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Ecological and demographic drivers of jellyfish blooms

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ABSTRACT: Jellyfish blooms are conspicuous demographic events with significant ecological and socio-economic impact, as they alter aquatic food webs. Despite worldwide concern about an increased frequency and intensity of jellyfish outbreaks, we are challenged to predict their booms and busts. To overcome this issue, we need to identify the ecological drivers of jellyfish blooms by taking into account the complex life cycle of scyphozoans (Cnidaria). Here we present demographic rates of all life stages of the cosmopolitan jellyfish *Aurelia aurita* s. l. within a stage-structured matrix model to investigate the life stage-dynamics of such complex populations under different environments. We illustrate how booms and busts of the medusa stage are highly influenced by non-medusa stage dynamics. We further point out increased food availability as an important ecological driver of jellyfish blooms, as it can shift the population structure of *A. aurita* away from the benthic polyp stage towards more medusae. Comparatively, our projected climate change scenario caused low fluctuations in population density. Overall, our

study reveals ecological and demographic key variables that regulate the intensity and frequency of jellyfish blooms, and thereby contributes to a better understanding of anthropogenic drivers of jellyfish mass occurrence, including habitat eutrophication and climate change.

KEY WORDS: Jellyfish blooms · Biodemography · Global change · Matrix model · Population structure

RUNNING HEAD: Drivers of jellyfish blooms

INTRODUCTION

Jellyfish blooms are demographic events in which the population density of one life stage, the medusa, shows distinct peaks over broad temporal and regional scales (Lucas et al. 2012). Such mass occurrence can have lasting ecological, economic and social consequences by changing the structure of pelagic ecosystems from fish-dominated to jellyfish-dominated food webs (Richardson et al. 2009). Jellyfish directly interfere with human activities; including tourism by stinging swimmers, fisheries by clogging nets, aquaculture by killing fish in net-pens, and power production by clogging cooling-water intake screens (Purcell et al. 2007, Purcell 2012). Concern about a global rise in jellyfish populations is widespread (Sanz-Martín et al. 2016), as jellyfish outbreaks are thought to be favored by several anthropogenic influences such as overfishing, eutrophication, habitat modification, species translocation and climate change (Richardson et al. 2009, Purcell 2012).

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Species involved in jellyfish blooms are primarily found within the cnidarian taxon Scyphozoa and feature complex life histories (Hamner & Dawson 2008). Their life cycles typically include a pelagic, sexually reproducing medusa and a benthic, asexually reproducing polyp stage (Lucas et al. 2012). Such life histories facilitate mass occurrence of the medusa stage and previous studies revealed high phenotypic plasticity of specific life stages under changing environmental conditions (e.g. Hamner & Jenssen 1974, Gröndahl 1989, Liu et al. 2009, Fu et al. 2014). Despite mortality of the planula, polyp and ephyra stage could be of major importance for the development of medusa populations (Lucas 2001), information on the demographic rates involved in the complex life cycles of bloom-forming jellyfish has remained extremely scarce (e.g. Xie et al. 2015). Quantification of demographic rates can identify the life stages that control jellyfish blooms in changing environments. Here, we used Aurelia aurita as model organism to reveal the basic demographic phenomena underlying jellyfish blooms and to point out how such drivers are influenced by environmental conditions. We first parameterized a stage structured matrix model based on data of all life stages (larvae, polyps, ephyrae and medusa) from laboratory experiments and the field. We compared the life history of A. aurita under different food levels with reference to two contrasting Danish water systems – the food limited Kerteminde Fjord/Kertinge Nor and the eutrophic Limfjorden – where zooplankton biomasses are <1-100 µg C l⁻¹ and >10-1000 µg C l⁻¹, respectively (Nielsen et al. 1997, Møller & Riisgård 2007). In Kerteminde Fjord/Kertinge Nor, high density of the local A. aurita medusa population exhausts the available food resources and thereby restricts individual growth to maximum umbrella diameters of only 3-7 cm (Olesen et al. 1994, Nielsen et al. 1997). In Limfjorden, umbrella diameters of A. aurita medusae are in the range of 10-25 cm (Riisgård et al. 2012a, b), as is commonly observed in open coastal areas

(Möller 1980, Schneider 1989). We used our matrix model to evaluate jellyfish population growth rates with respect to changes in survival, stage-transitions or fecundity, i.e., reproductive rates. We further assessed the impact of high and low food conditions on population dynamics in a life-table response experiment (LTRE). Finally, we investigated the effects of climate change by exploring responses in population structure for a projected winter warming trend. Our results highlight the demographic drivers underlying jellyfish blooms and reveal the role of changing food web structures in combination with increased winter temperatures on their intensity and frequency. Such information is essential to predict the impact of anthropogenic influences on the life histories of the cosmopolitan jellyfish *A. aurita* and other jellyfish species with typical boom and bust population dynamics.

MATERIALS AND METHODS

Model organism

The common jellyfish *Aurelia aurita* s.l. (Scyphozoa, Cnidaria) belongs to an ubiquitous, cosmopolitan genus with several species and subspecies (Dawson 2003). *A. aurita* occurs in a variety of coastal and shelf sea environments, particularly in northwestern Europe, the Black Sea, Japan and parts of North America (Lucas 2001). Its complex life history (Fig. 1a) involves pelagic medusae that reproduce sexually and asexually reproducing benthic polyps. After sexual maturation, medusae produce large numbers of planula larvae (Ishii & Takagi 2003), which spend 12 h to 1 week in the water column (Lucas 2001) prior to settlement and metamorphosis into polyps on suitable hard substrate (Keen 1987, Holst & Jarms 2007). Polyps grow and reproduce asexually by several modes, including the production of well protected chitin-covered resting stages called podocysts (Arai 2009), budding of new polyps (Ishii & Watanabe 2003),

and the release of ephyrae by polyps through polydisc strobilation (Lucas et al. 2012). After asexual propagation, polyps return to the polyp stage and the development of ephyrae into sexually mature medusae closes the life cycle (Lucas et al. 2012).

97 **Data collection**

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We cultivated planula larvae, polyps, ephyrae and medusae of A. aurita in the laboratories of the Marine Biological Research Centre, Kerteminde (Denmark), between August 2013 and March 2015. We set up separate experimental series for each life stage using filtered seawater (38 µm) with a constant salinity of 20 PSU and water temperature between 5 and 22 °C. Temperature conditions followed seasonal variability of the local temperate climate zone. We obtained planula larvae from female medusae which we collected in the fjord system Kerteminde Fjord/Kertinge Nor and the major water system Limfjorden (Denmark) in summer 2014. Directly after the release of planulae, we individually transferred larvae to small multi-well containers (n = 960). During daily counts, we determined survival rate and transition rate to the polyp stage by counting the number of living larvae and the number of larvae that had completed metamorphosis into polyps. We continued the counts until all larvae had metamorphosed into polyps or had died. We fed all life stages, except the planula larvae, with low or high amounts (corresponding to a ratio 1:10) of two-day old Artemia salina nauplii and regularly exchanged the seawater in our cultivation containers. We determined a mean organic weight (= AFDW; Vanhaecke & Sorgeloos 1980) of $2.8 \pm 1.3 \mu g \, C$ ind. for each prey organism using the dry weight (DW; 24 h at 60 °C) to ash weight (AW; 4 h at 550 °C) ratio of replicate series with 100 individuals each (n = 10). We adjusted the amount of carbon added through feeding to the body size and resulting energy demands of each life stage. Between summer 2013 and summer 2014,

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we cultivated polyps of A. aurita in the laboratory. For this purpose, we initially added planulae, released by female medusae collected in Kertinge Nor, to small (50 mL) containers with floating polystyrene substrate plates to facilitate settlement. After 24 h, we removed all but one newly settled polyp per substrate plate and replaced the seawater in the 50 mL containers. We subsequently raised and cultivated the individual polyps (n = 100) under high and low concentrations of A. salina nauplii. Polyps of A. aurita have maximum feeding rates of 10 µg C ind. -1 d-1 (Kamiyama 2011); our high and low food levels hence resembled 1 or 10 µg C ind. -1 d-1 ¹, respectively. Each week we counted the number of live polyps as well as the number of buds, podocysts and ephyrae produced by each polyp. To maintain the initial polyp cohorts, we removed all buds detached from "mother" polyps instantly from substrate plates. During the whole experimental period, no excystment of polyps from podocysts was observed. During spring 2014, we individually transferred ephyrae, which were released by polyps under both high (n = 33) and low food level (n = 33), to 250 mL flasks with constant water movement and air supply, and reared them continuously under either high or low concentrations of A. salina nauplii. Food levels for ephyrae corresponded to either minimum energy requirements for maintenance (low food treatment) or a 10-fold carbon load (high food treatment). Based on weekly measurements of inter-rhopalia ephyra diameters, we determined size-specific energy demands to cover minimum energy requirements for maintenance (Frandsen & Riisgård 1997, Båmstedt et al. 1999). We further calculated the number of food organisms offered to each ephyra by converting respiratory maintenance costs into energy demands (1 J = 50 μ l O₂; Eckert et al. 1988), assuming an average energy content of 0.055 J per A. salina nauplius (Bengston et al. 1991). Each week we counted the number of surviving ephyrae and determined the transition from ephyra to medusa; the latter we defined by the presence of fully developed intermediate

lappets and the final closure of adradial clefts. We performed these counts over 5 months until all ephyrae had either died or completed development into medusae. We tracked the newly transitioned medusae for an additional 9 months in weekly time intervals. During autumn 2014, we collected and individually transferred adult, female, larvae-carrying medusae from Kertinge Nor to 1-L aquaria with constant air supply. We kept these adult medusae either under complete starvation (n = 16) or under food concentrations of ~1000 µg C ind. $^{-1}$ d $^{-1}$ (n = 33) by adding 5 daily rations of 2000 *A. salina* nauplii over a 12-h period followed by 12-h without feeding. We stopped daily recording of medusa survival after 4 months, when all individuals in the low food treatment had died; at that time all medusae in the high food treatment were still alive.

Estimating stage-specific rates

We used the experimental data collected in the lab to parameterize age-stage-structured matrix population models. For these models, we used discrete time steps of one month to estimate survival, fecundity, and stage transitions of *A. aurita* between age *x* and age *x*+1. Stages included larvae (L), polyps (P), ephyrae (E) and medusae (M). Since the empirical data was either based on daily (L, M) or weekly records (P, E), we first estimated monthly demographic rates for each life-stage *i* under low and high food conditions, respectively. We determined the fecundity of polyps under low and high food conditions from monthly budding rates and ephyra release rates. We estimated the number of planula larvae carried per adult female medusa (N_L , ind. ind. ⁻¹) during the reproductive period (July to November; cf. Goldstein & Riisgård 2016) from average umbrella diameters (*d*, mm) of 40 mm and 200 mm for low and high food conditions, respectively. In this context, we further used the field observation-based exponential relationship $N_L = 160.8 \times e^{0.029 \times d}$ (Goldstein & Riisgård 2016). From these numbers of planula larvae carried

per medusa, we calculated the daily release of planula larvae (ρ , ind. ind. $^{-1}$ d $^{-1}$) as $\rho = 0.087 \times N_{\rm L}$ (Goldstein & Riisgård 2016). We converted the daily release rates into monthly fecundity of medusae, assuming a sex ratio of 1:1.

To convert any of the daily or weekly transition probabilities and fertilities to monthly rates, we assumed that individuals that died during a given month, had died halfway during this month, i.e., they have lived half the month on average. This assumption is needed since both transition and fecundity rates between subsequent time steps depend on survivorship within each monthly time step, as for instance, a medusa might still release larvae before its death during a given month. Some ephyrae completed transition to the medusa stage within less than a month after they were released by a polyp. We therefore corrected the monthly release of ephyrae by polyps (F_{PE}) not only for daily survivorship of ephyrae, but also for the transition probability to the medusa stage within a given month. Further details on estimating monthly stage-specific demographic rates of *A. aurita* are given in the Supplement.

Monthly stage-structured matrix model

Our matrix model included monthly stage-specific vital rates for each of the 12 month of a year. Age is determined by month, and the final stages are polyp, ephyra, and medusa. Individuals in each life stage have a certain monthly probability to survive in their current stage i (t_{ii}), or to transition to another stage j (t_{ij}). Stage transitions are possible from larva to polyp (t_{LP} , month⁻¹) or from ephyra to medusa (t_{EM} , month⁻¹). Fecundity includes asexual reproduction of polyps by budding (F_{PP} , ind. ind. ⁻¹ month⁻¹) or release of ephyrae (F_{PE} , ind. ind. ⁻¹ month⁻¹), and sexual reproduction of medusae by the release of planula larvae (F_{ML} , ind. ind. ⁻¹ month⁻¹; Fig. 1b). Since larval transition to the polyp stage was very fast (within days or even hours), we linked the

monthly transition from larva to polyp stage to fecundity of medusae ($F_{ML} \times t_{LP}$) to build the basic structure of an irreducible monthly stage-structured population matrix model (Fig. 1c). Using empirical datasets on all demographic rates of A. aurita under low and high food conditions, respectively, we constructed monthly stage-structured submatrices A_x (3 × 3) of the following structure:

$$\mathbf{A}_{x} = \begin{pmatrix} t_{\text{PP}} + F_{\text{PP}} & 0 & F_{\text{ML}} \times t_{\text{LP}} \\ F_{\text{PE}} & t_{\text{EE}} & 0 \\ F_{\text{PE}} \times t_{\text{EM}} & t_{\text{EM}} & t_{\text{MM}} \end{pmatrix}$$

where x denotes the age in months, t_{ij} the transition probability from stage i to j, with the survivorship of stage i expressed as $s_i = \sum_{j=1}^i t_{ij}$ and F_{ij} describes the fecundity of stage i to produce offspring of stage j (cf. Fig. 1c). Note that parameters $F_{PE} \times t_{EM}$ and $F_{ML} \times t_{LP}$ are part of the fecundity and not to be considered for estimating survival s_i . With the 12 monthly submatrices (cf. Figs. S1 & S2 in the Supplement), we constructed yearly population projection matrices A (36 × 36; time step = 1 month), one for each food level:

In some months we observed no survivorship for ephyrae or medusae, which would lead to reducible matrices (Caswell 2001); we therefore included a small transition probability of 10⁻¹⁰ as placeholder (Figs. S1 & S2 in the Supplement). Our results were robust to this assumption.

Demographic parameters and population projection

We analyzed our two (low and high food) population projection matrices using R version 3.1.3 (R Core Team 2015) and compared the demographic dynamics of A. aurita under both food levels. We used R packages 'popbio' (Stubben & Milligan 2007) and 'popdemo' (Stott et al. 2012) to compute population growth rate λ (dominant eigenvalue), stable stage distribution SSD (right eigenvector \mathbf{w} corresponding to dominant eigenvalue), reproductive value R (left eigenvector \mathbf{v} corresponding to dominant eigenvalue), the net reproductive rate R_0 , the generation time, the life expectancy e_0 , and the sensitivity of λ to perturbations of the matrix elements (only elements >0 are considered). We further used the two matrices to perform life-table response experiment (LTRE) analysis (Caswell 1996, Tuljapurkar & Caswell 1997) with focus on the sensitivity of λ to perturbations in stage-specific transitions when comparing high to low food levels.

Adjusting matrix parameters with respect to population growth rates

Our matrix model for low food conditions resulted in a population growth rate λ of 1.32 month⁻¹. Such growth rates are not realistic, at least not over extended periods in the field. The unrealistically high λ reflects our sheltered laboratory environment under which we collected the data to parameterize the models. As an important microzooplankton component, planula larvae are highly subject to predation under natural conditions. We therefore reduced the transition rates

from larva to polyp of the two annual population matrices to a level that resulted in a stationary population ($\lambda = 1 \text{ month}^{-1}$) under low food conditions. We applied the same calibration to the high food condition, leaving us with a still rapidly growing population under high food conditions.

Population projection in a climate change scenario

We altered the population matrices to investigate potential effects of climate warming on the population growth, stable stage distribution, net reproductive rate and life expectancy of *A. aurita*. Increased water temperatures from 5 to 10 °C during winter are predicted to benefit the production of *A. aurita* ephyrae by extended strobilation periods and enhanced ephyra release per polyp (Holst 2012). We established such a warming scenario by increasing ephyra production 1.8-fold during the entire strobilation periods (cf. Tables S1 & S2 in the Supplement). We additionally extended strobilation periods by one month (i.e., January or February for high and low food conditions, respectively). With these modified matrices we investigated the demographic effects of a predicted winter warming trend on *A. aurita* populations under low and high food conditions.

238 RESULTS

Demographic dynamics of Aurelia aurita under low and high food levels

The metagenic life cycle of *Aurelia aurita* (Fig. 1a) was resembled by cyclic (imprimitive)

structure of our matrix model, expressing annual peak and pit dynamics in projected population density (Fig. 2). The calibrated population growth rate under 10-fold increased food levels was λ = 1.63 month⁻¹, indicating a 345-fold yearly increase in population size compared to the

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calibrated stationary stable population under low food conditions ($\lambda = 1 \text{ month}^{-1}$). Similar to low food levels, the projection matrix showed cyclic annual structure with typical peak and pit dynamics in projected population density, but amplitudes of cycles were much increased under high food availability (Fig. 2a). Under low food conditions, population density increased to a peak during spring and summer and thereafter decreased to low densities during fall and winter. Under high food levels, peak density and dynamics differed significantly compared to low food conditions, by showing two distinct maxima in population density, one during spring and one during autumn, each followed by a subsequent minimum in summer and winter (Figs. 2a-b). The convergence time to stable stage, i.e. the damping ratio, of our modelled A. aurita population was 59 months (~5 years) under low food conditions and only 3 months for the high food population. The stable stage distribution under low food availability was vastly dominated by polyps (~0.94) that survived throughout the year. Ephyrae and medusae accounted for comparatively small fractions of the population (0.04 and 0.02, respectively) and showed typical seasonal patterns (Fig. 2b). These included the release of ephyrae during April/May and their survival into September. Only few ephyrae transitioned to the medusa stage under low food conditions, so that medusae were present from June to December and contributed only marginally to restocking polyp densities by the release of planula larvae during autumn and winter. Under high food supply, the stable stage distribution of the A. aurita population was characterized by less polyps (0.76) compared to low food conditions. Medusae played a more pronounced role (0.2) under high food levels and ephyrae accounted for a small fraction of the overall population (0.04) due to faster transition to the medusa stage. Compared to a year-round standing stock of polyps under low food conditions, the stable stage distribution under high food levels was characterized by

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lower proportions of polyps, which were predominant from September to March. Ephyrae were released in March/April, and medusae dominated the A. aurita population from April to June under high food levels. In autumn, the release of planula larvae restocked the polyp fraction (Fig. 2b). Our findings illustrate a shift from polyp-dominated A. aurita populations towards more medusae under increased food conditions (see Table S3 in the Supplement). The reproductive value describes the contribution of an individual in a given stage to future generations. It was low for ephyrae (0.03) under low food conditions, and sexual reproduction of medusae contributed about half as much (0.42) compared to the asexually reproducing polyp stage (1; the reproductive values are scaled for January). The importance of the medusa stage under high food conditions was reflected in a reproductive value that increased to 0.27 for ephyrae and to 9.13 for medusae (note scaling to 1 for polyps in January). A net reproductive rate of 1 indicated that each polyp is replaced by 1 new polyp at the end of its life under low food conditions which resulted in long generation times of (200 months) ~16.7 years. Under high food levels, the net reproductive rate increased to 57 polyps replacing each polyp at the end of its life, highlighting the production of increased numbers of offspring compared to low food conditions. Due to this high net reproductive rate, high food conditions also boosted the average time between two consecutive generations to 8 months. Life expectancy of polyps, estimated from the fundamental matrix as the mean time to death, was 33 years under low food conditions, while ephyrae and medusae revealed much shorter life expectancies of 2 and 3 months, respectively. In consensus with a more pronounced impact of the medusa stage on population structure under high food levels, life expectancy of polyps

decreased to 7 years, while the length of ephyra and especially medusa lives extended to 3 and 8 months, respectively (cf. Table S3 in the Supplement).

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Population growth λ under low food conditions was most sensitive to changes in polyp survival $(s_P = t_{PP})$, with constantly high sensitivity of 0.08 throughout the months of the year (Fig. 2c). Sensitivity analysis further highlighted a lower influence of fecundity on growth rates compared to survival rates. We detected sensitivities of λ between 0.01 and 0.02 for perturbations in ephyra production (F_{PE}) from March to May and of 0.01 to 0.05 for perturbed transition probability from ephyra to medusa $(t_{\rm EM})$ from March to June. Under high food conditions, the estimated sensitivity of λ was generally higher than under low food levels. The population growth rate was most sensitive to changes in medusa survival, with sensitivity remaining high, in a range of 0.07 to 0.16 from April to October, and reaching a maximum of 0.32 from June to July when medusa survival decreased. Changes in polyp survival affected λ particularly from October to April, with sensitivities ranging from 0.04 to 0.13. To investigate differences in the sensitivity of λ between food treatments, we performed a life table response experiment (LTRE) analysis. Our results showed a maximum sensitivity of 0.14 for the transition from ephyra to medusa stage in April/May, followed by high sensitivities of 0.08 for medusa survival in the same time period. We further observed high sensitivities of 0.05 to 0.07 for combined larva production and subsequent transition to the polyp stage from August to December (Table S4 in the Supplement).

Effects of climate change

For our winter warming scenario with extended strobilation periods and increased ephyra release, population growth rates of *A. aurita* increased to 1.03 month⁻¹ and 1.65 month⁻¹ under

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low and high food conditions, corresponding to relative monthly increases of 2.5 and 1.3 %, respectively. The projected winter warming trend enhanced the variability in inter-annual population structure in both food treatments. We also observed slightly more distinct peak and pit dynamics in population density (Fig. 2a). The stable population structure of A. aurita we predicted for winter warming shifted slightly towards ephyrae (0.08) and medusae (0.03), with polyps still dominating (0.9) under low food levels (Fig. 2b). In relative terms, we observed 1.2 % more medusae, 3.8 % more ephyrae and 5.0 % less polyps per year. Under high food levels, our increased winter temperature scenario shifted the stable stage distribution to 9.9 % more medusae, 2.3 % less ephyrae and 7.6 % less polyps (see Table S5 in the Supplement). As a consequence of an earlier onset of strobilation periods in the scenario, fast transition of ephyrae resulted in significant proportions of medusae already present in February and March under high food levels (Fig. 2b). Our winter warming scenario further increased reproductive values in both ephyrae and medusae by enhancing their relative contributions to the number of offspring of future generations by 4.3 % and 0.2 % per year under low and by 5.1 % and 2.8 % per year under high food levels, respectively. We found that winter warming increased the net reproductive rate by 1.35 polyps replacing every polyp at the end of its life under low food conditions, which corresponds to a relative difference of 35 %. Under high food conditions, we found a net reproductive rate of 62, indicating a relative increase by 8 % as a result of higher winter temperatures. The generation time under low food conditions was shortened to 12.3 years, corresponding to a difference of 4.4 years made up by the projected winter warming trend, while generation times were only marginally shortened by 2 days under high food conditions. Winter warming increased life expectancy of ephyrae by 12 day and 19 days, respectively (Table S5 in

the Supplement), whereas life expectancy of polyps and medusae remained unchanged under both low and high food levels.

DISCUSSION

We reveal the demographic dynamics and life history shifts underlying jellyfish blooms and show that the intensity and frequency of jellyfish booms and busts is subject to specific ecological triggers. We illustrate that the typical peak and pit dynamics of jellyfish populations are significantly enhanced by natural variations in the food regime, while other ecological changes, such as a predicted winter warming, has a comparatively low impact on the development of jellyfish blooms. Despite our winter warming scenario causing substantial changes in population stage structure and dynamics, it is not sufficient to shift the predominant mode of reproduction from asexual to sexual. We suggest this latter shift from polyp- to medusadominated populations as a key mechanism causing jellyfish outbreaks. Our findings support several previous studies which have identified metagenic life cycles as the base for boom and bust population dynamics (Hamner & Dawson 2008, Pitt et al. 2008, Brotz et al. 2012). Since jellyfish blooms have been suggested as indicators for ocean degradation (Schrope 2012), the present insights may provide important basic knowledge regarding the health of ecosystems worldwide.

The demographic dynamics behind jellyfish blooms

Our findings suggest food availability as a major constraint for the intensity and frequency of jellyfish blooms, confirming recent observations that have pointed out food supply as limiting factor for the development of ephyrae (Fu et al. 2014) as well as the survivorship and fecundity

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of medusae (Goldstein & Riisgård 2016). Further, prey removal has been suggested as highly efficient management strategy in reducing the population size of cubomedusae (Bordehore et al. 2015). Yet, to our knowledge, we present the first comprehensive quantitative study of how increased food availability can trigger demographic changes in Aurelia aurita populations. Such changes include sharpened seasonal dynamics and shorter generation times of the pelagic medusa stage. We show that jellyfish blooms are a natural consequence of complex life cycles and confirm the key role of benthic polyps in ensuring long-term survival of A. aurita populations (Boero et al. 2008, Lucas et al. 2012), particularly when food is scarce. Increased food levels are associated with boosted population growth and dramatic life history shifts in A. aurita, as expressed by longer-lived medusae with enhanced release of planula larvae, increased ephyra production, as well as faster and more successful development of ephyrae into medusae. Our study provides important basic knowledge on the consequences of habitat eutrophication in context with jellyfish blooms, as investigated food levels are motivated by eutrophic to hypereutrophic conditions (cf. Riisgård et al. 2008) which are observed in our study areas Kerteminde Fjord/Kertinge Nor (Olesen et al. 1994, Nielsen et al. 1997) and Limfjorden (Møller & Riisgård 2007, Riisgård et al. 2012b). Indicated by our life table response experiment analysis, which showed high sensitivities for the fecundity of medusae combined with the transition from larvae to polyps, an increased natural mortality risk is especially likely for the planktonic planula stage of A. aurita (cf. Lucas 2001). A transition probability of 0.001 month⁻¹ from larva to polyp matches well previous estimates (Xie et al. 2015) and was used to calibrate our presented matrix models for the A. aurita population under low and high food conditions. Our resulting stable population under low food levels is comparable to the local A. aurita jellyfish population in the semi-enclosed fjord

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system Kerteminde Fjord/Kertinge Nor, which has remained unchanged over the last 24 years (Olesen et al. 1994, Nielsen et al. 1997, Riisgård et al. 2010, Goldstein & Riisgård 2016). The rapid growth potential we show for our calibrated jellyfish population under constantly high food (optimum) conditions however is in contrast to rather low abundances of A. aurita medusae observed in open coastal waters (Möller 1980, Goldstein & Riisgård 2016) or in Limfjorden (Hansson et al. 2005, Møller & Riisgård 2007), where A. aurita may currently experience the process of being outcompeted by the invasive ctenophore *Mnemiopsis leidyi* (Riisgård et al. 2015). This implies additional factors regulating polyp, ephyra and medusa density in less protected ecosystems. According to our model for high food levels, an increased polyp mortality of 0.3 d⁻¹ (Xie et al. 2015; i.e., a survivorship of 2.3×10^{-5} month⁻¹) would decrease population growth rates to 1.44 month⁻¹. Reduced medusa survival to 0.1 month⁻¹ (corresponding to a mortality of 0.07 d⁻¹; cf. Xie et al. 2015, Table 1 therein) or alternatively, a lowered transition probability from ephyra to medusa stage to 0.0001 month⁻¹, would result in remarkably decelerated population growth rates of $\lambda = 1.10 \text{ month}^{-1}$ and 1.09 month⁻¹, respectively. Since population growth of A. aurita is less dependent on the survival of benthic polyps under increased food levels, our estimates suggest that transition from ephyra to medusa stage and survival of medusae are the most critical parameters for the development of jellyfish blooms in exposed coastal regions. This finding is additionally supported by the results of our life table response experiment analysis. Besides food availability, intra- and inter-specific competition, predation (Kakinuma 1975, Hernroth & Gröndahl 1985, Hansson 1997, Arai 2005), substrate availability, hydrodynamic currents (Johnson et al. 2001, Xie et al. 2015) and extreme environmental conditions (Lucas et al. 2012) should be considered as important constraints for the development of jellyfish blooms.

In particular, population density-dependent mechanisms controlling individual size (Schneider & Behrends 1994, Lucas 2001, Riisgård et al. 2010, Goldstein & Riisgård 2016) could further play a major role for the demography of *A. aurita*. Especially size-structure should hence be taken into account in future population models (cf. Bordehore et al. 2015) to evaluate the importance of individual density for the development of jellyfish blooms.

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Climate change and jellyfish blooms

Warm temperature can lead to enhanced asexual reproduction and has therefore been associated with greater numbers of medusae for most temperate scyphozoan species (Purcell 2005). Our findings do not support rather qualitative arguments about increased water temperature boosting the boom and bust dynamics of jellyfish blooms (Richardson et al. 2009, Xie et al. 2016). At least winter warming has comparatively weak effects on the growth of A. aurita populations due to resulting seasonal changes in population composition and structure. Our winter warming scenario led to only slightly enhanced seasonal variability in population structure which involved changes towards less polyps and more medusae but did not cause a shift from asexual polyp- to sexual medusa-dominated A. aurita populations as observed under increased food conditions. Winter warming however can increase the population growth of temperate jellyfish populations, which may be especially pronounced for populations in regions with limited food availability. As a consequence of enhanced strobilation periods and increased numbers of ephyrae released per polyp (Holst 2012), warmer winter temperatures shorten generation times due to more medusae with increased reproductive value. Climate warming has previously been suggested to benefit the population size of temperate scyphozoan species, but not tropical or boreal species (Holst 2012, Purcell et al. 2012). Based on recent climate analyses, rises in sea surface temperature by 5 °C

are expected within the next 100 years (Belkin 2009). Our results indicate a maximum increase in jellyfish density by 1.1 during this warming period, while long-term observations in Kertinge Nor have shown that fluctuations by a factor of up to 1.6 are common among years (Olesen et al. 1994, Riisgård et al. 2010, Goldstein & Riisgård 2016). Further taking into account the requirement of a cold trigger to stimulate strobilation in *A. aurita* (Holst 2012), we suggest that winter warming has marginal effects compared to regionally observed increases in food level.

CONCLUSIONS

Our study highlights that food availability drives and specifically shapes the booms and busts of jellyfish blooms. We show that predicted changes in water temperature likely cause much less dramatic increases in the density of *Aurelia aurita* medusae than already established food concentrations in several eutrophic regions around the world. The importance of natural competitors and predators for keeping jellyfish blooms in check needs further exploration, but our findings might serve as a promising starting point. We conclude that a combination of environmental triggers, such as habitat eutrophication, overfishing, artificial settlement substrates and climate change, can promote jellyfish outbreaks severely. Our findings emphasize the fundamental importance of an integral, quantitative view on jellyfish life histories. We believe such inclusive knowledge, comprising species-specific, life stage-specific and age-specific traits, is an inevitable prerequisite in providing perspective for future management of jellyfish blooms.

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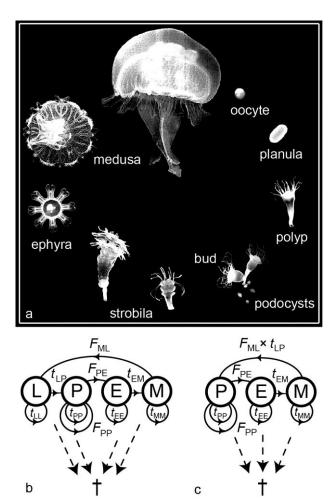


Fig. 1. Life history of the common jellyfish *Aurelia aurita* (Scyphozoa, Cnidaria). (a) Metagenic life cycle with alternation of generations between the sexually reproducing medusa and the asexually reproducing polyp stage. (b) Life cycle graph for *A. aurita*, including transition probability *t*, fecundity *F* (solid arrows) and mortality † (broken arrows) of all life stages, i.e., planula larva (L), polyp (P), ephyra (E) and medusa (M). (c) Reduced life cycle graph for polyps, ephyrae and medusae considering fast (daily) transition from larva to polyp stage.

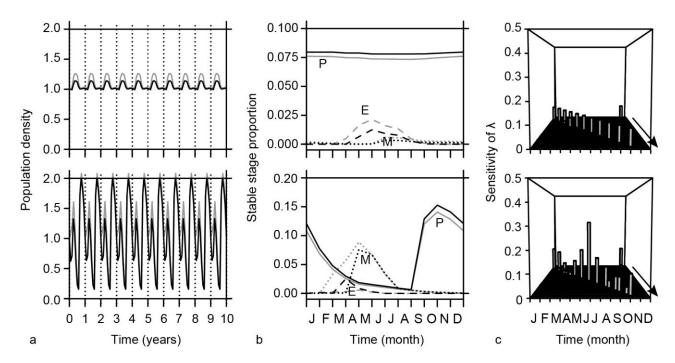


Fig. 2. Population matrix model for *Aurelia aurita* under low (upper panels) and high food (lower panels) conditions (1:10). (a) Matrix projection over a 10-year period (dotted lines), standardized to the long-term effects of population growth rates $\lambda = 1$ month⁻¹ and $\lambda = 1.63$ month⁻¹, respectively, for present temperature conditions (solid black lines) and for a predicted winter warming trend (solid grey lines). (b) Stable stage proportion of polyps (P), ephyrae (E) and medusae (M) for present temperature conditions (black lines) and for a winter warming scenario (gray lines). (c) Sensitivity of λ to changes in monthly stage transitions.