected for organisms that do not actively search for prey, such as filter feeders. Most vertebrate and invertebrate predators exhibit a hyperbolic functional response that increases to an asymptote due to limits associated with prey handling, ingestion, and metabolism (Type II functional response) (Hassell et al. 1977). Predators that feed upon cryptic or otherwise hard-to-find prey exhibit a sigmoidal functional response, where consumption rates increase slowly at low prey densities (Type III functional response) (Holling 1959). Prey that avoid predators can achieve a low-density refuge; thus, the functional response can explain the distribution of prey resources, and it can be used to predict the persistence of prey species at low densities (Eggleston et al. 1992).

Many mathematical models can be used to predict predator–prey dynamics (Briggs & Hoopes 2004). These models contain nonlinear functions describing the density-dependent interactions between predator and prey. Due to nonlinearities, model behavior often includes shifts to alternative stable states (Drake & Griffen 2010). The states may include extinction of one or both species, or coexistence steady states where both predator and prey are able to coexist at densities predicted by the model. Multiple stable coexistence states are possible in fairly simple predator–prey functions (Mumby et al. 2007, Kramer & Drake 2010).

A dynamic systems approach can predict coexistence states in natural and managed systems. Steady states analysis has been used to develop optimum harvesting and conservation strategies for forest and rangeland systems (Hritonenko et al. 2013, Bauch et al. 2016). The theory surrounding steady states and bifurcations is well developed, but there are very few tests of the theory using empirical data. To date, coexistence steady states in predator–prey systems have been observed in populations of small organ-

isms such as unicellular organisms, yeast, small crustaceans, and bacteria (Luckinbill 1973, van den Ende 1973, Drake & Griffen 2010). Identifying or collecting data from coexisting predator–prey systems that can be used to test model predictions has proven difficult, especially for macro-organisms. Of the few examples of steady states that focus on macro-organisms, none examines interactions between natural populations of predators and prey (Kramer & Drake 2010, Jiang et al. 2018, McNickle & Evans 2018).

In this study, we examined evidence for a storm-driven shift to a low-density state for the soft-shell clam *Mya arenaria*, which was once a biomass-dominant species in Chesapeake Bay, USA, in the face of predation by the blue crab *Callinectes sapidus*. The objectives of the study were to (1) analyze over 30 yr of *M. arenaria* and *C. sapidus* time series data for changes in abundance following a major storm event; (2) develop a Lotka-Volterra model of the crab–clam predator–prey system from parameters available in the literature and analyze for steady states; and (3) examine the agreement between model predictions and empirical data using previously published data from a field survey and a field experiment.

History of the predator–prey system. Tropical Storm Agnes, which reached and remained in the Chesapeake Bay watershed from 21 to 23 June 1972, has long been suspected of causing long-term changes to the Bay (Orth & Moore 1983). Tropical Storm Agnes was a '100-year storm' that caused sustained, extremely low salinities and increased sedimentation throughout Chesapeake Bay (Schubel 1976, Schubel et al. 1976). This storm has been blamed for accelerating the loss of seagrass *Zostera marina* (Orth & Moore 1983) and oysters *Crassostrea virginica* (Haven et al. 1976) in Chesapeake Bay, although these losses were already in motion before the storm occurred.

Unlike seagrass and oysters, declines in abundance of the soft-shell clam *M. arenaria* were uniquely related to the storm. *M. arenaria* was abundant enough to support a major commercial fishery throughout Chesapeake Bay prior to 1972 (Haven 1970). Widespread mass mortality of *M. arenaria* occurred after the storm (Cory & Redding 1976), and the fishery never recovered in lower Chesapeake Bay (Virginia) (Glaspie et al. 2018). Attempts to revive a commercial fishery in Virginia waters were never realized after the passage of the storm. The commercial fishery for soft-shell clams in the Maryland portion of the Bay is characterized by variable and low harvest (Dungan et al. 2002); the fishery declined by 89% after the storm and has been near collapse since (NMFS Annual Commercial Landing Statistics, <https://www.st.nmfs>).

noaa.gov/commercial-fisheries/commercial-landings/annual-landings/index).

The failure of *M. arenaria* to recover from storm-related declines has been attributed to predation, habitat loss, disease, rising temperatures, and over-fishing (Glaspie et al. 2018). The Virginia and Maryland portions of Chesapeake Bay have different habitats, disease dynamics, climates, and fishing pressure; therefore, these factors are unlikely to explain the inability of *M. arenaria* to recover from low density in both regions (Dungan et al. 2002, Glaspie et al. 2018). More recently, disease has been blamed for an added minor decline in *M. arenaria* (Dungan et al. 2002); however, there is no evidence that disease prevalence or intensity are correlated with *M. arenaria* density (Glaspie et al. 2018).

Experimental evidence suggests that on a local scale, interactions between *M. arenaria* and its major predator, the blue crab *C. sapidus* (Meise & Stehlik 2003), are capable of keeping clams at low densities (Lipcius & Hines 1986, Seitz et al. 2001). *M. arenaria* burrows deeply in sediments, and although crabs can readily detect and excavate clams when they are at high densities, crabs are unable to detect their presence when clams are at low densities (Lipcius & Hines 1986). The result is a low-density refuge for *M. arenaria*, driven by disproportionately low predation, which is characteristic of a sigmoidal functional response (Lipcius & Hines 1986, Seitz et al. 2001). Given this evidence regarding a potential mechanism for the decline in *M. arenaria* and maintenance of the population at low density, this study examines the empirical evidence for a low-density steady state in this predator-prey system, and the impact of Tropical Storm Agnes on basin-scale population dynamics in *M. arenaria*.

2. MATERIALS AND METHODS

Change-point analysis of time series (Killick & Eckley 2014) was conducted using R statistical software (v. 3.6.0) on *Mya arenaria* landings (NMFS 2017) and log-transformed average adult female *Callinectes sapidus* abundance (Virginia Institute of Marine Science [VIMS] trawl survey) in the Chesapeake Bay from 1958–1992, with an Akaike's information criterion (AIC) penalty and using the segment neighbor algorithm (Auger & Lawrence 1989). Change-point analysis identifies the optimal number and location of changes in the mean of the data using an algorithm, in this case the segment neighbor algorithm, which produces an exact estimate of the change-point

(maximizing accuracy) at the expense of computing power (Auger & Lawrence 1989, Killick & Eckley 2014). The time period 1958–1992 was chosen for analysis because it begins when *M. arenaria* landings data first became available and ends before the slow decline in landings in the early 1990s due to fisheries collapse. Before and after change points, correlation between predator and prey time series was examined using cross-correlation function estimation in R. Correlation coefficients outside the 95% confidence intervals were considered evidence for temporal lags in correlation between predator and prey.

Predator-prey ordinary differential equation (ODE) models for the system including *M. arenaria* and *C. sapidus* were developed based on Lotka-Volterra models. The ODE for prey density (N) was modified with a Type III functional response:

$$N(t) = rN \left(1 - \frac{N}{K} \right) - f(N) P \quad (1)$$

where N is the density of prey, P is the density of predators, r is the intrinsic per capita growth rate, K is the carrying capacity, and $f(N)$ takes the form of a sigmoidal functional response:

$$f(N) = \frac{N^2 b T}{1 + cN + bT_h N^2} \quad (2)$$

where T is the time available for foraging, T_h is handling time, and b and c are components of the attack rate in a sigmoidal functional response (Hassell et al. 1977). The ODE model for predator density (P) was considered to be unrelated to prey density, because *C. sapidus* are generalist predators that may feed on other species (Hines et al. 1990, Williams 1990); however, *M. arenaria* are a preferred prey item for *C. sapidus* (Blundon & Kennedy 1982). For the purposes of this modeling effort, predator density was held constant at $P = 0.06 \text{ m}^{-2}$ (MD DNR Blue Crab Winter Dredge Survey, <https://dnr.maryland.gov/fisheries/Pages/blue-crab/dredge.aspx>).

Models were parameterized using data from the literature as follows: $r = 1.75 \text{ yr}^{-1}$ (Brousseau 1978), $K = 200 \text{ m}^{-2}$ (Abraham & Dillon 1986), $T = 1 \text{ yr}$, $T_h = 0.0015 \text{ yr}$ (Lipcius & Hines 1986), $b = 26.30 \text{ yr}^{-1}$ (Lipcius & Hines 1986), and $c = 0.14$ (Lipcius & Hines 1986). Analytical solutions of steady states were calculated using Matlab statistical software. Stability of each coexistence steady state was determined by examining the sign of eigenvalues.

The clam densities predicted at stable coexistence steady states were compared to observed densities of *M. arenaria* in Chesapeake Bay from a published field survey (Glaspie et al. 2018). Clam mortality

rates predicted by the ODE model were compared to mortality rates of *M. arenaria* exposed to *C. sapidus* predation in a field predation experiment (Glaspie & Seitz 2018). To examine mortality rates, we solved the equation for number consumed:

$$N_E = N - f(N)P \quad (3)$$

where N_E is the number of clams eaten calculated for a period of 8 d (0.022 yr) at an initial density of $N = 48 \text{ m}^{-2}$ to match the field predation experiments (Glaspie & Seitz 2018). We then calculated mortality as:

$$M = \frac{N - N_E}{N} \times 100\% \quad (4)$$

where M is percent mortality. Density of predators (P) was allowed to vary to achieve $M = 76.3\%$ (Glaspie & Seitz 2018), and the resultant P that achieved observed mortality rates of juvenile *M. arenaria* was compared to published *C. sapidus* densities for Chesapeake Bay. Data and R code files are available in the Knowledge Network for Biocomplexity repository (Glaspie 2019).

3. RESULTS

Change-point analysis identified an abrupt shift in clam abundance in 1972, the year of Tropical Storm Agnes, and this was the only change-point identified (Fig. 1). Before the storm, clam abundance was positively correlated with crab abundance at a lag of 1 yr ($r = 0.66$, $p = 0.01$; Fig. 2a). After the storm, clam abundance was negatively correlated with crab abundance with a lag of 1 yr ($r = -0.48$, $p = 0.04$; Fig. 2b). These were the only temporal lags for which there was evidence from cross-correlation analysis.

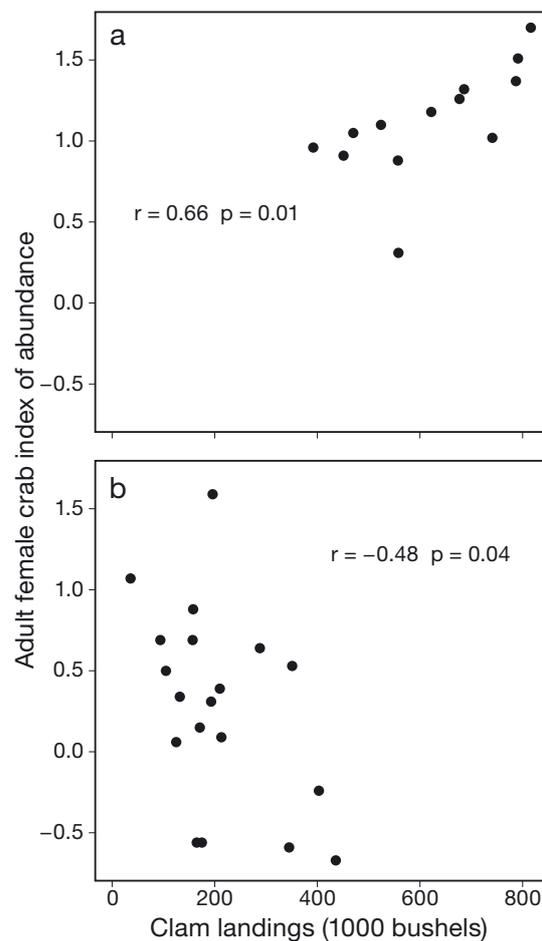
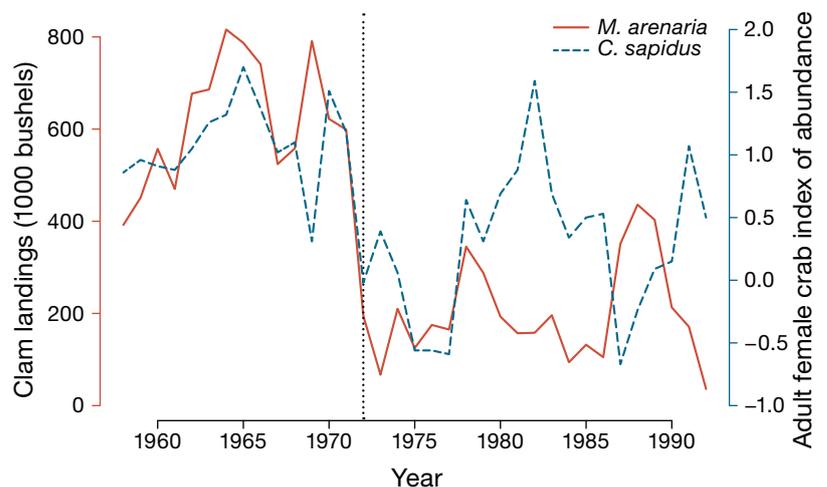


Fig. 2. Pre- and post-storm relationship between crab index of abundance (log-transformed average female abundance per tow, VIMS trawl survey) and clam landings (fisheries-dependent data). (a) Before the storm, crab abundance was positively correlated with clam abundance (1 yr lag). (b) After the storm, clam abundance was negatively correlated with crab abundance (1 yr lag)

Fig. 1. Predator–prey time series for soft-shell clam *Mya arenaria* landings (red) and adult female blue crab *Callinectes sapidus* index of abundance (blue). Blue crab data are log-transformed average female abundance per tow (VIMS trawl survey). *M. arenaria* data are fisheries landings (1000 bushels) (NMFS Annual Commercial Landing Statistics). Vertical dotted line represents Tropical Storm Agnes (1972) and the location of the change-point from the time series analysis



Predator–prey modeling confirmed the presence of high-density (near carrying capacity at 173.99 clams m^{-2}) and low-density (at 1.41 clams m^{-2}) steady states separated by an unstable steady state at 20.93 clams m^{-2} (Fig. 3). The low-density steady state predicted by the predator–prey model is similar to observed densities of *M. arenaria* in Chesapeake Bay, 0.4–1.73 m^{-2} (95% CI) (Glaspie et al. 2018). Mortality rates observed in the field were comparable to mortality rates predicted by the model for a blue crab density of 4.8 m^{-2} , which is a typical density for juvenile crabs in the summer months in Chesapeake Bay (Ralph & Lipcius 2014).

4. DISCUSSION

The observations, theory, and mechanistic basis indicate that *Mya arenaria* was subjected to a storm-driven shift to a low-density alternative stable state, which has been maintained by blue crab predation in Chesapeake Bay. Before the storm, clams were likely prey for juvenile crabs that recruited to the fishery at 1 yr of age, resulting in a positive correlation between crab and clam abundance with a lag of 1 yr (Fig. 2a). After the storm, crabs were likely consuming juvenile clams that would have recruited to the fishery a year later, resulting in a negative correlation between clams and crab abundance with a lag of 1 yr (Fig. 2b). Both *M. arenaria* and *Callinectes sapidus* recruit to their respective fisheries after 1–1.5 yr (Newell & Hidu 1986, Lipcius & van Engel 1990), providing an explanation for the 1 yr lag. This is consistent with a shift from a system controlled from the bottom up by prey resources, to

a system controlled from the top down by predation pressure on bivalves.

The results of the time series analysis would be difficult to explain with any other mechanism impacting *M. arenaria* population dynamics. Alternative explanations for persistent declines in *M. arenaria* density include disease, habitat loss, and environmental variability. *M. arenaria* is commonly infected by the parasitic protist *Perkinsus* sp., but infection intensity is not correlated with *M. arenaria* densities in Chesapeake Bay (Glaspie et al. 2018). Habitat loss in the form of declining seagrass has had a negative impact on *M. arenaria* (Glaspie et al. 2018) and *C. sapidus* alike (Anderson 1989), and would not produce the negative correlation between crabs and clams in the years after the storm. Environmental variability in particular is likely to influence population dynamics of both species, with mortality of *M. arenaria* in warm summers (Glaspie et al. 2018) and mortality of *C. sapidus* in cold winters (Rome et al. 2005). However, the effect of environmental variability is not density-dependent, and would not be expected to act differently on the 2 populations before and after the storm, as seen in this study. Predation remains the most likely explanation for the results obtained here.

Predator–prey models with these 2 species alone were capable of reproducing observations of clam densities and mortality rates, consistent with the idea that blue crabs are a major driver of *M. arenaria* population dynamics (Lipcius & Hines 1986, Seitz et al. 2001, Meise & Stehlik 2003). We propose that *M. arenaria* existed in Chesapeake Bay at high densities until perturbed past the unstable steady state in 1972 by Tropical Storm Agnes. Thereafter, it was able to persist at low density due to the low-

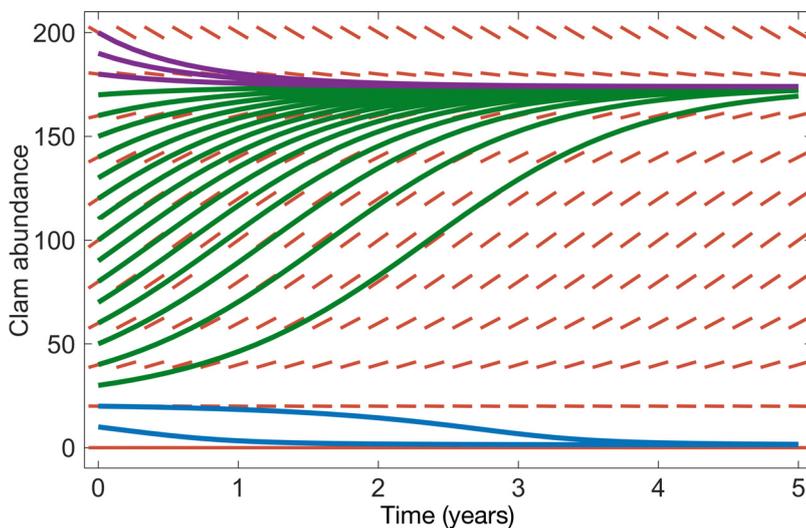


Fig. 3. Slope field diagrams for predator–prey models. Red lines indicate slope field. Trajectories of *Mya arenaria* density either approach a high-density stable steady state at carrying capacity (green and purple lines) or a low-density stable steady state at 1.41 clams m^{-2} (blue lines). Trajectories diverge from an unstable steady state at 20.93 clams m^{-2} (horizontal dashed line). The solid red horizontal line at 0 represents extinction

density refuge from blue crab predation (Lipcius & Hines 1986, Seitz et al. 2001), rather than collapsing to local extinction. This shift to an alternative stable state is different from post-storm declines in seagrass and oysters because it is most likely maintained by density-dependent processes, specifically predation. Unfortunately, *M. arenaria* is unlikely to rebound to high abundance without a beneficial disturbance, such as a considerable recruitment episode or substantial reduction in predation pressure, which propels it above the unstable steady state and concurrently allows it to overcome the exacerbated disease burden.

Extreme weather events are costly, and they are likely to become even more common with predicted increases in the intensity and frequency of extreme events due to anthropogenic climate change (Settele et al. 2014). When examining the cost of extreme weather, ecological impacts are rarely considered, even though the impacts of such events on the ecosystem may be severe (Thomson et al. 2012). Evidence for storm-driven shifts to alternative states in coral reefs (Mumby et al. 2007), kelp ecosystems (Byrnes et al. 2011), and now soft-sediment communities (current study) suggests that management of ecosystems should include an examination of nonlinear interactions and the potential for shifts to alternative stable states.

To our knowledge, this is one of only a handful of examples of agreement between predictions from predator–prey theory and empirical data from natural systems. More evidence is needed to fully evaluate the predictive potential of steady-state analysis involving predator–prey models, and the value of a dynamics systems approach to ecosystem modeling efforts. However, the steady-state analysis approach used here can be adapted to a variety of predator–prey systems with available estimates of life history parameters, population density, and distribution. The approach may also be expanded to include more than 2 species, or encompass additional complexity, such as metabolic functions.

The concepts represented by this crab–clam predator–prey system, including population self-limitation, consumer–resource oscillations, and the functional response, are widespread in nature. These concepts may be considered ‘laws’ of population ecology (Turchin 2001). Each concept listed above introduces nonlinear dynamics into population models, especially those with multiple interacting species. Given the preponderance of multispecies interactions exhibiting nonlinear dynamics, multiple steady states may be a general ecological phenomenon.

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