Spatiotemporal patterns in early life stage winter flounder (*Pseudopleuronectes*

- *americanus***) highlight phenology changes and habitat dependencies**
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- 4 M. Conor McManus^{1,2,*}, Joseph A. Langan², Richard J. Bell³, Jeremy S. Collie², Grace Klein-
- 5 MacPhee^{1,2}, Michael D. Scherer⁴, Richard G. Balouskus¹
- ¹ Rhode Island Department of Environmental Management, Division of Marine Fisheries, Jamestown
- RI 02840
- ² Graduate School of Oceanography, University of Rhode Island, Narragansett RI 02882
- ³ The Nature Conservancy, Narragansett RI 02882
- 10 ⁴ Normandeau Associates, Falmouth, MA 02540
- * [conor.mcmanus@dem.ri.gov;](mailto:conor.mcmanus@dem.ri.gov) (401) 423-1941
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- Running Title: Winter flounder larvae
- Abstract
- Decadal changes in the life history events of marine species are becoming increasingly important
- to identify under a warming climate, yet many long-term monitoring programs do not collect
- data at the spatiotemporal resolution needed to describe them. Such data are vital for
- understanding the southern New England/Mid-Atlantic winter flounder (*Pseudopleuronectes*
- *americanus*) stock, a species hypothesized to be adversely impacted by warming waters via
- increased temporal overlap between its early life stages and predators. To provide insight into
- winter flounder early life history dynamics and context for the stock's resiliency, we examined
- ichthyoplankton data collected from two monitoring programs within Narragansett Bay, RI, USA
- one spatially comprehensive across the Bay proper (2001-2008, 2016-2017), and the other a longer time series centered in one of the Bay's sub-estuaries (Mount Hope Bay, 1972-2017). By
- leveraging these datasets together, we conducted a spatiotemporal synthesis of early life stage
- winter flounder by evaluating changes in larval phenology, decadal coherence in larval spatial
- patterns, and correspondence between larvae and their subsequent life stage. We identified
- changes in larval phenology via earlier seasonal peaks in density through time. Results also
- indicated stable larval spatial patterns during a period of larval decline, as well as spatial
- coherence between larval and young-of-the-year stages. Using winter flounder as a model
- species, our results highlight the importance of high resolution spatiotemporal ichthyoplankton
- sampling to identify changes in phenology and site-fidelity for marine fishes.
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- Key Words: winter flounder, larvae, Narragansett Bay, young-of-the-year, phenology, settlement habitat
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1 1. Introduction

 Long-term ecological datasets provide an opportunity to understand the timing of life cycle events among species and assess how changes in phenology or variability over time may impact ecosystem dynamics. Phenology changes have been documented in many processes important to the function of ecosystems, including alterations in primary production patterns (Ji et al. 2010), as well as shifts in marine species seasonal distributions (Perry et al. 2005, Mueter and Litzow 2008), migration timing (Schindler 2019, Cherry et al. 2013, Langan et al. 2021), molting season (Groner et al. 2018, Daniel et al. 2013), and spawning phenology (Poloczanska et al. 2016). Climate-linked variations in spawning phenology have been documented in marine (Asch, 2015, Fincham et al. 2013, Sims et al. 2004) and anadromous fishes (Bal et al. 2017, Dempson et al. 2017, Kovach et al. 2013), sea birds (Mallory et al. 2010, Gaston et al. 2009, Daunt et al. 2006), and marine mammals (Jemison and Kelly 2001, Rode et al. 2018). In marine fishes, such changes can lead to early life history mismatches for prey availability (Asch et al. 2019, Llopiz et al. 2014) and the diminished presence of requisite physical and environmental habitat requirements for successful development (Flitcroft et al. 2016, Krabbenhoft et al. 2014, Falke et al. 2010).

 Habitat dependencies for the early life history of marine species may be represented as a continuously changing mosaic of multi-dimensional necessities (e.g., prey availability, predator absence, physical oceanographic conditions); even small shifts in a species phenology may have measurable impacts on survivorship and population productivity (Hovel et al. 2019). For many species, specific habitat conditions are critical for the development of early life history stages (Able and Fahay 2010, Collette and Klein-MacPhee 2002). If these conditions deteriorate with environmental alterations or phenological shifts in predator-prey overlap, population bottlenecks

 and declines may result. Therefore, spatial variability in species abundance often highlights essential habitats and abiotic requirements needed to complete their life cycles. Such is particularly true of species inhabiting temperate estuaries, which can experience relatively rapid and extreme environmental changes over short time periods.

 To properly study changes in the phenology and habitat requirements of marine species, spatially resolved long-term datasets are required. Long-term monitoring of fish larvae provides insight into changes in fish populations over time (Llopiz et al. 2014, McClatchie et al. 2014), with applications including understanding interdecadal changes in larval fish assemblage (Morson et al., 2019) and distribution (Walsh et al. 2015), estimating spawning stock biomass (Richardson et al., 2010, 2020), inferring larval habitat requirements (McManus et al. 2018), and discerning variability in recruitment (Payne et al., 2009). However, many of these long-term ichthyoplankton programs do not collect data at spatial scales that allow for assessing fine-scale spatial patterns and their ecological drivers (Brosset et al. 2020) or for developing spatial management measures. Further, few long-term fisheries datasets implement the sampling frequency necessary to detect phenology changes over time (Asch 2015, Staudinger et al. 2019), despite the fact that marine species are particularly prone environmental conditions altering the timing of their life cycle events (Edwards & Richardson 2004).

 As a particularly vulnerable species sensitive to climate change within the Northeast U.S. Continental Shelf ecosystem (Hare et al. 2016), winter flounder (*Pseudopleuronectes americanus*) represent an ideal species to explore environmentally driven changes to phenology and early life history habitat dependencies. A flatfish species once abundant throughout southern New England, the winter flounder population has markedly declined since the 1980s (Bell et al. 2014, 2018). While fishing mortality has been attributed to initiating these declines, the southern

 environments (Scanes et al. 2020), estuarine-dependent species such as winter flounder may be exceptionally vulnerable to phenology changes and the adverse effects of climate change.

 Here, we use winter flounder as an example of a climate sensitive marine fish species to assess early life history temporal patterns and spatial distribution that provide insight into responses to a changing climate. To address these considerations, we collated multiple ichthyoplankton sampling program datasets of varying spatial extent, time series length, and periodicity of sampling from Narragansett Bay, Rhode Island, USA to examine winter flounder early life history. To understand phenology changes in winter flounder larvae, we examined temporal patterns in larval density in the context of the thermal environment. We then characterized the spatial distribution of winter flounder larvae to identify areas of significant spawning in Narragansett Bay and the stability of these spawning patterns across two decades. This spatial assessment was then complemented with an analysis of the spatial coherence between annual larval densities and those of young-of-the-year winter flounder from a juvenile fish seine survey to ascertain whether larval flounder are predictive of subsequent settlement and if this correspondence varies spatially. Through these analyses, we provide further insight into the ecology and population dynamics for this well-studied and historically significant species challenged by climate change as scientists and managers aim to rebuild the stock.

2. Materials and Methods

2.1 Ichthyoplankton Sampling

 Winter flounder larval data were compiled from two ichthyoplankton surveys to understand early life stage spatiotemporal patterns. Mount Hope Bay ichthyoplankton sampling began in 1972 to understand possible adverse impacts to fishes from operation of the Brayton Point power plant's cooling water system (BPE 2016). Annual sampling varied over time, but

 during the larval winter flounder season, sampling always occurred once in February, then 2 weekly $(1972 - 1992)$ or every 4 to 5 days $(1993 - 2014)$ from March through at minimum mid- May. The number of sampling locations also varied over the time series - 23 (1972 – 1979), 7 (1980 – 1985, 1993 – 2014) or 5 (1986 – 1992). Previous analysis of shifting from 23 to 7 sampling stations indicated that the spatial sampling variability was smaller than that of temporal variability. As such, changes in the number of sampling stations over time are not believed to influence inferences on changes in densities or phenology over time. For further details on the stations sampled through time, see BPE (2016). Single oblique tows were made from bottom to surface during the day using a paired 60‐cm diameter bridleless bongo frame for 5 minutes, with the goal of sampling at least 100 $m³$ of water. The primary net mesh used changed from 505-µm to 333-µm in 1995, but dual bongo tows using the two meshes from 1995-2017 were conducted to allow for estimating annual larval extrusion and standardize to a 333-µm density estimate over the time series (BPE 2016). Annual average larval densities prior to 1995 were standardized to 333-µm for a comparison to other surveys that used the same mesh (Supplement 1A). Hereafter these data are referred to as the "Mount Hope Bay survey." A more spatially holistic ichthyoplankton survey of Narragansett Bay was conducted by the Rhode Island Department of Environmental Management (RIDEM) from 2001-2008, and later renewed by RIDEM and the University of Rhode Island's Graduate School of Oceanography (URI-GSO) in 2016 and 2017 (Figure 1). The 2001-2008 RIDEM survey performed biweekly tows for six minutes at 15 stations within Narragansett Bay year-round. In the winter period (January through April), bongo nets were equipped with 335 µm mesh, and 22 from May through December 500 µm mesh. The collected samples were identified to the species

23 level and reported as densities per 100 m^3 of water. The 2016-2017 RIDEM and URI-GSO

 survey closely followed the methods of the earlier study, performing weekly 6-minute tows of 335 µm bongo nets at 15 stations within Narragansett Bay, but targeting only the spawning period of winter flounder (February through April). Fourteen of the stations sampled during the 2016-2017 survey matched those sampled from 2001-2008, with one station relocated from the mouth of Narragansett Bay to Mt. Hope Bay. As described in Bourne and Govoni (1988), both surveys towed nets for 2 minutes each near the bottom, at mid-depth, and at the surface at 7 stations > 11 m depth, and 3 minutes each near the bottom and at the surface at stations ≤ 11 m depth. Field data including date, location, depth, volume filtered by the nets, and surface/bottom water temperature were recorded for each tow. Larval counts were converted to densities (#/100 m^3) using volume filtered estimates. In 2016 and 2017, larval lengths were measured for winter flounder larvae to the nearest tenth millimeter. Hereafter, this 2001-2008 and 2016-2017 dataset is referred to as the "Narragansett Bay survey." Additional details on Narragansett Bay survey ichthyoplankton sampling are found in Langan *et al.* (2020). Analyses were constrained to February through April data to allow for continuity between 2001-2008 and 2016-2017 data (both temporally and plankton net mesh size used), and specifically focus on the winter flounder spawning period.

2.2 Temporal Analyses

 The Mount Hope Bay survey data were examined for long-term seasonal trends in winter flounder larvae with linear regression. The weighted mean week of winter flounder larvae occurrence (*Wt*) was calculated by multiplying a given week of year by its observed larval density, with these values summed over weeks of a given year, and then divided by the sum of the larval density of the entire sampling period:

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W_t = \frac{\sum_{1}^{n} w_{n,t} \times d_{n,t}}{\sum_{1}^{n} d_{n,t}}
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 [1]

1 where $w_{n,t}$ is the week number in year *t*, and $d_{n,t}$ is the mean larval density for that week and year. A seasonal cumulative density curve of larvae was calculated from the weekly larval density data to determine the fifth and ninety-fifth percentiles of *Wt*. The beginning of the larval season was 4 considered the $5th$ percentile of the cumulative density curve and the end of the season was 5 considered the $95th$ percentile. The annual mean week of winter flounder larval occurrence was then regressed against year to determine if the slope was significantly different from zero, and thus whether the annual mean week has changed through time. The time series was examined with the Durbin-Watson statistic to ensure a potential change in mean week through time was not 9 driven by autocorrelation. If the statistic was less than two $(D < 2)$, an autocorrelation term was included in the model and fit with generalized least squares to account for any potential autocorrelation.

 Mount Hope Bay annual average larval densities and the mean week of larvae occurrence were also compared to various temperature metrics to determine if annual variability in larval density or phenology corresponded to the thermal environment. Temperature data used for the analysis were from the concurrent larval sampling conducted in Mount Hope Bay. Previous work has suggested that the influence of temperature on winter flounder may vary with ecological processes and the metrics used to represent those processes (Bell et al. 2014). To comprehensively examine prospective impacts of temperature in winter flounder larvae, we evaluated five thermal metrics for comparison with annual winter flounder larval mean density and occurrence indices: mean annal temperature during the spawning period of approximately 21 February through April (T_{mean}), mean annual temperature when larvae were present ($T_{present}$), 22 annual mean temperature weighted by larval density (T_{weight}) , the annual minimum temperature 23 observed during the spawning period (T_{min}) , and the annual maximum temperature observed

1 during the spawning period (T_{max}) . Winter flounder and thermal time series were compared using linear regressions. Correlations with p-values < 0.05 after Bonferroni correction were deemed significant (Table 1).

 Generalized additive models (GAMs) were used to model the winter flounder larval densities from the Mount Hope Bay survey to understand whether timing or thermal constraints may be a greater determinant for larvae. GAMs use smooth, additive functions to construct non- linear relationships between independent and dependent variables, resulting in curvature or splines in the predictions (Hastie & Tibshirani 1986), and are often used to describe the relationships between fish densities and covariates (Venables & Dichmont 2004). For the Mount Hope Bay survey, concurrently measured temperature at three meters depth and week of the year for the samples were used as explanatory variables for larval density, representing timing and thermal niche drivers. While week of year and sea temperature can be collinear, our focus for this analysis to understand the variables' influence on density rather than predict larval density and risk overfitting concerns. Larval density was calculated as the sum of larval densities across all stations within Mount Hope Bay in each sampling week and divided by the number of stations sampled from Mount Hope Bay during the given period. Similarly, temperature experienced by larvae was calculated as the mean temperature across all sampled stations on a given date weighted by larval density on that date for the Mount Hope Bay dataset. Given the high proportion of zeros in the data, the GAM was run with a Tweedie error distribution using R package 'mgcv' (ver. 1.8-2.2). The Tweedie distribution is a compound distribution that is well suited to data that have both large numbers of true zeros and positive continuous values. *2.3 Spatial Analyses*

Region designations were roughly adapted from Keller et al. (1999), and regional comparisons

 were only conducted where Narragansett Bay larval survey data and RIDEM seine data co- occurred. Seine data annual averages from June through August were used, as these months with the greatest winter flounder presence (i.e. over >86% of winter flounder observed in the survey during these months). Annual larval densities and young-of-the-year abundances were natural log-transformed and compared for individual regions using linear regression. In addition, Bay- wide correspondence between life stages using all regions' data was examined using a generalized linear mixed model to account for fixed sites representing the larger population of the Bay, with region modeled as a random effect. The mixed model was fit with a Gaussian error distribution and implemented using R package 'glmmTMB', ver 1.2.12 (Brooks et al. 2017).

3. Results

3.1 Temporal Analyses

 Annual larval densities from Mount Hope Bay were greatest and highly variable during the 1980s through the 1990s, after which densities declined to low and stable numbers (Figure 2). Annual average densities from the Narragansett Bay dataset over the two decades also showed a decline from 2004 to 2005, and a nearly eight-fold decline from the 2000s to 2010s. Larval densities from Mount Hope Bay sampling standardized to the 333-μm mesh were typically greater than those from the Narragansett Bay data, yet overlapping time series data 18 were not correlated $(R^2=0.15, p-value=0.26)$.

 The weighted mean week of larval occurrence was around mid-March, but in some years was as late as mid-April and became earlier over time. Larvae were present in Mount Hope Bay for four to eleven weeks with a mean of 7.4 weeks (standard deviation = 1.4). Larvae were first 22 detected (defined as the $5th$ percentile) around mid-February, though in some years they did not appear until late March. Larvae tended to be absent from sampling by the end of April, but in

some instances extended into early May $(95th$ percentile; Figure 3). No significant changes were seen over time in the duration of larval occurrence over time. The weighted mean timing of larvae in Mount Hope Bay fit with a temporal autocorrelation term exhibited a significant trend, indicating a shift of larval presence earlier in the year over the time series (slope = -0.055 weeks per year, p-value = 0.005; Figure 3). Through time, larvae became present in the Bay increasingly earlier in the year and disappeared earlier. The Durbin-Watson statistic (1.55, p-7 value = 0.048) indicated autocorrelation (ϕ = 0.27) necessitating the use of generalized least squares regression.

 Weak correlations were observed between annual larval densities and mean week of year and temperature metrics from the Mount Hope Bay survey (Table 1). Annual larval density was negatively correlated with the annual maximum temperature during the spawning period. The mean week of year for larvae was negatively correlated with the annual mean and minimum temperature during the spawning period, but positively correlated with the weighted mean 14 temperature (Table 1).

 In the GAM using Mount Hope Bay survey data, temperature and week of the year exhibited significant relationships with larval density and accounted for 51% of the deviance in larval density. When using only temperature to predict larval density, 31% of the deviance was explained, and using week of year explained 51% of the deviance. The model using both covariates indicated that positive effects on larval density were observed at water temperature 20 below 10° C and from mid-March through early May (Figure 4). A preferred temperature range was apparent, but the temperature range indicated little contrast in preference within this optimal thermal window. The week of year preference indicated a more specified response (Figure 4),

 represented as the strong contrast in week of year response from the GAM, suggesting that the seasonal timing may be a stronger determinant of larval densities than the observed temperature. *3.2 Spatial Analyses*

 Larval densities varied through space and time within Narragansett Bay (Supplement 1B, 5 Figure 5). Annual station average larval densities ranged from 0.29 to 313.16 larvae per 100 m^3 . Larval densities tended to be greatest in northern portions of Narragansett Bay; however, comparing average densities by station across years indicated only stations 9 and 13 were 8 statistically different from each other (One-way ANOVA: F=2.56, p=0.002, d.f=15). The strong decline in larval densities evident in annually averaged data was also apparent between the two decades across stations (Figure 5). While the total winter flounder larval density over Narragansett Bay decreased from the 2000s to the 2010s, the stations of relative importance did not. Across both decades, the general proportion of larvae at each station remained the same (Figure 6). While Densities in the 2010s were lower than those of the 2000s (Figures 5 and 6). Statistically, mean station densities in years 2001-2008 were significantly different than 2016, years 2001-2004 and 2006-2008 were different from 2017, and 2016 and 2017 were different 16 from each other (One-way ANOVA: $F=18.9$, $p<0.001$, d.f=9).

 The correspondence between winter flounder larval densities and young-of-the-year abundance varied spatially within Narragansett Bay (Figure 7). The upper portions of Narragansett Bay, such as the Providence River, Upper Bay, and northern East and West Passages exhibited significant correlations between annual average larval densities and young- of-year flounder (Table 2). Greenwich Bay and southern areas of the Bay (except the lower East Passage) indicated little correspondence between the two life stages. Of the areas with significant life-stage correlations, the Providence River and Upper Bay had higher rates of correspondence

 (or young-of-the-year flounder per larval flounder) than lower regions (northern East and West Passages). Varying from this pattern is the lower East Passage, with the highest rate of larval correspondence to YoY of the Bay's regions (Table 2).

4. Discussion

4.1 Changes in Spawning Phenology

 Over the course of the time series, earlier seasonal timing in peak winter flounder larval densities was found within the Mount Hope Bay sub-estuary (Figure 3). This finding adds to the growing literature on identifying changes in marine fish phenology of life cycle events (Asch 2015, Henderson et al. 2017, Staudinger et al. 2019, Thaxton et al. 2020). Phenological changes in migration for marine fishes has also been observed locally for Narragansett Bay. Using data from a weekly trawl survey, Langan et al. (2021) found that 12 species' residence periods changed by as much 118 days, with those of warm-water species having expanded and those of cold-water species contracted. While migration phenology was not detectable for winter flounder (a known estuary-dependent species with limited migration patterns), changes in phenology may manifest in other life history traits, such as spawning. Previous studies have suggested that warming waters have negatively affected the winter flounder stock through increased predation 17 from a stronger temporal overlap between early life stage flounder and their predators (Taylor & Collie 2003a, Keller & Klein-MacPhee 2011). The observed earlier timing of larvae may be a product of spawning initiated earlier in the year, increased predation on larvae spawning later in the season, or a combination of both. For this predation hypothesis to be true for marine species, it requires the predators' phenology to track that of its prey (Thackeray et al. 2016, Bell et al. 22 2017). The earlier timing of larvae could be impacted by, or have ramifications on, life history traits, such as larval growth. Larval length measurements from the Narragansett Bay survey

 during a mild (2016) and cold (2017) year highlight interannual disparity in larval densities at size through the season (Supplement 1C). Given larval growth can vary with temperature (Houde 2008), it is still to be determined how earlier spawning dynamics or larval presence could be due to, or synergistically interact with, environmentally driven changes in egg incubation and larval growth. Changes in larval dispersal pathways could also influence larval fish phenology, as interannual spatiotemporal larval densities could be influenced by changes in larval transport (DeCelles et al. 2015). However, the life history traits for larval winter flounder favor local retention (Pereira et al. 1999) and are less favorable for distant larval transport. Further hydrodynamic modeling would be needed to assess the degree of which larval transport influences the observed larval phenology.

 Winter flounder larvae have been observed over a range of water temperatures (Pereira *et al*. 1999), with evidence of reduced survival at temperatures greater than 10°C (Able and Fahay 1998). The observed presence of larvae in these colder waters aligns well with previous reports (Figure 4). The thermal window for winter flounder larvae in Mount Hope Bay was similar to that from data over the larger Narragansett Bay system, which indicated a thermal preference of \pm 4-10 \degree C (Supplement 1D). There may also be additional factors beyond water temperature, such as day length and moon phase, that are important for the time of spawning. The actual dates when larvae were present were relatively constrained. Larvae were common by mid-March and were effectively absent by early May. By week 20 (mid-May), it appears that larvae have settled out of the water column or suffered mortality. With the stronger response in the week covariate than temperature in the GAM (Figure 4) and deviance explained results when excluding each variable, seasonal timing appears to be a better predictor of spawning rather than the thermal conditions (Helm et al. 2013).

the thermal environment is likely not the only factor influencing winter flounder larvae,

 particularly given the weak correlations. For example, temperature may be serving as a proxy for other ecological processes, such as predation (Keller and Klein-MacPhee 2000, Taylor and Collie 2003b). Additionally, the decline in larval density tracks similarly to adult or spawning stock trends, suggesting that reproductive capacity for the stock is also influencing observed larval densities.

4.2 Spatial Distribution of Early Life Stages

 Spatiotemporal patterns of winter flounder larval densities emerged for Narragansett Bay. While annual larval densities were variable (Supplement 1B), northern areas of the Bay tended to have greater annual larval densities than lower reaches of the Bay (Figures 5 and 6, Supplement 1D). Historical studies representing annual snapshots of spatial abundance for winter flounder larvae in Narragansett Bay have provided varied findings. For example, winter flounder in 1973 followed a gradient of declining densities toward the Bay's mouth, whereas the same pattern was not observed in 1990 (Keller et al. 1999). Pearcy (1962) noted seasonal progressions in larval density spatially, with those in the Mystic River estuary greatest in the upper estuary earlier in the spawning season, and larval development progression leading to greater abundances in the lower estuary. It is hypothesized that more upstream areas of estuaries may provide ichthyoplankton with regions of reduced predation and increase the likelihood of retention in preferable inshore areas (Barletta-Bergan et al. 2002). The interannual variability in winter flounder larval density found here highlights the challenges in identifying patterns from individual years. For example, some years indicated a similar spatial gradient as found in 1973, while other years resembled the more homogenous 1990 survey (Supplement 1B.) Statistical tests did not identify many larval differences by station over the recent years; however,

 examining the annual densities by site between the two decades highlighted the importance of the upper Narragansett Bay in supporting higher larval densities (Figure 6), and that discerning patterns in larval data with noise can be resolved via aggregation over longer time scales.

 The strong correlations between larvae and young-of-the-year winter flounder in Narragansett Bay highlights the importance of individual spawning sites supporting winter flounder early life history (Table 2, Figure 7.) As noted in other systems, larval- juvenile coherence offers insights for both spawning grounds and nursery areas (Pereira et al. 1999), with these results suggesting spawning locations are important for winter flounder through their first year of life. The importance of larval densities in predicting young-of-the-year abundances later in the season was significant across regions of Narragansett Bay (Table 2), with upper reaches of the Bay often having stronger correspondence between the life stages than southern areas (Figure 9), further highlighting the importance of areas further north in the estuary.

 The high connectivity between larval and young-of-the-year stages for subregions provides additional context for hypotheses on winter flounder subpopulations in Narragansett Bay (Collette and Klein-MacPhee 2002). Tagging studies have suggested that three discrete winter flounder spawning groups exist in Narragansett Bay: Mount Hope Bay, Sakonnet River, and the East Passage/Upper Narragansett Bay (Saila 1962, Powell 1991). Genetic information on Narragansett Bay larval and young-of-the-year winter flounder have indicated that those from the same or nearby locations share genetic signatures (Buckley et al. 2008). These genetic results have suggested 16 distinct subpopulations within the Bay, which is more discrete than tagging studies. The geographic designations presented herein represent a spatial scale in between these two studies yet allows for corroboration of both ideas put forth previously. All East Passage and Upper Bay correlations between life stages were found to be statistically significant (Table 2),

 and physical oceanographic conditions that promote larval retention, earlier settlement, and suite of environmental conditions that improve survivorship. These environmental habitat preferences for early-life winter flounder can provide predation refuge; when conditions are sub-optimal, such as at higher salinities, predators such as striped sea robin and summer flounder can enter these nursery areas (Manderson et al. 2006).

 The spatial heterogeneity in early life winter flounder abundances and connectivity suggest that only certain locations within the Bay may promote growth and survival and be best in describing Narragansett Bay winter flounder metapopulation dynamics. As previously suggested (Yencho et al. 2015), these metapopulation dynamics should be considered in local fisheries management plans. Further analyses on quantifying the factors influencing the settlement rates to the early juvenile stage winter flounder may improve predictions of future year-class strengths for the Narragansett Bay population (Able et al. 2011). Given that the relationship between larval supply and the juvenile stage contributes to the rate and variability in recruitment (Myers and Barrowman 1996), this analysis suggests additional spatial life-cycle analyses across multiple stages are imperative to understand population bottlenecks for Narragansett Bay winter flounder.

5. Conclusion

 The coupling of ichthyoplankton and young-of-year fish monitoring at varying spatiotemporal scales allowed us to provide new insights into a species life history and population structure for a system that has experienced ecosystem changes. Using winter flounder as a model species, we found evidence of an increasing shift toward earlier larval presence, potentially due to changes in spawning phenology and/or predator-prey interactions. Analysis of spatiotemporal larval densities indicated the importance of upper bay habitats. Importantly, this

 research also found high connectivity between larval and young-of-year stages across Bay habitats indicating strong subpopulation dependencies, and likely differences in habitat suitability. The phenological changes and discrete habitat dependencies highlight critical areas of concern for temperate estuary-dependent fish species. Identifying the population bottleneck in declining species will require examination of many factors including physical and environmental spawning triggers, predator-prey match/mismatch theory (Cushing 1990), and assessment of critical habitats for early life history stages, which is only made possible with the appropriate long-term monitoring at fine spatiotemporal scales. Our work highlights the value of such datasets and supports improved monitoring internationally to address these questions endemic to marine fishes.

 The findings presented herein for winter flounder apply globally to species with productivity that continues to decline with ecosystem changes. The prospects of rebuilding the SNE/MA winter flounder stock to past high levels of abundance appear doubtful, as winter flounder population models incorporating future climate conditions project that even fishing moratoria may not suffice to rebuild winter flounder to previous stock levels (Bell et al. 2018). Despite ominous stock-wide outlooks, fisheries managers should not be deterred from developing informed strategies for species negatively impacted by climate. Identifying the habitat dependencies of such environmentally sensitive species that display strong site fidelity and metapopulations, as well as how these dependencies and life histories have changed through time, is critical in constructing effective management tools that protect the species future productivity. Such scientifically informed measures will be paramount in conserving these species vulnerable to environmental change.

6. Acknowledgements

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- ichthyoplankton phenology from the longest continuous larval fish time series on the east

- 2 Table 1. Correlation coefficient (^r) between the Mount Hope Bay survey winter flounder larvae
- 3 and temperature metrics. Winter flounder metrics included annual average density of larvae (D_t)
- 4 and average week of year of larval presence (W_t) . Thermal metrics included mean annal
- 5 temperature during the spawning period (T_{mean}) , mean annual temperature when larvae were
- 6 present (T_{present}), annual mean temperature weighted by larval density (T_{weight}), and both the
- 7 annual minimum and maximum temperatures observed during the spawning period $(T_{min}$ and
- 8 T_{max} , respectively). Bold values represent correlations with p-values < 0.05 after Bonferroni
- correction.
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1 Table 2. Regional correspondence between natural-log transformed winter flounder larval and

2 young-of-the-year indices (n=10 per region). Regions are defined by the larval (L) and seine

 3 (YoY) survey stations used to represent the area. Linear regression slopes (mean \pm standard

4 deviation, R^2 and p-values) from the correlation between regional annual larval and young-of-

5 the-year catch are presented, with significant relationships in bold. Slopes represent the rate of

6 settlement of young-of-the-year densities per larval abundance (YoY/L). The Bay-wide

7 correspondence used all sites' data (n=80), but was performed using a generalized linear mixed

- 8 model with site as a random effect.
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Figure 1. Narragansett Bay winter flounder survey stations: ichthyoplankton (larvae, blue) from

the Narragansett Bay survey and seine (young-of-the-year [YoY], red). The Mount Hope Bay

long-term ichthyoplankton sampling area is indicated by the box. Local regions of the system

referenced in the text are identified. The locus map indicates the study area location within

southern New England, USA.

Figure 2. Annual average larval densities from Narragansett Bay (2001-2008, 2016-2017) and Mount Hope Bay (1973-2017) sampling. Mount Hope Bay samples from 1995-2017 were collected using 333-μm mesh gear; data in years prior were collected with 505-μm mesh but then calibrated to 333-μm equivalent densities based on dual mesh extrusion work (Supplement 1A) for comparison to the Narragansett Bay Survey. Annual average larval densities for the Narragansett Bay survey only included stations that were sampled all years.

Figure 3. Mount Hope Bay survey-derived winter flounder larval phenology calculated as larval density-weighted mean date of occurrence for each year and the period which contains 90% of total abundance (5-95%ile). The trend in the larval density-weighted mean is also presented (blue dashed line; W_t = -0.055 \times year + 123.6 + ϕ). Time periods are expressed as both week of year and date.

Figure 4. Observed Mount Hope Bay winter flounder larval densities (\# per 100m^3) by temperature and week of year (top row), and the estimated relationships between abundance and these covariates from the Tweedie GAM (bottom row). Left and right figure columns represent temperature and week of year, respectively.

Figure 5. Narragansett Bay survey larval densities averaged by station for the two decades: 2000s (left) and 2010s (right). Size of points reflect differences in larval densities (no. larvae per 100m³).

Figure 6. Average natural log-transformed larval densities by station compared between the 2000s (2001-2008) and 2010s (2016-2017). Data points indicate the station number. The dashed line represents the linear relationship between the two periods $(R^2=0.69, p-value<0.001)$.

Figure 7. Spatial correspondence between winter flounder larval densities (Larvae, number per 100 m³) and young-of-the-year abundances (YoY, number per tow) in Narragansett Bay. Data were natural log-transformed prior to correlations. Red lines within panels indicate the linear regressions where the correlations are significant. Regional groupings of stations by survey are presented in Table 2.