Fitness-related traits are maximized in recently introduced, slow-growing populations of a global invasive clam

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ABSTRACT

Many species are shifting their ranges being forced to rapidly respond to novel stressful environmental conditions. Colonizing individuals experience strong selective forces that favor the expression of life history traits notably affecting dispersal and reproductive rates in newly invaded habitats. Limited information is currently available on trait variation within the invasive range despite being critical for understanding ecological and evolutionary factors that drive the process of range expansion of invasive species. Here we evaluated life history shifts of the widely introduced Asian clam *Corbicula* within its invaded range. Through an exhaustive literature search, we obtained data for 17 invasive *Corbicula* populations from different ecosystems worldwide to test the relationship between population and individual parameters relevant to the process of range expansion. Our main results are that (i) recently introduced *Corbicula* populations are characterized by low conspecific density and low rate of population increase, (ii) clams reproduce earlier in slow-growing populations, and (iii) density had no effect on exponential population increase. All invasive *Corbicula* populations analyzed in this study (Form A/R) are mostly fixed for one genotype. Our results therefore suggest that adaptive phenotypic plasticity favored the expression of traits that maximize fitness in recently established populations facing stronger *r*-selective forces, thereby increasing their chances to overcome difficulties associated with low densities and low population increase in newly invaded areas.

1. INTRODUCTION

The ongoing increase in frequency of extreme climatic events, caused by anthropogenic climate change, and global transport of people and goods is altering ecosystems by triggering a wide array of irreversible consequences (Bossdorf et al., 2005; Hoegh-Guldberg & Bruno, 2010; IPCC, 2014; Walther et al., 2002). Changes in local conditions, in particular those driven by human-induced climate change, are predicted to affect ecological processes across several levels of organization, likely over the coming decades. Such changes are altering the performance of individual organisms, the dynamics of populations and the distribution of species, ultimately modifying ecosystem properties and function (Hoffmann & Sgró, 2011; Lonhart et al., 2019).

Many species are forced to rapidly respond to novel stressful environmental conditions when shifting their distribution and colonizing new areas. If non-native species have difficulties in adapting to such conditions and fail to track abrupt environmental changes, populations become vulnerable to decline and extinction (Hill et al., 2011; Hoffmann & Sgró, 2011). Phenotypic plasticity is the ability of a single genotype to express different phenotypes when exposed to different environmental conditions (Travis, 1994; West-Eberhard, 2003). In a new environment, plasticity rapidly favors the appearance of a novel phenotypic variant, promoting successful colonization and increasing the likelihood for the long-term persistence of populations facing new environmental stress brought about by range shifting (Hill et al., 2011; Pfennig et al., 2010; Pigliucci et al., 2006). Adaptive plasticity that places populations close enough to the new favored phenotypic optimum for directional selection to act provides the first step in adaptation to new environments on ecological time scales (see review in Ghalambor et al. 2007). Individuals at the expanding edge of the species distribution experience strong selective forces (*r*-selection) that can have profound impacts species life

histories. These forces can, for instance, favor the expression of traits that optimize population growth and survival from offspring to maturity (e. g. reduced age of maturity and reproduction, increased growth rate, increased energy allocated to reproduction; Stearns, 1976, 1977), thus affecting dispersal and reproductive rates (Phillips, 2009; Shine et al., 2011).

Colonization events through either continuous range expansion or jump dispersal are usually driven by a small number of individuals that are the first to arrive at a new point of the distribution, where they benefit from increased space open for colonization and low conspecific density, below carrying capacity (Cole, 1954; Phillips et al., 2010; Sakai et al., 2001). Hence, the density gradient occurring at the expanding range (i. e. grading from zero to carrying capacity) creates different selective environments where individuals from a recently established population face stronger *r*-selection compared to those from long established ones, which are exposed to selective *K*-type forces (e. g. Phillips et al., 2010). According to theory, low densities favor the expression of life-history traits related with *r*-type selections (Burton et al., 2010; MacArthur & Wilson, 1967; Phillips, 2009), leading to exponential (non-density regulated) population growth in recently established populations while denser, long-established ones follow a density-regulated population growth (Brook & Bradshaw, 2006).

Recently established populations are characterized by low conspecific density and low rate of increase, often leading to lag-times between initial colonization and the onset of rapid population growth (Sakai et al., 2001). The duration of the lag phase is mediated, among others, by a number of non-mutually exclusive ecological and evolutionary mechanisms (e. g. low encounter rate for reproduction, density-dependent effects, adaptation, and selection of new genotypes; Bossdorf et al., 2005; Davis, 2005; Ricciardi, 2013). An effective mechanism

that increases the rate of population growth is early reproduction, which results from fast growing individuals (Cole, 1954; Lewontin, 1965; Roff, 1993). Individuals that grow faster will attain reproductive sizes earlier and bring down generation times, promoting a rapid build-up of population numbers (Cole, 1954; Lewontin, 1965; Phillips, 2009; Roff, 1993) and accelerating primary invasion and secondary spread of invasive species (Lockwood et al., 2007; Phillips et al., 2006; Skellam, 1951). Several studies have compared life-history traits between native and invaded ranges ('home-and-away' comparisons). Such comparisons could lead to erroneous interpretations of the patterns observed because it would be difficult to disentangle what were the processes (e.g. propagule bias and/or evolution in populations at spatial (dis)equilibrium) causing the shifts in life-history traits (Phillips et al., 2010 and references therein). Interestingly, only a few studies have looked for such trait changes in introduced species (Gaston, 2009), despite knowledge on life-history trait variation within the invasive range being critical for better understanding the process of species range expansion.

Bivalves of the genus *Corbicula* (Megerle Von Mühlfeld 1811) are native to Southeast Asia, the Middle East, Australia and Africa (Araujo et al., 1993) but they have colonized a large part of the Americas and Europe (Crespo et al., 2015; Mouthon, 2001; Schmidlin & Baur, 2007). The successful invasion of *Corbicula* clams has been mainly attributed to their rapid growth and maturation, high fecundity, and high dispersal making this bivalve genus one of the most relevant faunal non-indigenous groups in aquatic ecosystems (reviewed in Sousa et al. 2008a). Invasive freshwater *Corbicula* populations are constituted by a small number of lineages composed of hermaphroditic individuals reproducing through androgenesis, for which the most widespread and abundant genotype in the invaded area is the lineage so-called Form A/R, widely known as *C. fluminea* (Komaru et al., 1998; Pigneur et al., 2012, 2014). The current taxonomic status of the invasive *Corbicula* lineages in the Americas and Europe

is still largely unresolved despite several morphological and genetic studies (reviewed in Pigneur et al. 2014). In the present paper, we deal with this form and refer to it as *Corbicula* to avoid entering unsettled taxonomic debates.

Here we investigated how different life-history traits of the Asian clam *Corbicula* responded to different selective pressures occurring at varying population contexts within the introduced range: (i) newly- vs. long-established, (ii) low vs. high density, and (iii) low vs. high rates of population increase; all of them being linked to the geographic range shifting of species' distribution (i. e. front/core populations). To do so, we searched for peer-reviewed articles providing enough information to estimate time since population introduction as well as different population and individual processes and characteristics including population density, individual growth rate, minimum age at sexual maturity, and lifespan of global populations of this worldwide invader. Different studies suggest that *Corbicula*'s population dynamics are chiefly regulated by environmental conditions (e. g. Crespo et al., 2015, 2017; Gama et al., 2017), whereby population declines are typically driven by periodic mass mortality due to extreme hydro-meteorological events (Ilarri et al., 2011; McDowell et al., 2017; McDowell & Sousa, 2019). This strongly suggests that invasive *Corbicula* populations would primarily follow a non-density regulated population increase and, hence, they might not level off at carrying capacity as happens with many other species (e.g. Brook and Bradshaw, 2006). We focused our predictions on life history strategies exhibited by individuals from recently established populations, which typically occur at the invasion front, and likely subjected to strong *r*-selective pressures with respect to those from older, long-established ones. We tested the hypotheses that population density and population increase are directly related with time since introduction. *Corbicula* is widely known to exhibit rapid population growth, being able to reach extremely high densities (reviewed in McMahon, 2002). We expected density to have

no significant effect on population increase even within a wide range of densities in nature, though high densities are considered likely to have a negative effect on individual growth rates. We also asked whether clams that grow faster and reproduce earlier are favored in slow-growing populations at the expanding front. A related prediction is that clams that grow rapidly would have a shorter lifespan with respect to individuals exhibiting slow growth.

2. METHODS

2.1 Literature search

We exhaustively searched the literature for peer-reviewed articles on *Corbicula* (*Corbicula fluminea sensu lato*) providing enough information to estimate the individual growth rate, population density, rate of exponential population increase, time since colonization, water temperature, and conductivity of global populations of this clam. To do this, we first identified relevant studies from ISI Web of Knowledge (Web of Science Core Collection, 1900 through 2017) by conducting a first search using the field tag "topic (TS)" in the "Advanced search" option and the search string: TS="Corbicula fluminea" OR TS="asia* clam" (search #1). We conducted a second search including topics TS="population dynamic*" OR TS="population size*" OR TS="population densit*" OR TS="population structure" (search #2). We ran a third search using TS="growth" OR TS="individual growth" OR TS="population growth" (search #3). Finally, we combined searches #2 and #3 into search #4 (#4= #3 OR #2), and restricted the latter by search #1 to obtain a final set of candidate studies (search #5= #4 AND #1).

This literature search was finished on 27 September 2018 and this protocol identified 215 candidate studies (database search: 215 candidates; gray literature and unpublished data: 0 candidates). To identify papers potentially missing in this list, we subsequently scanned the

references in these papers and conducted a Google Scholar search using keywords mentioned above and added 8 studies to this candidate pool (Fig. S1). A total of 67 studies that belonged to research areas unrelated to the topic of the current study (*e. g.*, applied chemistry, toxicology, paleontology) were excluded from this candidate pool by refining the set based in 'Web of Science Categories'. The resulting studies (n = 156) were finally inspected individually in detail to assess their eligibility for our analyses according to the following inclusion criteria: i) Studies that assessed invasive populations of *Corbicula* clams (i. e. studies exclusively providing estimates for native populations were excluded); ii) Studies that reported or provided data enough to extract the population density and age/size distribution for >6-months (see below); iii) Studies that assessed populations in reasonably "average" natural environments, such that populations from heavily impacted sites (e. g. thermal plumes) were excluded. We finally retained a sample of 16 studies, covering 17 invasive populations of *Corbicula* from different ecosystems worldwide (n = 140 out of 156 were excluded; see Fig. S1 for further details on screening and eligibility criteria). Whenever population data was only graphically presented, we contacted directly the authors. However, if data were no longer available or authors did not respond, we used the WebPlotDigitizer software (http://arohatgi.info/WebPlotDigitizer/) to extract data and accurately estimate population density and growth rates. This step-by-step criterion for the literature search and data collection attempted to reduce any potential selection bias in the long-term populations that might occur because old high-density populations (with higher growth rates) could be more studied than young low-density populations (with low growth rates), which would be more likely to be extirpated if population characteristics were not changing with time.

2.2 Population density and exponential population increase

For each population selected from our literature search, we obtained the mean population

density (ind m^2) for the entire study period or extracted temporal values to estimate it whenever this value was missing in the text. *Corbicula* clams generally exhibit a bivoltine juvenile release pattern (i. e. producing two generations a year McMahon, 2002; Sousa, Antunes, et al., 2008), leading to one or more high-density peaks related to recruitment events yearly. However, density in the field fluctuates almost constantly generating several peaks and valleys that do not necessarily indicate significant recruitment events. *Corbicula* populations exhibit rapid growth (Franco et al., 2012; McMahon, 2002; Sousa et al., 2006; Sousa, Antunes, et al., 2008). Hence, the rate of exponential population increase of *Corbicula* can be expressed by deriving the classical exponential growth equation:

$$r = \frac{\ln\left(\frac{\mathrm{Nt}}{\mathrm{N0}}\right)}{\mathrm{t}}$$

One of the goals of this study was to compare population growth rate during periods of substantial growth across populations, and it is such capacity that is of interest to better understand *Corbicula* invasiveness. Hence, to estimate the rate of exponential population increase (*r*, hereafter population growth rate), we set a criterion to determine relevant recruitment peaks in each study by using the average density throughout the study period as a threshold. By doing so, we excluded time periods where population growth was extremely low likely due to mass mortality due to extreme hydro-meteorological events (e.g. a period of dramatic demographic decline), usually characterized by a lag phase (where population growth can be negligible) followed by an exponential increase (Ricciardi, 2013). Likewise, this approach allows for exclusion of large time periods resulting from *Corbicula*'s inability to reproduce year-round as they generally exhibit a uni- or bivoltine juvenile release. We therefore considered the lowest density value (N_0), the highest peak above the average density (N_t), and the time period elapsed between the valley and the peak (t). Intermediate density values comprised in each period (i. e. defined by N_0 and N_t) were used to fit the

growth equation and obtain *r*. Population rates of increase were estimated using GraphPad Prism version 7.00 for Windows, GraphPad Software, La Jolla California USA, www.graphpad.com. Note that periods defined by non-relevant peaks (i. e. below the average density threshold) were discarded, as were periods of progressive density decline between a relevant peak and the next N_0 value (see Fig. S2 for further details on the step-by-step procedure used to estimate population increase). When populations exhibited more than one relevant density peak throughout the study period, the population increase rate for such populations was calculated as the average of single *r* estimations. Then, we standardized estimates on an annual basis to make them comparable. Note that the calculated *r* values provide a relative growth rate, that is, one that can be used to compare different populations to one another, rather than a population growth rate that can be used to forecast future population sizes.

2.3 Individual growth rates and parameters

For studies that only presented data on shell length-frequency distributions (shell length [mm] being the greatest distance between the anterior and posterior margins of the shell valve), we identified age cohorts using that information and applied the Bhattacharya's method available in FISAT II software (Version 1.2.0, FAO-ICLARM Fish Assessment Tools; (Gayanilo et al., 2002). To confirm each component of normal distributions from the modal progression analysis, we used the NORMSEP method also available in the FISAT II software (Pauly & Caddy, 1985). Once we had cohorts identified for all the studies, we calculated individual growth parameters using the same equations for all the studies considered in order to make sure that results were fully comparable across studies. For each cohort present in each of the populations considered in this study, we fitted the von Bertalanffy-growth function with seasonal oscillations (hereafter SVBGF) proposed by

Pauly and Gaschutz (1979), Hoenig and Hanumara (1982), and Somers (1988):

$$L_t = L_\infty (1 - exp^{-K[(t-t_0)+T_1-T_2]}),$$

$$T_1 = C \sin(2\pi (t - t_s)) / 2\pi,$$

and

$$T_2 = C \sin(2\pi (t_0 - t_s)) / 2\pi,$$

where L_t is the predicted shell length at age t, L_∞ is the asymptotic shell length, K is the growth constant of dimension time (year^{-1} in most seasonally oscillating growth curves) expressing the rate at which L_∞ is approached, t_0 is the theoretical 'age' the clam has at shell length zero, C expresses the relative amplitude of the seasonal oscillation and varies between 0 and 1 (0 indicating lack of summer-winter differences in growth, it was constrained to be less than or equal to 1 during model fitting), and t_s is the starting point of the oscillation.

The parameters of the function were estimated by the modeling method available in JMP statistical software (v9.0 SAS Institute). When preliminary results of the SVBGF failed to converge to estimate the asymptotic shell length (L_∞), we used the maximum shell length (*Lmax*) observed or reported at/for each study site to calculate the asymptotic shell length following the equation suggested by Taylor (1958):

$$Lmax / 0.95 = L_\infty$$

These values were fixed when we performed a second fit.

Because of the negative correlation between growth parameters (K and L_∞) prevents making comparisons based on individual parameters (Ramón et al., 2007; Vakily, 1992), we used the growth-performance index (*GPI*), which reflects the growth rate of a given organism of unit

length. In other words, *GPI* can be viewed as the (theoretical) value of *K* that would occur in organisms with an L_∞ value of 1 unit of length (Munro & Pauly, 1983), and it was defined by Pauly and Munro (1984) as:

$$GPI = 2 \log_{10} L_\infty + \log_{10} K$$

We calculated the *GPI* values for cohorts from each study site based on growth parameters (L_∞ and *K*) obtained after fitting the SVBGF using the *Growth Performance Indices* application of the FISAT II software (Gayanilo et al., 2002).

2.3.1Minimum age at sexual maturity

The minimum age at sexual maturity was estimated as the time (months) elapsed until clams reach the minimum size (shell length) at which invasive *Corbicula* clams become sexually mature, which, on average, is 10.0 mm ± 0.8 SE (n = 13 reported sizes). This value was obtained from shell sizes reported in the literature (Aldridge & McMahon, 1978; Cao et al., 2017; Eng, 1979; French & Schloesser, 1991; Gardner et al., 1976; Heinsohn, 1958; Ituarte, 1985; Kennedy & Van Huekelem, 1985; Kraemer, 1979; Kraemer & Lott, 1977; McMahon, 2002; Rajagopal et al., 2000). We acknowledge that in some conditions and populations, clams might mature at a different pace and thus introduce a degree of error in such cases. While this error is unavoidable until more detailed data is available, we are confident that the 10-mm estimate applies to many real-life situations and thus our results based on it are robust.

2.3.2 Lifespan

As *t0* is the theoretical ‘age’ the clam has at shell length zero, *tmax* can be defined as the theoretical ‘age’ the a clam reaches its maximum shell length (*Lmax*). Since *Lmax* is dependently related to a maximum point in time (years), it is assumed that *Lmax* implicates

tmax (*Lmax* ⇒ *tmax*) (Bonel & Lorda, 2015). Thus, the theoretical maximum lifespan of a clam cohort can be approximately expressed as:

$$Lifespan = tmax - t0$$

2.4 Time since colonization and key environmental variables

For the same 17 populations and sites studied, we calculated the time elapsed since population foundation (hereafter time since colonization) as the difference between the sampling dates in the first ecological survey and the first report of introduction. The accuracy of the introduction date is generally very difficult to know, however, many of the studies used report range expansions of a conspicuous invader across well studied regions, where the appearance of the species might have been anticipated and readily detected by ongoing monitoring actions in the initial stages of invasion. Therefore, we consider that the possible error introduced should be moderate to low and the value obtained can be viewed as a good approximation. Moreover, we obtained water temperature and conductivity values for each site averaging monthly mean values for the same period used to estimate growth rates in each study. Other biotic and abiotic variables are known to interact with temperature to drive growth rates of invasive bivalves including food availability, dissolved oxygen, and heavy metal concentration (Bonel & Lorda, 2015; Vohmann et al., 2010). However, we only considered water temperature (ºC) and conductivity (µS/cm) in the analyses because they have been reported to play a key role on the *Corbicula*'s biology and ecology (Crespo et al., 2017; Franco et al., 2012; Magnussen, 2007). Indeed, temperature has emerged as the most important variable explaining the current distribution and, also, predicting that climate change will favor the expansion of *Corbicula*'s into colder areas at higher latitudes (Gama et al., 2017). Likewise, salinity has been considered a key factor influencing the success and velocity of the invasion of new estuarine ecosystems (e.g. Sousa et al., 2006). As *Corbicula*

populations considered herein belonged to a broad range of freshwater and estuarine habitats, we therefore included conductivity to serve as a *proxy* for salinity levels. In order to keep values seasonally unbiased, only entire years were used in subsequent calculations. Whenever these parameters were not reported in the original paper, we obtained values for the same sites and years from the literature and reliable online databases (e. g. https://waterdata.usgs.gov/nwis/qw). When data was only graphically presented, we extracted it following the procedure mentioned above.

2.5 Statistical analyses

We ran five separate multivariate weighted linear regressions in order to test ecological processes and characteristics at three different (but non-mutually exclusive) levels: 1) population level, 2) population and individual level, and 3) individual level. In this sense, a response variable at the population level can be considered as an explanatory variable when analyzing a response at the individual level. In all cases, we computed the variance of the effect size from different populations, which were then used as weights in linear regressions. The inverse-variance weighting allows combining the results from independent measurements giving more importance to the least noisy measurements and *vice versa* (Borenstein et al., 2009).

We were interested in testing individual effects of predictors, which are related to life-history traits of *Corbicula* (hereafter reduced models). However, we added annual mean water temperature and conductivity as quantitative variables in each model (hereafter full models). Preliminary analyses showed that latitudinal effects are strongly correlated with water temperature ($r = –0.84$). Hence, it was excluded from the models to avoid multicollinearity. In some cases, we also included interaction terms between explanatory variables. We used the

partial *F*-test to compare models—that is, full models with and without the interaction term, and full *versus* reduced models. As we had strong *a priori* hypotheses as to the direction of effects, we used one-tailed tests. Variables were Ln-transformed to fulfill the models' assumption of normality and homoscedasticity, which were tested using the *gvlma* package (Peña & Slate, 2006). Prior to analyses, we standardized explanatory variables to zero mean and a unit of variance in order to get directly comparable effects. All analyses were carried out in R version 3.6.1 (R Development Core Team 2019). Values are given as means ± SE unless otherwise stated.

3. RESULTS

3.1 Literature search

Populations analyzed in this study belonged to Europe, North and South America (Fig. 1; Table 1). These included eight populations in the United States, three in France, three in Argentina, and three in Portugal (see Table 1 for detailed location, bibliographic source, and code number used herein to refer to the populations, for each of the 17 *Corbicula* populations selected, and Table S1 for additional information).

3.2 Density, growth, time since colonization, and key environmental variables

Overall, mean density of *Corbicula* from 17 locations worldwide spanned 32 to *ca*. 9,500 ind. m^{-2}, population growth rate (*r*) varied from 0.37 to 189, individual growth rate (*GPI*) ranged 2.53-3.49, age at first reproduction 3.1-22.7 months, and lifespan 2.3-8.0 years (Table 1). On the other hand, time since colonization ranged 2 to 28 years in the populations studied (Table 1) and yearly water temperature ranged from 13.2 to 20.7 ºC and conductivity from 68 to 8,457 µS/cm (see Table S1 of the Supporting Information for details). In all cases, we found no evidence that adding an interaction term with variables of interest nor temperature and

conductivity in each model significantly increased its predictive power (for details see '*Model comparison*' in Supporting Information). We therefore report estimates and statistics from the reduced models in Table 2.

At the population level, we found that younger *Corbicula* populations (i.e. newly established) were characterized by low density and slow population growth rate with respect to older, long established ones (Fig. 2a, b). We found that density had no significant effect on population increase (Fig. 2c). When analyzing responses at the interaction between population and individual processes and characteristics, we found that clams tended to grow faster (marginally non-significant) in low-density and slow-growing populations (Fig. 2d, e). In contrast, we observed a strong positive response of the minimum age at sexual maturity as a function of population growth rate, clearly indicating that clams reproduce earlier in slow-growing populations (Fig. 2f). This is directly linked to processes occurring at the individual level, where clams with higher growth rates reach the minimum reproductive size earlier (10 mm) and have shorter life span than clams that grow more slowly (Fig. 2g, h).

4. DISCUSSION

This is the first study that evaluates how population and individual parameters, involved in the process of range expansion, varied in response to different selective pressures occurring within the introduced range of the worldwide invasive Asian clam *Corbicula*. Our main results showed that (i) recently introduced *Corbicula* populations are characterized by low density and reduced population growth rate, (ii) clams reproduce earlier in slow-growing populations, and (iii) population growth rate was unfettered by density (summarized in Fig. 3).

Density of *Corbicula* populations increased with time since colonization, which is consistent with the standpoint that recently established populations typically occur at low conspecific density, below the carrying capacity (Cole, 1954; Sakai et al., 2001). Likewise, we found that population growth rate and time since colonization were directly related, suggesting that, in newly colonized areas, populations could have experienced stronger *r*-type selection favoring fitness-related traits effectively increasing the rate of population growth, as it was observed in long established populations (Burton et al., 2010; MacArthur & Wilson, 1967; Phillips, 2009). Population density did not predict changes in the rate of population increase, as variation in population growth remained relatively low even at high-density values. By definition, long established and denser populations tend to be density regulated as they approach to carrying capacity (logistic population growth; e. g. Phillips et al., 2010). Considering the so-called *r*/K selection continuum, this implies that such populations are predicted to shift from the *r*-towards the K-endpoint (MacArthur & Wilson, 1967)—that is, selection shifts a population from density-independent to density-dependent regulation. However, *Corbicula*'s population growth rate did not decrease with increasing density despite the fact that some of the populations included in this study exhibit extremely high densities exceeding, in some cases, 100,000 individuals per square meter (e. g. Eng 1979; Table 1).

Why was population growth unfettered by density regulation? A large number of studies have analyzed density dynamics of *Corbicula* and commonly assumed that, following exponential growth, the clam population achieves densities approaching the carrying capacity of the environment (e.g. Sousa et al., 2006). However, to our knowledge, no studies have ever conducted (field or laboratory) experiments to test whether population growth is limited by increasing conspecific density, which, in turn, is expected to reduce local biotic resources. In other words, there is a complete lack of knowledge on the true effect of density on population

growth rate. On the contrary, different studies have shown that *Corbicula*'s dynamics are chiefly influenced by environmental variables (e. g. water temperature, conductivity, pH, and/or dissolved oxygen) and seem to be less influenced by population's density (e. g. Crespo et al., 2015; Gama et al., 2016). Indeed, periodic mass mortality due to extreme hydro-meteorological events (e.g. extremely high or low temperatures, drought) could play an important role with infrequent, but catastrophic population declines (>99% in some cases; Ilarri et al., 2011; McDowell et al., 2017; Mouthon & Parghentanian, 2004). These events have been described (though not quantified) in several other systems as well (e. g. McDowell & Sousa, 2019; Vaughn et al., 2015). Despite that a given *Corbicula* population might approach or even grow beyond the carrying capacity of the ecosystem, evidence from previous studies (and those cited herein) suggests that population size significantly varies with time. This means that the population is at non-equilibrium and that such fluctuations (or boom-bust dynamics; Strayer et al., 2017) are likely driven by density-independent factors, as the above-mentioned hydro-meteorological events. Thus, it does seem reasonable to infer that *Corbicula*'s population growth rate is non-density-regulated and will not level off at carrying capacity, as expected from a highly invasive organism as the Asian clam (McMahon 2002). Our results suggest that this bivalve seems to be far from approaching carrying capacity, even at extremely high densities, supporting the idea that individuals from long established denser populations do not face a *K*-selective environment but a less strongly *r*-selective pressure.

Individuals tended to grow faster in low-density populations experiencing low rates of growth. Likewise, *Corbicula* clams reproduced earlier in slow-growing populations. In recently introduced populations, periods of negligible population increase can be driven by non-mutually exclusive ecological and evolutionary adjustments such as Allee effect, changing abiotic conditions, and adaptation and selection of new genotypes imposed by novel

environments conditions in a new habitat (Bossdorf et al., 2005; Davis, 2005; Ricciardi, 2013; Sakai et al., 2001). Such conditions can lead to a strong selection on life-history traits that optimize population increase and survival to maturity of offspring (Stearns, 1976, 1977). As predicted by life-history theory, individuals from recently established populations (typically occurring at the invasion front) face stronger *r*-selection relative to conspecifics from older populations (MacArthur & Wilson, 1967), favoring traits that accelerate the range expansion. For instance, some empirical observations have revealed that individuals from recently colonized areas grow faster, and thus, reproduce earlier, than do conspecifics from older, long-established populations (e. g. butterflies: Hanski et al., 2006; Australian cane toads: Phillips, 2009; tallow trees: Siemann & Rogers, 2001). Accordingly, our results for *Corbicula* clams showing fast growth and, thus, early reproduction support the idea that such populations faced strong *r*-selection, which favored traits associated with high reproductive rates in recently introduced, slow-growing populations.

The increase in individual growth was associated with a decrease in the clams' life span likely due to trade-offs between different life-history traits. Such trade-offs can be interpreted as an effective mechanism to bring down generation times, promoting a rapid build-up of population numbers, increasing population growth in newly colonized areas (Cole, 1954; Lewontin, 1965; Roff, 1993) and accelerates primary invasion and secondary spread of invasive species (Lockwood et al., 2007; Phillips et al., 2006; Skellam, 1951). Collectively, our results suggest that reduced population increase of the current *Corbicula* population triggers the shortening of the time that the clams need to reach maturity by favoring clams that grow the fastest. Thus, as early-maturing clams reproduce earlier, this has the effect of fostering an increase in size of the future population. In this way, incipient *Corbicula* populations could increase their chances of overcoming the difficulties associated with low

densities and survive to the population establishment stage, which can clearly subsidize the bivalve's invasive abilities in newly colonized areas.

In conclusion, in this study we have provided evidence supporting the idea that life-history traits of individuals can respond to selective pressures occurring under varying population contexts (e.g. varying time since population foundation, low vs. high population density, and low vs. high rates of population growth). All invasive *Corbicula* populations analyzed in this study are mostly fixed for one genotype (Form A/R). Our results, therefore, provide reasonable ground for advocating that clams from recently established, slow-growing populations experienced a stronger *r*-selective environment than those from long-established ones; and that adaptive plasticity can maximize the expression of fitness-related traits, such as individual growth rate and the minimum age at sexual maturity, in these populations. While these responses could be interpreted as an effective mechanism to decrease the risk of stochastic extinction associated with low densities at the invasion front, population growth rate seems to be independent of population density in a wide range of environmental conditions within *Corbicula*'s invasive range. Although other relevant variables could not be considered in our models, this study represents a first step in the attempt to understand in a wide geographical perspective how different selection pressures affect fitness-related traits of the Asian clam during range shifting.

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TABLES

Table 1. Summary of ecological and growth variables (mean ± SD) of worldwide invasive populations of the Asian clam *Corbicula* reviewed from the literature. *Density* is the estimated abundance of clams at the study site (ind./m^2), *Pop.growth* is the rate of exponential population derived from the classical form of the exponential growth equation, *Time* is the number of years elapsed since population colonization at the time of the original study (years), *Ind.growth* is the individual growth rate (*GPI*, a *proxy* for individual growth rate), *mASM* is the minimum age at sexual maturity (months), and lifespan (years), *ID#* is a number assigned arbitrarily for identification purposes. The country, specific study location, and reference to the source paper of the original study used to build the information are shown for each population. See the main text for details on how data was collected and variables were built from the literature. For extended information on each population, see Table S1 of the Supporting Information.

Country	Study location	*Time*	Population parameters		Individual parameters			*ID#*	*Reference*
			Density	*Pop.growth*	*Ind.growth*	*mASM*	*Lifespan*		
Argentina	Punta Atalaya, Río de la Plata	3.5	339±405	2.20±0.01	2.87±0.07	4.30±0.13	3.83±0.57	1	Ituarte (1985)
	Paraná de las Palmas River	25.5	1,053±768	14.44±0.31	2.73±0.01	9.23±1.17	6.82±0.05	2	Cataldo and Boltovskoy (1999)
	Río Negro estuary	15.5	94±8.3	7.85±0.10	2.94±0.23	5.32±1.66	3.55±0.41	3	Hünicken et al. (2019)
France	Saone River	12.5	302±186	14.65±0.01	2.54±0.13	12.73±2.42	7.55±2.17	4	Mouthon (2001)
	Loire Lateral Canal	17.0	992±1209	41.17±0.12	2.80±0.24	8.93±5.95	5.17±3.00	5	Mouthon and Parghentanian (2004)
	Canal of Roanne	17.0	626±897	25.50±0.01	2.77±0.16	8.25±3.43	4.27±1.58	6	Mouthon and Parghentanian (2004)
Portugal	River Minho estuary	16.5	840±387	11.89±0.06	3.32±0.05	3.06±1.20	2.28±0.30	7	Sousa et al. (2008b)
	Mondego estuary	6.9	5,017±5,489	7.67±0.04	–	–	–	8	Franco *et al.*, 2012
	Casal de São Tomé	10.3	3,255±1,115	4.06±0.04	3.14±0.12	3.83±1.69	2.75±0.51	9	Rosa et al. (2014)
USA	Altamaha River	4.5	712±1,613	0.37±0.00	–	–	–	10	Gardner et al. (1976)
	Lake Arlington	2.0	32±11	11.94±0.03	3.49±0.05	5.80±0.60	2.59±0.20	11	Aldridge and McMahon (1978)
	Delta-Mendota canal	28.0	9,470±7067	4.53±0.03	2.77±0.17	9.69±4.07	8.03±2.75	12	Eng (1979)
	New River (Station 1)	2	316±654	9.40±0.02	–	–	–	13	Graney et al. (1980)
	Clear Fork, Trinity River	7.0	4,474±3,899	19.34±0.04	3.00±0.16	6.76±3.62	7.07±2.83	14	McMahon and Williams (1986)
	Mechums River	10.5	669±636	188.52±0.62	2.56±0.15	9.20±4.41	3.31±1.21	15	Hornbach (1992)
	Ogeechee River	10.0	1,398±1,096	6.23±0.00	2.52±0.12	22.72±9.17	6.85±1.63	16	Stites et al. (1995)
	Lake Tahoe	8.0	1,809±723	16.64±0.04	2.80±0.12	6.90±1.90	2.70±1.32	17	Denton et al. (2012)

Table 2. Summary of results from reduced models of weighted linear regression analyses of invasion and growth variables of global invasive populations of *Corbicula*. Parameter values are means ± SE. We report one-tailed *P* values indicating significant results in bold characters. Note natural logarithmic transformations of variables. For details on descriptive statistics of full models and results on partial *F*-tests see the Supporting Information. All variables were Ln-transformed and standardized to zero mean and a unit of variance (see *Statistical Analyses* for details).

Response variable	Predictor	Estimate ± SE	Intercept ± SE	R^2	*F*-statistics	*num.df, den.df*	*one-tailed P*-value
Density	*Time*	0.454 ± 0.196	-0.906 ± 0.241	0.263	5.34	1, 15	**0.018**
Population growth rate	*Time*	0.936 ± 0.323	-0.518 ± 0.216	0.376	4.03	2, 14	**0.006**
	Density	-0.522 ± 0.407					0.220†
Individual growth rate	*Density*	-0.422 ± 0.265	0.098 ± 0.264	0.283	2.17	2, 11	0.070
	Population growth rate	-0.473 ± 0.344					0.099
minimum Age at sexual maturity	*Individual growth rate*	-0.505± 0.179	-0.192 ± 0.171	0.601	8.28	2, 11	**0.008**
	Population growth rate	0.460 ± 0.208					**0.024**
Lifespan	*Individual growth rate*	-0.718± 0.184	-0.056 ± 0.192	0.559	15.20	1, 12	**0.001**

† two-tailed *P*-value.

FIGURES

Figure 1. Global distribution of invasive *Corbicula* clams indicating the native range of the genus (dark grey), the distribution of *Corbicula* inside (black/grey shading and black dots) and outside of the genus' native range (color dots), which are mostly represented by forms A/R (formerly designed as *Corbicula fluminea sensu lato*; see main text for detailed explanation). Numbered stars are invasive worldwide populations reviewed in the present study. Years indicate estimated date of first introduction. See Table 1 and main text for references and details on the invasive populations reviewed here. See the Supporting Information for detailed criteria and data used to build the genus native and invasive distributions.

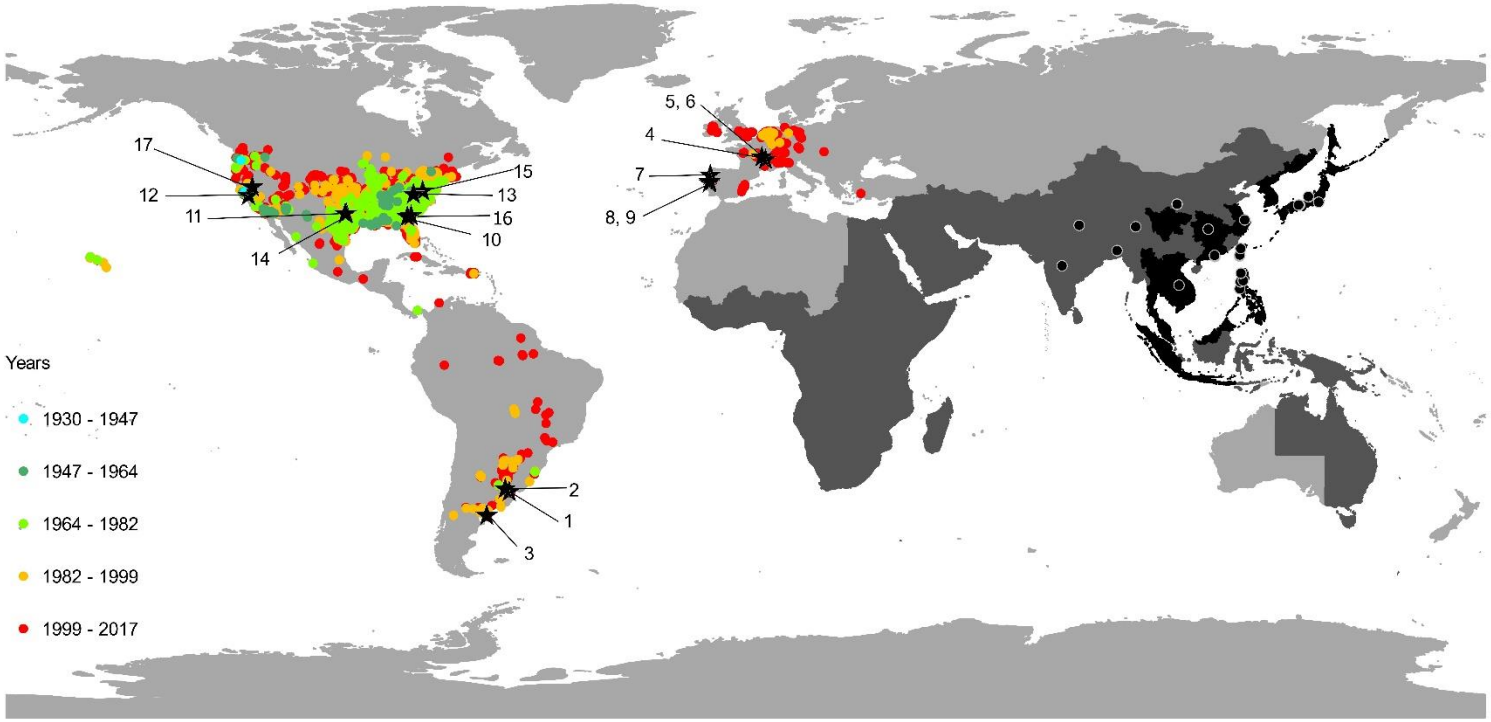

Figure 2. Relationships between processes and characteristics at different levels of organization of 17 global invasive populations of the Asian clam *Corbicula*. At the population level: a) the effect of time since colonization (*Time*, years) on population *Density* (ind./m^2) and b, c) the simultaneous effect of *Density* and *Time* on *Population growth rate*. At the interaction between population and individual level: d, e) the simultaneous effect of density and population growth rate on individual growth rate, and f) the effect of *Population growth rate* on the minimum *Age at sexual maturity* (months). At the individual level: g) the effect of *Individual growth rate* on the minimum *Age at sexual maturity*, and h) on the *Life span* (years). When plotting the fit for one variable after testing for simultaneous effects, we removed the effect of the variable not considered (b, c, d, e). Dashed lines show fits when marginally non-significant 0.05> α <0.10. Solid lines show linear regression fits when statistically significant at α < 0.05. The size of circles correspond to the inverse of the variance of mean value of each observation of the response variable, used to weigh effect sizes in the regression analyses. All variables were Ln-transformed and standardized to zero mean and a unit of variance (see *Statistical Analyses* for details).

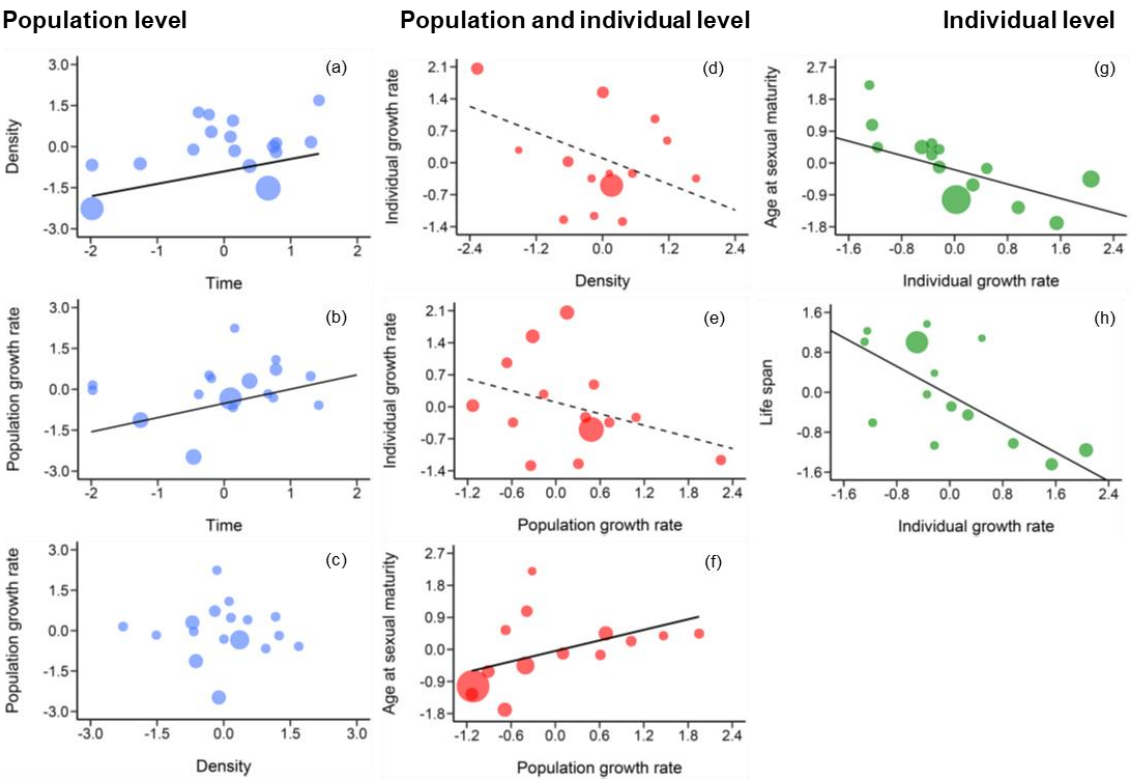
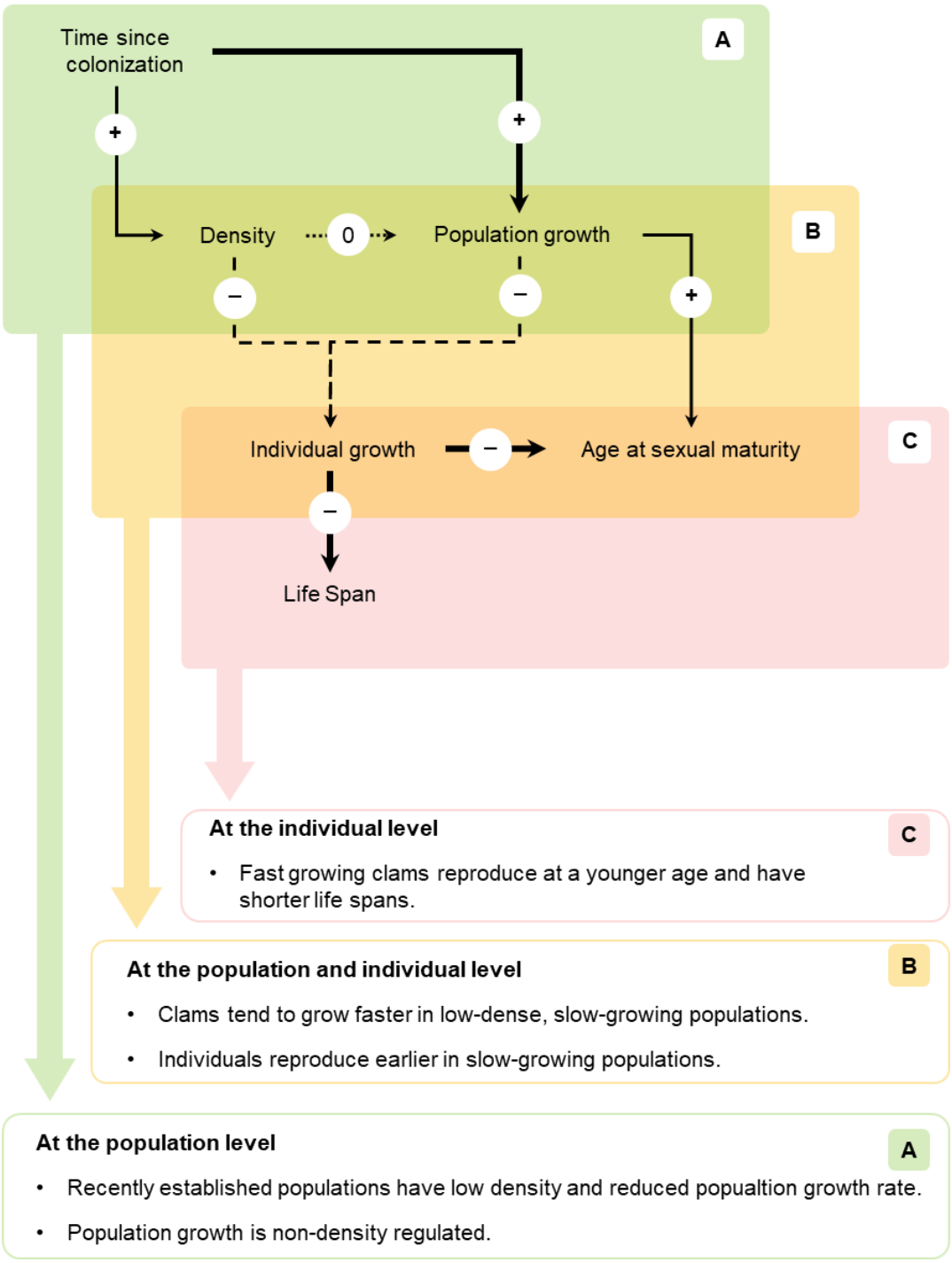
Figura 3. Summary of the relationships between processes and characteristics at different levels of organization of 17 global invasive populations of the Asian clam *Corbicula*, showing positive (+), negative (–), and no relationships (0) after fitting weighted linear regressions. Thick solid lines (⟶) show linear regression fits when statistically significant at $\alpha < 0.01$. Thin solid lines (⟶) show linear regression fits when statistically significant at $\alpha < 0.05$. Dashed lines (– –➤) show fits when marginally non-significant $0.05 > \alpha < 0.10$. Dotted line (·····➤) show no significant relationship $\alpha > 0.10$. Relationships are shown at three different levels: A) at the population level, B) the interaction between processes and characteristics at the population and individual level, and C) at the individual level.