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Chamelea gallina growth declined in the Northern Adriatic Sea during the Holocene Climate Optimum

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The fossil record of past climate transitions offers insights into future biotic responses to climate change. Here, we compare shell growth dynamics, specifically linear extension and net calcification rates, of the bivalve *Chamelea gallina* between Northern Adriatic Sea assemblages from the Holocene Climate Optimum (HCO, 9–5 cal. kyr B.P.) and today. This species is a valuable economic resource, currently threatened by climate change and numerous anthropogenic stressors. During the HCO, regional sea surface temperatures were warmer than today, making it a potential analog for exploring ecological responses to increasing seawater temperatures predicted in the coming decades. By combining standard aging methods with reconstructed sea surface temperatures, we observed a significant reduction in linear extension and net calcification rates in warmer HCO assemblages. During the HCO, immature *C. gallina* specimens developed a denser shell at the expense of a linear extension rate, which was significantly lower than modern specimens. This resulted in an average delay of 3 months in reaching sexual maturity, which is currently reached after 13–14 months or at a length of ~18 mm. This study sheds light on the natural range of variability of *C. gallina* over longer time scales and its potential responses to near-future global warming.

Keywords Bivalvia, Adriatic Sea, Growth dynamics, Climate change, Fossil assemblages

Studies on the capacity of marine organisms to acclimatize or adapt to temperature changes have increased during the past few decades as the severity of climate warming has been recognized^{1–6}. Some coastal areas have experienced changes in mean seawater temperature, in some cases equal to or greater than the conditions originally predicted by future ocean scenarios^{1–5}.

The venus clam *Chamelea gallina* has been recently used as a model organism for evaluating adaptation responses to near-future environmental changes^{7,8}. This species is abundant in shoreface settings at depths ranging from 0 to 12 m up to 1–2 nautical miles in the Mediterranean Sea⁹. It is particularly abundant in the western Adriatic Sea (Mediterranean Geographical Subarea—GSA 17), where the massive outflow of the Po (along with that of other minor rivers) and the anticlockwise currents in the North Adriatic Sea (NAS) provide, along the Italian coasts (and particularly around and south of the Po Delta), abundant nutrients, particles, and organic matter¹⁰. Moreover, *C. gallina* represents an economically and ecologically important species in this basin, and its harvesting in GSA 17 accounts for over 98% of 'venus clams' production at the national level¹¹. Previous studies have mainly focused on population dynamics, shell growth, and composition of this species in living populations of the western Adriatic coasts^{12–16}. It has been observed that the shell properties of *C. gallina* populations are influenced by environmental changes, with variations in shell density, thickness, and growth primarily related to varying water temperatures and secondarily to salinity and food availability along a spatial gradient^{12,13}. Along the latitudinal gradient in the Adriatic Sea, shells of the warmest populations were characterized by lighter, thinner, more porous, and fragile morphotypes¹². However, a solid understanding of long-term biotic dynamics in the context of major environmental shifts is required to reduce the uncertainty regarding organismal responses to ongoing and near-future climate (and derived environmental) changes. Nevertheless, detecting long-term population dynamics in field-based ecological studies can be time-prohibitive.

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Temporally well-resolved (i.e., low time-averaged) fossil-rich assemblages retrieved in sedimentary successions might record ecological responses to past climatic-driven environmental shifts, far beyond the limited timescales of direct ecological monitoring, typically restricted to the most recent decades^{17–23}. Given their extensive fossil record, marine calcifying organisms are unique recorders of past environmental changes^{24–29}. In particular, bivalve shells record historical data due to their seasonal deposition of carbonate material, retaining high-resolution long-term temporal records of the ambient physical and chemical conditions during growth^{30–32}. *C. gallina* holds significant, yet largely untapped, potential in this regard. Despite its ecological and economic importance, the use of *C. gallina* dead assemblages to reconstruct millennial-scale intraspecific dynamics in relation to climate change remains virtually unexplored. To date, only one study (Cheli et al.,³³) investigated the long-term variability in shell microstructural features—specifically, bulk density, microporosity, and micro-density—in response to past climate-driven environmental changes, comparing Middle and Late Holocene assemblages with modern counterparts from the NAS. That study found that modern assemblages displayed less dense and more porous shells than fossil ones, likely due to lower present-day aragonite saturation states. While Cheli et al.³³ focused on shell microstructure and mineralogical properties, the present study uniquely investigates growth performance metrics—specifically, linear extension and net calcification rates—during the Holocene Climate Optimum (HCO) and today, in the NAS south of Po Delta. Here, we aim to evaluate how warming temperatures during the HCO, a potential analog for near-future climate conditions, affected growth dynamics compared to the present day. To this end, we estimated linear extension and net calcification rates on four of five shoreface-related *C. gallina* assemblages previously investigated in Cheli et al.³³: two from the present-day NAS (namely MGO and MCE; Fig. S1) and the two from the HCO (i.e., CO1 and CO2, respectively dated 7.6 ± 0.1 ky B.P., and 5.9 ± 0.1 ky B.P.; Fig. S1), when regional sea surface temperatures were higher than today (see supplementary material “Environmental Parameters”).

This study addresses the knowledge gap concerning the natural range of variability of *C. gallina* at temporal scales beyond those typically considered in ecological monitoring, providing valuable insights into the potential responses of *C. gallina* to near-future global warming.

Results

Specimen age-length von Bertalanffy growth curve

In all examined assemblages, there is no significant difference in the extrapolated ages of fossil and modern *C. gallina* specimens between estimations based on external and internal rings (Mann-Whitney *U* test, $p > 0.05$). Moreover, Kimura’s likelihood ratio tests indicated no significant differences between the growth curves obtained from the external and internal rings. Therefore, a generalized VBGF curve was obtained for each assemblage using the combined data (Fig. 1; Fig. S2). Measurements of stable oxygen isotopes ($\delta^{18}\text{O}$) validated the extrapolated ages from the aging methods by fitting the VBGF curves (Fig. 1).

Differences in the observed values for the estimated mean asymptotic specimen length (L_{inf}) (anterior-posterior) and the growth constant (k) in the targeted assemblages were not statistically significant (Kruskal-Wallis test, $df = 3$ and $p > 0.05$; Table 1). Shell length and estimated life span were comparable among assemblages for mature and all shells (Kruskal-Wallis test, $df = 3$ and $p > 0.05$; Table 2). The age at the onset of sexual maturity resulted higher in the warmer HCO assemblage (i.e., +28% in CO1) compared to modern ones, thus showing a delay in reaching sexual maturity in the former (Kruskal-Wallis test, $df = 3$ and $p < 0.05$; Table 2).

Specimen linear extension and net calcification parameters

In each assemblage, the examined shell growth parameters (i.e., linear extension and net calcification rates) were negatively correlated with shell age (Fig. 2). Linear extension and net calcification rates differed significantly among assemblages (Table 2; Fig. 3). Bulk density was significantly different for immature specimens and hence for the whole dataset among assemblages (Table 2; Fig. 3; Fig. S3). The variation of the two above-mentioned growth parameters was then analyzed in relation to temperature-driven changes between the HCO and the present day. Linear extension rate and net calcification rate showed a significant but moderate negative correlation with SST in the whole dataset ($r_s = -0.311$ and -0.292 , respectively, $p < 0.001$; Fig. 3). Correlations with the SST were also performed separately for immature and mature specimens, with both sub-datasets exhibiting a significant negative correlation for linear extension and net calcification rates (Fig. 3). The two assemblages positioned at the highest and lowest temperature range over time (i.e., CO1–18.6 °C and MGO–17.1 °C) displayed stronger variations in immature specimens, with a 2.3 mm (~18%) per year difference in mean linear extension rate and a 0.47 g/cm² (~15%) per year difference in mean calcification rate and (Table 2; Fig. 3).

Discussion

The radiocarbon analyses of *C. gallina* specimens revealed that fossil and modern dead assemblages have a comparable time averaging (i.e., the cumulative amount of time in which individuals forming an assemblage have lived). Radiocarbon dating of seven *C. gallina* specimens from a nearby shoreface dead assemblage showed an interquartile range (IQR) of 50 years¹⁷. Along the same line, the radiocarbon dating of eleven *C. gallina* specimens from sample CO1 resulted in an IQR of 85 years (Table S1). This suggests that the fossil and modern dead assemblages have accumulated over similar time spans (centuries). As a result, the trends in biomineralization observed in the analyzed assemblages reflect long-term, mediated responses to environmental changes and buffer the results from possible short-term or uncommon environmental perturbations that could have affected a specific cohort of the *C. gallina* population.

As for *C. gallina* growth parameters retrieved in this study, and despite the variation in temperature between the Holocene Climate Optimum (HCO) and present-day conditions, the maximum expected length of the population (L_{inf}) and the rate at which it is approached (k) do not show significant changes between fossil and

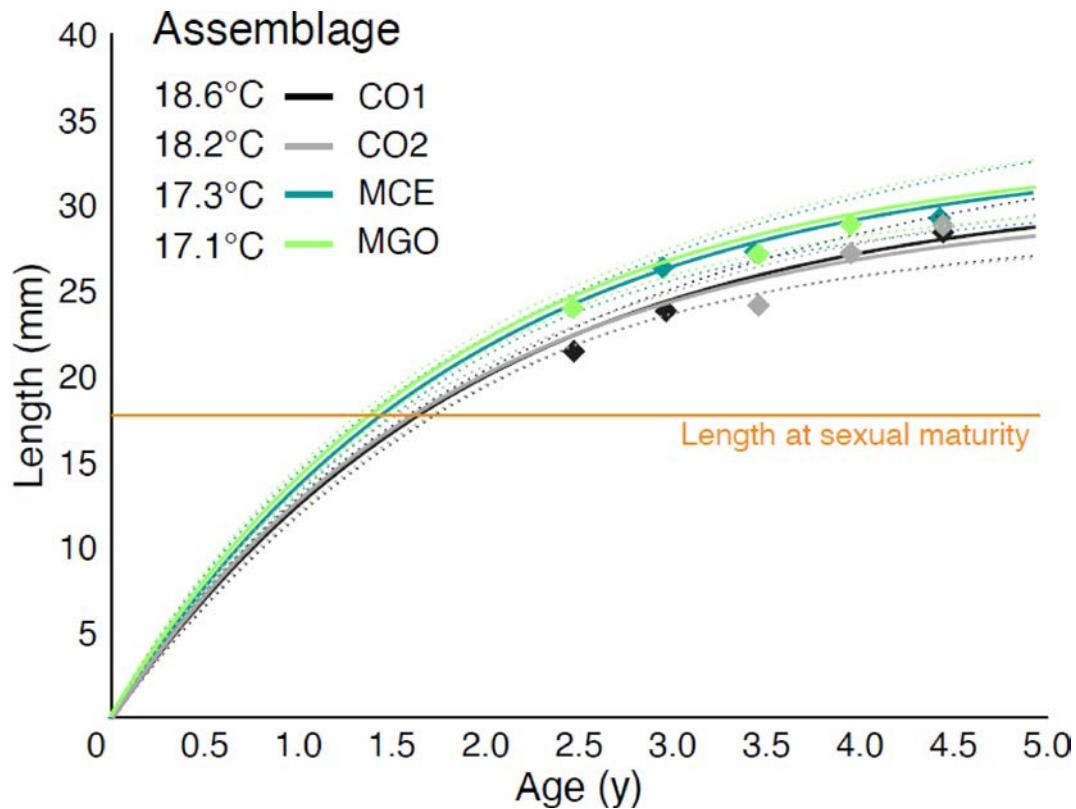


Fig. 1. Von Bertalanffy growth curves. The generalized age-length curve for each assemblage is presented alongside the specimen lengths associated with maturity. The von Bertalanffy curves were created by merging data from the two aging methods (counting external and internal rings). Diamonds represent the ages obtained from the $\delta^{18}\text{O}$ profiles along the shell growth axis and validate the counting aging methods by fitting the curves.

Assemblage	n	^{14}C Age (ky B.P.)	SST (°C)	Generalized VBG function			
				L_{inf}	CI	k	CI
CO1	49	7.6 ± 0.1	18.6 ± 0.4	31.67	29.03–34.92	0.51	0.42–0.59
CO2	59	5.9 ± 0.1	18.2 ± 0.3	30.66	28.67–32.86	0.54	0.47–0.63
MCE	68	modern	17.3 ± 0.1	33.62	30.81–37.04	0.53	0.44–0.62
MGO	73	modern	17.1 ± 0.1	33.07	31.07–36.78	0.54	0.45–0.63
K-W			***	NS		NS	

Table 1. Calibrated radiocarbon age, sea surface temperature (SST), and von Bertalanffy growth function (VBGF) parameters. Values for each assemblage in chronological and temperature order. Radiocarbon measurements are reported in years 'before present' (B.P.); HCO temperature estimates derive from alkenone data collected in the Gulf of Lion (refer to supplementary material for further details). L_{inf} and k were estimated from the generalized VBG function by pooling the data for external and internal rings. N = number of collected specimens. K-W = Kruskal-Wallis rank test; NS not significant, *** $p < 0.001$. Assemblage labels, number of right valves examined (n), calibrated ages ^{14}C Age), and Sea Surface Temperature (SST) are after³³.

modern assemblages (Fig. S2; Table 1). The estimated mean asymptotic shell lengths (L_{inf}) for fossil and modern assemblages were lower than those reported in previous studies for modern living populations of *C. gallina* from the Adriatic nearshore settings^{13,15,34–37}. However, considering the lower confidence interval for L_{inf} observed in the sites around the Po Delta, the modern shells from this study were comparable with the modern ones from Mancuso and collaborators²⁹. The observed values of von Bertalanffy growth constants (k) for all assemblages were consistent with nearly all previous data obtained in the Adriatic Sea^{13,34,38}.

A lower L_{inf} but the same k suggests that while individuals in one population will not grow as large as those in another, they grow at a comparable rate during their life span. This could be influenced by a combination of environmental, genetic, and ecological factors that limit the maximum achievable size without affecting the intrinsic growth rate. On a decadal scale, the specimens of the present study might have lived in conditions of

	Assemblage	n	Length (mm)	Age (y)	Bulk density (g/cm ³)	Linear extension rate (cm/y)	Net calcification rate (g/cm ² y)
Immature shell	CO1	24	12.44 ± 0.79	1.02 ± 0.16	2.45 ± 0.03	1.26 ± 0.02	3.07 ± 0.04
	CO2	31	12.08 ± 0.58	0.95 ± 0.12	2.52 ± 0.02	1.30 ± 0.04	3.29 ± 0.05
	MCE	38	11.54 ± 0.57	0.82 ± 0.10	2.47 ± 0.01	1.44 ± 0.04	3.56 ± 0.05
	MGO	38	11.45 ± 0.62	0.80 ± 0.11	2.39 ± 0.03	1.49 ± 0.04	3.54 ± 0.06
	K-W	NS		*	***	***	***
Mature shell	CO1	25	22.08 ± 0.62	2.51 ± 0.35	2.59 ± 0.02	0.93 ± 0.03	2.39 ± 0.06
	CO2	28	22.77 ± 0.57	2.69 ± 0.37	2.59 ± 0.01	0.90 ± 0.03	2.32 ± 0.07
	MCE	30	23.13 ± 0.54	2.30 ± 0.23	2.55 ± 0.01	1.04 ± 0.02	2.65 ± 0.05
	MGO	35	23.66 ± 0.63	2.40 ± 0.28	2.56 ± 0.01	1.04 ± 0.03	2.67 ± 0.07
	K-W	NS	NS	NS	***	***	***
All shells	CO1	49	17.56 ± 0.85	1.81 ± 0.29	2.52 ± 0.02	1.08 ± 0.03	2.71 ± 0.06
	CO2	59	17.15 ± 0.81	1.78 ± 0.29	2.55 ± 0.01	1.11 ± 0.03	2.83 ± 0.08
	MCE	68	16.65 ± 0.81	1.48 ± 0.21	2.51 ± 0.01	1.26 ± 0.03	3.16 ± 0.07
	MGO	73	17.30 ± 0.84	1.56 ± 0.24	2.47 ± 0.02	1.27 ± 0.03	3.12 ± 0.07
	KW	NS	NS	***	***	***	***

Table 2. Mean value and standard error of shell skeletal and growth parameters. Values for each assemblage in chronological and temperature order. K-W = Kruskal-Wallis rank test; NS not significant, * $p < 0.05$, ** $p < 0.01$. Assemblage labels, length (anterior-posterior maximum distance) and bulk density are after³³.

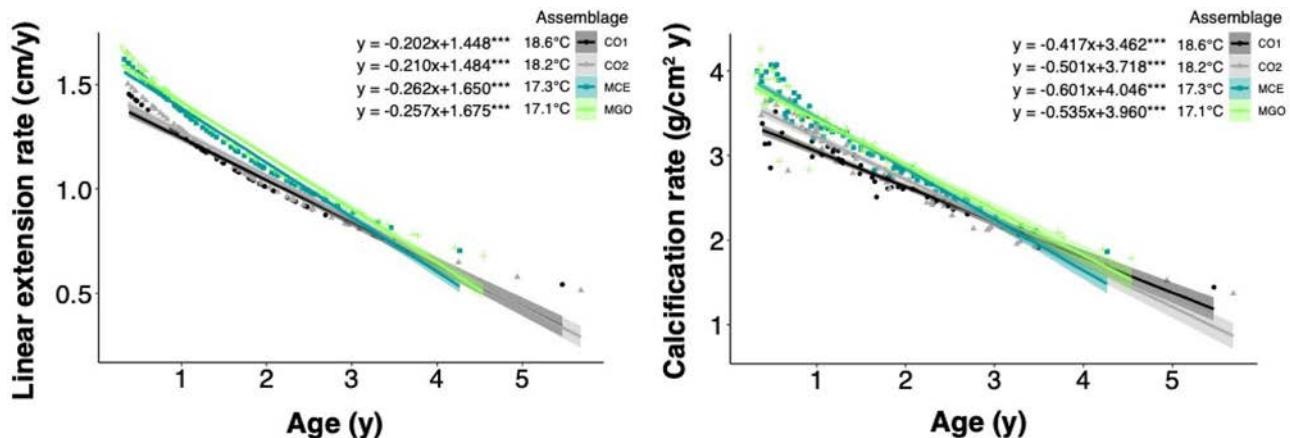


Fig. 2. Correlation of the growth parameters with shell age in the four different assemblages. The regression line and 95% confidence interval for predictions from a linear model ('lm') are visualized using the geom_smooth function of the R ggplot2 package. *** $p < 0.001$.

lower nutrient availability and higher competition for resources (i.e., higher density)³⁹ which affected the L_{inf} value but not their intrinsic growth rate. Thus, at a broad scale (specimen life span), our results support the general physiologic decrease in linear extension rates (and also net calcification rates) as a function of specimen age observed in mollusks and other organisms^{40,41}. For example, in living specimens of *C. gallina* retrieved along the targeted sector of the NAS¹³ the linear extension rate of *C. gallina* decreased with increasing age, highlighting fast growth in the first year of life, with a reduction of around 20% in the second year and around 40% in the third year. Hence, the power-shaped relation between growth parameters and age is confirmed in dead assemblages from modern NAS and HCO samples (Fig. 1). This suggests that the overall growth dynamics did not change even when temperatures were higher (1.5 °C) than present-day and comparable to temperature estimates forecasted for the NAS in the next few decades (i.e., before the end of this century)⁴².

Significant differences in growth patterns emerge when data are analyzed at the subgroup level, especially during the juvenile phase. During the HCO, *C. gallina* occupied warmer northern Adriatic waters where fluvial influence was limited due to the estuarine configuration of the Po and other rivers. Fossil shells from these horizons are consequently denser and display lower linear extension and net calcification rates than modern shells from sites south of the Po Delta, which now experience strong freshwater input (Fig. S1; Table 2). From a thermodynamic point of view, higher temperatures generally promote CaCO_3 deposition by decreasing the energetic costs of shell formation with an increasing aragonite saturation state⁴³. Our data suggest a possibly higher saturation state scenario in the NAS during the Holocene; such a scenario favors the precipitation of a denser shell in warmer seawater conditions. Similarly, the net calcification rates of *C. gallina* during the HCO

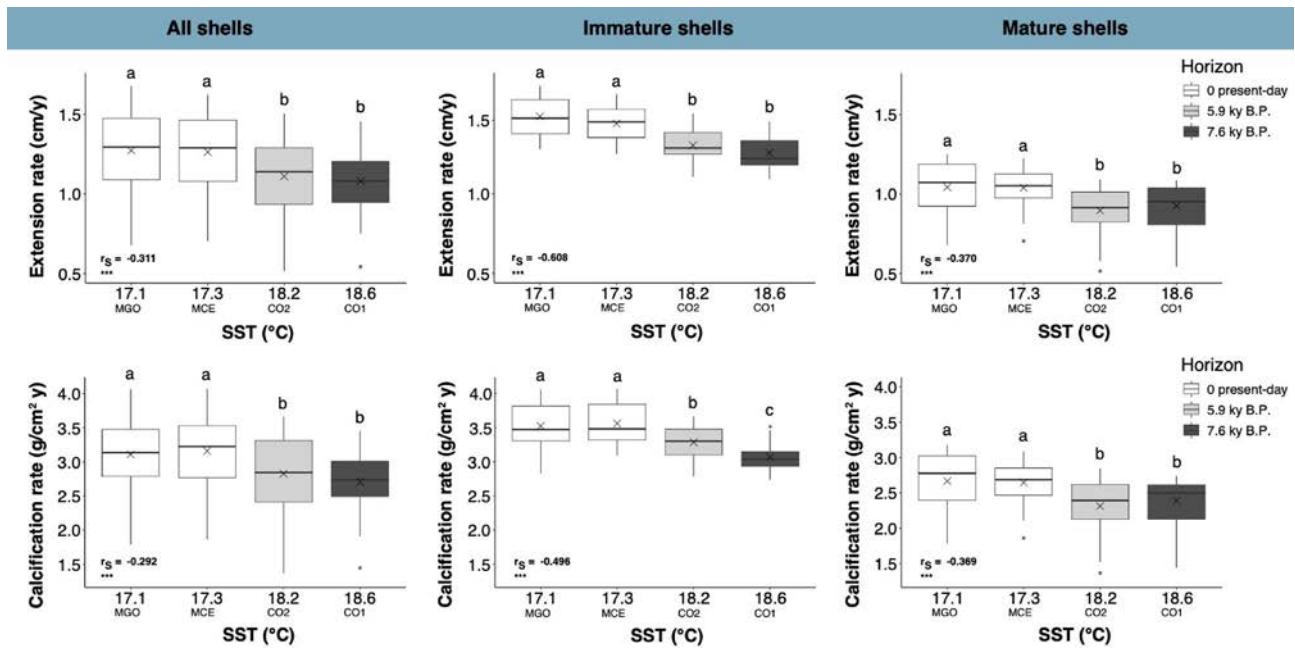


Fig. 3. Variation of shell growth parameters in relation to different SST (Sea Surface Temperature). Three groups were considered: correlations in all shells, correlations in immature shells (< 18 mm), and correlations in mature shells (> 18 mm). The box colors represent the age of the assemblages, ranging from the oldest fossil horizon (dark grey boxes) to the present-day horizon (white boxes). The boxes indicate the 25th and 75th percentiles, the lines within the boxes mark the medians, and the crosses mark the mean values. Black points represent outliers. Different letters indicate statistical differences among assemblages ($p < 0.05$; the number of clams measured for each assemblage is reported in Table 1). r_s , Spearman's rho coefficient. *** $p < 0.001$, NS, not significant.

are significantly depressed during the juvenile and mature stages of life. So, at present, juvenile clams in the investigated area may be allocating more energy towards faster growth (linear extension rate) at the expense of skeletal density to reach size at sexual maturity. Comparatively, juvenile specimens of *C. gallina* from the HCO reached maturity with a delay of around three months, with respect to the modern juvenile specimens that attained maturity after thirteen-fourteen months in the NAS²⁹. This delay could represent an important factor for the near-future management of this economically valuable resource. Higher temperature usually coincides with greater and faster gonadal development in bivalves, although this is mainly due to a greater amount of food ingested⁴⁴. However, a previous study conducted on the clam *Ruditapes philippinarum* found that at higher temperatures (18 °C respect to 14 °C), the clam had a lower ingestion rate. This led to a negative energy balance, causing slower gonadal development that takes place at the cost of the animal's energy reserves⁴⁴. This behavior could explain the growth pattern of *C. gallina* in HCO due to physical factors dependent on a warmer sea (influencing the aragonite saturation state), combined with lower nutrient run-off from the Po River (with respect to present-day).

It is also interesting to note that within the same study area, we have been assisting in the last centuries to a generally decreasing trend in predator-prey interactions, especially between shell-drilling gastropods and their molluscan prey, as testified recently by Zuschin and colleagues on slightly deeper (prodelta) environments from the same targeted area⁴⁵. Here, an unprecedented (relative to the Holocene) increase in dimension and population of *Varicorbula gibba* (another infaunal bivalve) was observed, and it was hypothesized to be driven by ecological release from predation⁴⁵. Thus, the recent removal of predators (naticids) also feeding on *C. gallina* and other mollusks⁴⁶ might have released *C. gallina* and favored those populations characterized by a faster linear extension rate to reach sexual maturity faster. Conversely, denser shells retrieved in *C. gallina* specimens from HCO can provide a buffer against drilling predators⁴⁷ and could also lead to a selective advantage, especially during their juvenile stages, where predation is frequent. However, this resulted in an overall decrease of linear extension rates in *C. gallina* that lived in a warmer Adriatic Sea, thus taking more time to reach the maturity size.

In addition and based on the current knowledge of the main limiting factors acting on this species' growth and the results obtained by *C. gallina* cohorts that lived mainly during the early HCO, we can discuss the principal factors affecting *C. gallina* populations in the near future, that is seawater temperature and nutrients.

Low tolerance to high seawater temperatures. Previous studies highlighted the relatively low tolerance of *C. gallina* to high temperatures compared with other bivalve species, showing a great influence on the overall physiological responses and heavy stress conditions when exposed to high temperatures, demonstrating that temperature could be a tolerance limit for this species^{48–50}. Indeed, specimens of *C. gallina* exposed to high summer temperatures (28 °C) showed reduced energy absorption and increased energy expenditure via respiration, negatively affecting the energy balance⁴⁸ and probably growth, as found during the summer season

(temperature $> 27^{\circ}\text{C}$) in specimens from the eastern coast of Spain³⁶. Generally, reduced summer growth is concentrated in low temperate zones, particularly in water bodies where bivalves are limited in their northward range migration, such as the Adriatic, due to their low tolerance for high seawater temperatures⁵¹. Moreover, oxygen depletion due to high temperatures may harm the clam's physiological performance, as observed in the bivalve *Ruditapes decussatus*⁵². This study evaluated the growth patterns of *C. gallina* in a warmer context than present-day conditions ($\sim 1.5^{\circ}\text{C}$); this would imply for HCO longer time, especially during summer seasons where temperatures could have intercepted the threshold of suboptimal growth, thus depressing linear extension rates of *C. gallina* during the summer season. Following⁴² the mid-twenty-first century climate scenario for the northeast Adriatic region depicts an increase in sea surface temperatures reaching $+3.5^{\circ}\text{C}$ in summer and early autumn⁴². Thus, *C. gallina* growth should be strongly negatively affected by climate warming during the summer, given that present-day summer average SST temperatures (June-September) in the targeted portion of the Adriatic are $25\text{--}26^{\circ}\text{C}$. Therefore, *C. gallina* could show a strong contraction ($> 90\%$) of its distributional range well before the end of the century⁷.

Nutrient supply. The growth of *C. gallina* is also affected by the concentration of the available food supply, as is the case for all filter-feeding organisms^{53,54}. Chlorophyll concentration is a good food proxy for clams, based on the assumption that phytoplankton is the main component of suspension-feeding bivalves' diet⁵⁵. Nowadays, the northern area of the western Adriatic basin is characterized by the presence of Po River run-off, which makes the NAS a eutrophic area with the highest average primary production and Chlorophyll *a* concentration in the Adriatic basin^{13,56}. The higher growth rates of *C. gallina* found in present-day assemblages thus could also result from the current high food availability compared to past environmental settings of HCO, where, especially for early HCO, the geomorphologic setting of the study area was completely different. Indeed, wide estuaries with lagoon barrier systems developed along the targeted coastal areas⁵⁷. Thus, the modern study area shows a river-influenced delta where the freshwater and nutrient-rich plumes mix with nearshore marine waters. Early HCO settings were characterized by estuarine settings with barrier island/lagoon complexes, where freshwater input was mainly mixed within the estuaries and lagoons⁵⁷. Thus, sampled shoreface settings showed a reduced fluvial influence. Increasing temperatures will be associated with increased sea level in the coming decades^{58,59}, reducing the plumes of nutrient-rich freshwater in the Adriatic, with mixing occurring more within the river outlet.

Together with previous (mainly ecological-based) research, our study suggests that *C. gallina* can adapt to (limited) environmental changes resulting from human societal development (see also a recent review by Grazioli and collaborators¹⁵) and highlights its ability to persist in a scenario of limited global warming in the near future (e.g., $\sim 1.5^{\circ}\text{C}$ change relative to present-day conditions in our study area).

Conclusion

This study analyzed the variation in shell growth (i.e., linear extension and net calcification rates) of the bivalve *C. gallina* from assemblages from present-day and Holocene Climate Optimum when the SST in the investigated area of the Adriatic Sea was supposedly 1.5°C higher than today. The comparison between Holocene sub-fossil records and modern assemblages provided insights into the growth dynamics of this bivalve on a millennial temporal scale, overcoming the time limits imposed by laboratory studies. Unlike our previous study (Cheli et al.,³³), which examined shell density and porosity in relation to environmental change, this study quantifies explicitly growth dynamics over the organism's life cycle. We demonstrate for the first time that elevated Holocene Sea surface temperatures coincided with a significant reduction in both linear extension and calcification rates, particularly during early life stages—a finding not previously addressed. Shell growth decreased per unit of time with SST, showing a reduced linear extension rate and calcification in fossil horizons from HCO. Faster linear extension rates and higher calcification rates were found in present-day *C. gallina* populations. Specifically, the present-day environmental context appears more favorable for *C. gallina* growth than the HCO. Given that the HCO may serve as an analog of near-future climate-environmental dynamics, we hypothesize that *C. gallina* will suffer a reduction in growth rate mainly due to climate warming and related environmental changes. Further investigation into the genetic variability of *C. gallina* in HCO assemblage and living populations along the latitudinal gradient in the Adriatic Sea could give insights into a selection of morphotypes more adapted to warmer waters and provide a better understanding of the local adaptation of this species in the face of anthropogenically derived climate warming.

Materials and methods

Study area and sampling

The NAS is a semi-enclosed, shallow basin characterized by a wide shelf with a low topographic gradient of 0.02°a and an average depth of 35 m ⁵⁹. It extends approximately 350 km southward from the Gulf of Trieste and is bordered westward by Italy and eastward by the Balkan Peninsula. The coasts along the Adriatic western (Italian) sector are primarily sandy and strongly influenced by the Po River outflow, which affects the water circulation in the northern Adriatic Sea and plays a fundamental role in the bio-geochemical processes of the basin^{60,61}. The Po is the largest Italian river, and currently, it supplies over 50% of freshwater input to the NAS⁶² and about 20% of the total river discharge in the Mediterranean Sea⁶³. However, the Adriatic sedimentary succession records major changes in the basin configuration in relation to the last glacial-interglacial transition^{64–66}. During the early Holocene Climate Optimum (HCO, 9–7 cal. kyr B.P.), the north-western Adriatic coast was characterized by estuary systems bounded seaward by a series of lagoons that limited riverine plumes into the Adriatic Sea⁵⁷. By contrast, during the last part of the HCO (i.e., between 7.0 and 5.0 cal. kyr B.P.), the area transitioned firstly to a wave-dominated and, after 2.0 cal. kyr B.P., to a river-dominated deltaic system⁵⁷. The last geomorphologic configurations led to the progressively increasing influence of the riverine processes that control present-day coastal dynamics and storage-release of sediments and nutrients. The enhanced freshwater discharge in the

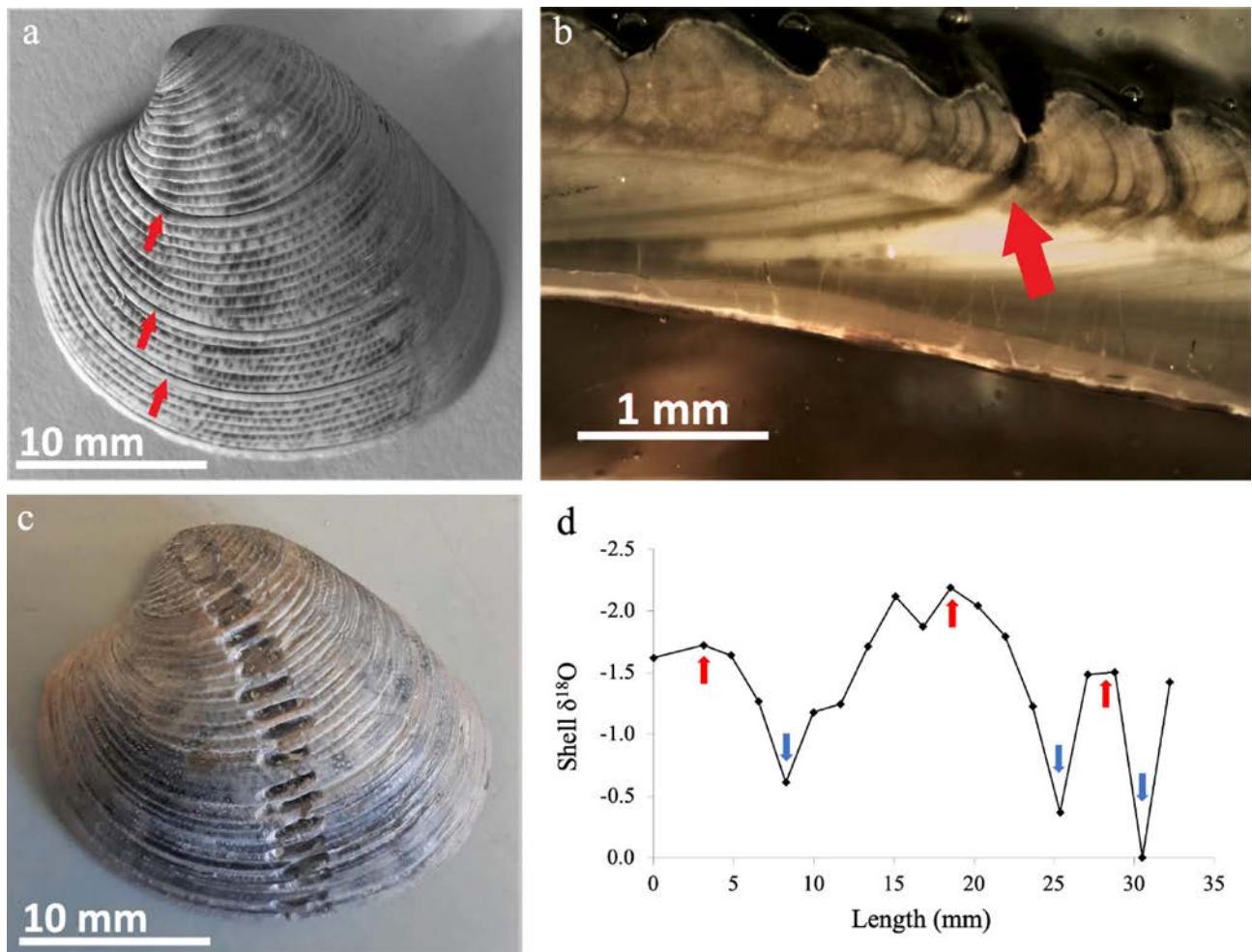


Fig. 4. Shell ageing methods. **(a)** external growth rings on the surface of *C. gallina* after shell scanning; **(b)** internal growth rings in the shell section; **(c)** samples of *C. gallina* with drilling “spots” from the umbo to the ventral edge to determine $\delta^{18}\text{O}$ values along the shell growth axis; **(d)** sinusoidal sequence of lower (summer, red arrows) and higher (winter, blue arrows) $\delta^{18}\text{O}$ values recorded by the shells.

nearshore area, especially during the last millennia, resulted in a strong progradation and the upbuilding of the modern Po Delta⁵⁷.

Our analysis focused on four *C. gallina* assemblages reported by³³ and grouped in two specific time intervals: (1) the HCO (at ~ 7.6 and ~ 5.9 kyr cal. B.P.) documented in subsurface cores and (2) the present day represented by the surfic al dead assemblages of the NAS (Fig. S1). Details on age determination methods, sampling design, specimen collection and preparation, and an estimate of environmental parameters can be found in³³ with a brief description provided in the supplementary material.

Age determination of *C. gallina* specimens

The specimen's age was estimated in a subsample of 30 shells of different sizes in each assemblage using three aging methods: shell surface growth rings, shell internal rings, and (partly) stable oxygen isotope composition (Fig. 4). The growth curves were obtained by counting the total number of visible external and internal rings in each examined valve and then fitted with the von Bertalanffy growth function (VBGF): $L_t = L_{\text{inf}}[1 - e^{-k(t)}]$

L_t is the individual length (the maximum distance on the anterior-posterior axis) at age t , L_{inf} is the asymptotic length (the maximum expected length in the population), k is the growth constant, and t is the specimen's age. Hence, two growth curves for each specimen belonging to a studied assemblage were produced, and a chi-square test of maximum likelihood ratios was used to examine the significance of differences in growth functions between the two aging methods. If differences are present between VBGF curves from valve external and internal ring counting methods, the growth curve will be based on internal ring counting, which is considered more reliable¹⁴. Whereas, if no differences are revealed between VBGF curves from the two-counting methods, a generalized growth curve will be constructed for specimens of each assemblage by merging the growth curves from both methods (Table 1; Fig. 1; Fig. S2). The resulting generalized VBGF curves were used to extrapolate the age in all the other shells by applying the inverse of the generalized VBGF.

$$t = \frac{1}{k * \ln \left(\frac{L_{inf}}{L_{inf} - L} \right)}$$

To validate the data from the two aging methods and the generalized VBGF curves, $\delta^{18}\text{O}$ measurements on three valves per assemblage were carried out at the Stable Isotope Biogeochemistry Laboratory, Department of Earth, Environmental, and Planetary Sciences, Washington University in St Louis (MO-USA). $\delta^{18}\text{O}$ measurements were performed on “spot” samples collected from the prismatic and cross-lamellar layers of the shell, sequentially drilled along the growth axis using a single diamond drill (Fig. 4). The $\delta^{18}\text{O}$ values were plotted against the distance from the umbo, producing a sinusoidal pattern that reflects seasonal temperature variation (Fig. 4). The value closest to the umbo was taken as the starting point (year 0), corresponding to a specific season depending on the $\delta^{18}\text{O}$ value (e.g., maximum for winter, minimum for summer, or intermediate for spring/autumn). One full year of growth was defined as a complete seasonal cycle in the isotopic curve (e.g., from winter to winter or from summer to summer). Subsequent years were counted by identifying these repeating seasonal cycles. The age-length key obtained from this isotope-based method was plotted alongside the age-length key from the two aging methods (Fig. S2).

Determination of linear extension and net calcification parameters

In *C. gallina* specimens, linear extension rates were obtained from the valve length/age ratio (cm y^{-1}), while the net calcification rate (mass of CaCO_3 deposited per year per unit area, $\text{g cm}^{-2} \text{y}^{-1}$) was calculated for each valve by applying the formula: bulk density (g cm^{-3}) \times linear extension rate (cm y^{-1}). The bulk density of each valve examined in our study was sourced from the literature (see Table 2 in³³). Correlation analyses between linear extension rates, net calcification rates, and SST were performed to test for any significant relationship developed over geological time as a function of temperature-driven environmental changes. Differences in the growth of *C. gallina* specimens were also examined in relation to sexual maturity, which is reached at around 18 mm in modern times¹⁵ to account for potential variations in the biomineralization process during different stages of the bivalve’s life cycle.

Statistical comparison of shell parameters among assemblages

Since assumptions for parametric statistics were not fulfilled, the non-parametric Kruskal-Wallis equality-of-populations rank test was used to test for differences in shell parameters among assemblages, followed by Dunn’s post hoc tests. Mann-Whitney U tests were applied to determine the differences between ages obtained from external and internal growth rings within each assemblage. For each assemblage, the age-length functions were fitted with the VBGF using a non-linear model, which estimated the VBGF parameters through bootstrapping. Kimura’s likelihood test was then used to compare VBGF functions among assemblages. Regarding age in each assemblage, logarithmic regression models were fitted to shell bulk density, while linear models were fitted to linear extension and net calcification rates. Spearman’s rank correlation coefficient (rho) was used to evaluate trends between shell growth parameters and sea surface temperature. All statistical analyses were computed using RStudio software⁶⁶.

Data availability

The codes and dataset generated and analysed during the current study are available from the corresponding authors upon request.

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Author contributions

A.C., A.M., S.G., and D.S.: designed the research and collected data; A.C., A.M. and A.R. wrote the initial draft and performed analyses. A.C., A.M., F.P., A.R., G.F., S.G., and D.S. contributed to the final version.

Declarations

Competing interests

The authors declare no competing interests.

Additional information

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