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Distribution of native and non-indigenous bivalves and their settlers along an urban gradient

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Marine coastal habitats are often characterized by strong gradients of anthropogenic disturbance such as pollution, typically most severe at urban waterfronts. These variations create stress for local organisms leading to their distribution along the disturbance as a function of their tolerance. Non-indigenous species (NIS) are considered more tolerant to anthropogenic disturbances than their native counterparts, thriving in urban areas, where native species are sparser. It is however not yet entirely clear if these distribution patterns are due to larval behavior and preferential settlement or post-settlement processes. In the present study, we investigated the abundance of adults and settlers of two native (*Ostrea edulis*, *Mytilus galloprovincialis*) and two non-indigenous (*Magallana gigas*, *Xenostrobus securis*) bivalves along a strong marine urban environmental gradient. Oysters had sparse abundances of both adults and settlers at all sites, with no obvious distributional gradients. The two mussel species showed different settler-adult distributions along the gradient. Both settlers and adults of the native mussel *M. galloprovincialis* strongly decreased moving from the outermost periurban site to the innermost urban site, consistent with preferential settlement of larvae according to environmental conditions. The non-indigenous adult mussel *X. securis* showed a distribution pattern opposite to that of *M. galloprovincialis*, markedly increasing in abundance from the outermost to the innermost site. This was not paralleled by the distribution of *X. securis* settlers, which established over a larger area, including sites where adults were essentially absent. The mismatch between settler and adult distributions for *X. securis* suggests that post-settlement factors dictate spatial adult distribution.

KEYWORDS

marine urbanization, propagule pressure, invasive species, post-settlement, recruitment, metabarcoding, larvae, pollution

1 Introduction

Coastal urbanization is on the rise, with urban and periurban marine areas currently making up more than 1.5% of the global exclusive economic zones (Airoldi et al., 2021; Bugnot et al., 2021). Urbanization results in the permanent modification of abiotic and biotic conditions around artificial structures (Dafforn et al., 2015; Bugnot et al., 2021), often generating a variety of disturbance sources (Todd et al., 2019). Marine urban areas and other anthropic (*i.e.* manmade) habitats are in fact typically characterized by high levels of chemical pollution (Lam and Todd, 2013; Gauff et al., 2022), physical disturbance (Atalah et al., 2007), sediment load and turbidity (Airoldi, 2003), and increased temperature variability (Menniti et al., 2020) compared to non-urban areas. This set of specific conditions largely influences the distribution and functionality of organisms in these habitats.

The fouling communities colonizing the most urbanized artificial habitats differ from those of less impacted natural habitats, being characterized by distinct faunal and floral communities, lower species and genetic diversity (Fauvelot et al., 2009; Megina et al., 2016), altered ecological interactions (Klein et al., 2011) and functions (Dafforn et al., 2015) and a notorious prevalence of non-indigenous species (NIS) over native species (Mineur et al., 2012; Rondeau et al., 2022). NIS thrive in marine urban habitats due to their often-higher performance in stressful environments compared to native species (McMahon, 2002; Lenz et al., 2011). The underlying drivers of native and NIS distributions are, however, not yet fully understood (Airoldi et al., 2015; Gauff et al., 2022; Rondeau et al., 2022). Identifying the ecological forces and processes shaping marine biodiversity along gradients of urbanization is considered a priority for a better planning, design, and management of the most urbanized marine spaces, such as ports and city waterfronts (Airoldi et al., 2021), especially since NIS are considered a critical biosecurity issue (Pyšek et al., 2020), causing loss of biodiversity and ecosystem services (Walsh et al., 2016), species extinctions (Blackburn et al., 2019), and significant economic loss (Diagne et al., 2021).

Disturbance in marine urban spaces could be attenuated at the most flushed areas, where the hydrodynamic conditions create dilution effects (Schiff et al., 2007; Menniti et al., 2020). Flush attenuation induces disturbance gradients potentially affecting NIS prevalence (Floerl and Inglis, 2003; Gauff et al., 2022; Rondeau et al., 2022) and propagule pressure, which tends to be correlated to greater settlement and invasion success (Simberloff, 2009). Many marine taxa show active larval behaviors ranging from substrate preferences (Pinochet et al., 2020) to actively seeking out biofilm and/or conspecifics (Scheltema et al., 1981; Von Der Meden et al., 2010), which could vastly impact recruitment patterns. However, whether the distribution of NIS and native species along urban gradients is related to factors affecting the settlement or subsequent recruitment of propagules is not clear yet.

We took advantage of the urban gradients along the channel harbor of Ravenna, North Adriatic, Italy to explore whether the distribution of target adult NIS and native species colonizing artificial structures is related or not to settlement patterns

(Walters, 1992). We quantified how the gradient of urbanization affected a range of relevant environmental parameters, and then compared the distributions of adult and settlers of two native and two functionally similar NIS of the same family of bivalves along that environmental gradient. Based on previous knowledge (see above), we expected higher abundances of the native species at the most flushed periurban entrance of the harbor canal and higher abundance of the two NIS at the most confined inner waterfront. We hypothesized that these gradients in adult distribution would be paralleled by settler abundances as environmental constraints should lead to preferential settlement in suitable habitat. Alternatively, a larger settlement potential compared to the final adult distribution would suggest that other post-settlement factors such as predation or competition (often overlooked in marine urban areas) could better explain the NIS adult distribution.

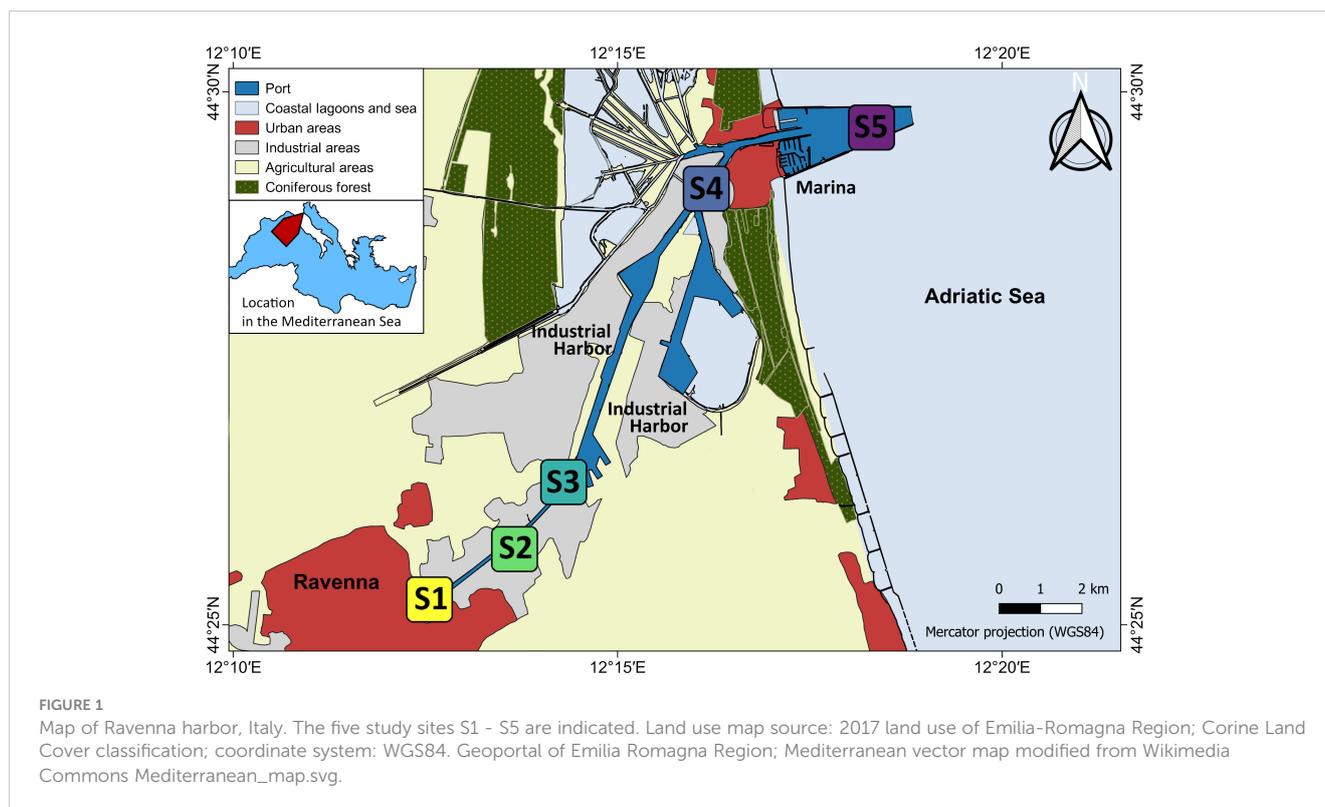
2 Materials and methods

2.1 Sampling site

The Ravenna harbor (Figure 1) is one of the largest industrial and commercial ports in Italy (Airoldi et al., 2016). It is an 11 km long channel going from the highly urbanized Ravenna city center, through a large industrial complex to the Adriatic Sea, where urban influence is diluted by high water circulation. Its entrance is protected by 2 large 2.4 km long converging jetties. The connection with the lagoon increases the tidal water flow, reducing the risk of siltation at the harbor entrance. We selected five study sites (Figure 1) along the inner-outer gradient representative of the different levels of urbanization and anthropogenic pressures, as well as different levels of confinement: the innermost, most confined, fully urban site (S1); three intermediate sites with decreasing intensity of urbanization and increased flushing (S2, S3, S4); and the outermost, more flushed, periurban site S5, which is on a long breakwater leading out to the sea and thus more distant from the urban center and its disturbances (for a definition of urban and periurban, see Airoldi et al., 2021). Due to access prohibition at the industrial harbor S3 and S4 are more distantly spaced than other sites.

2.2 Study species

We focused on four bivalve species with similar ecological niches and functions (mussels and oysters) occurring on the harbor seawalls, two of which were native and two of which were non-indigenous species (NIS). We expected the NIS to favor the innermost parts of the canal as seemingly specifically adapted to and dominant in urban habitats (Gauff et al., 2022; Rondeau et al., 2022). *Mytilus galloprovincialis* Lamarck, 1819 is a Mediterranean native species (Montes et al., 2021). Its non-indigenous counterpart *Xenostrobus securis* (Lamarck, 1819), is an Australian mussel species that was introduced in the north-western Adriatic Sea in 1994 (Garci et al., 2007). The native oyster *Ostrea edulis* Linnaeus, 1758 is a pan-European oyster species that was originally abundant



from the low intertidal into water depths of about 80 m, but it was replaced by the Pacific oyster *Magallana gigas* (Thunberg, 1793) introduced for aquaculture purposes. At the end of the dispersal phase, all four species colonize intertidal and shallow subtidal habitats, including artificial ones (Bataller et al., 1999; Gomes et al., 2016; Stagličić et al., 2020; Montes et al., 2021).

2.3 Sampling design and treatment

At each study site, environmental conditions were monitored through time, mussel and oyster adults were sampled once, and their settlement was followed through the entire study. Environmental conditions were monitored at each site from March to July 2017 on a two-week basis, for a total of eight times (T0 - T7). Seawater pH, temperature, salinity, chlorophyll-*a*, dissolved oxygen, and turbidity were measured using a multiparametric probe (Hanna Instruments Hi 9828), and water samples were collected for subsequent analysis of ammonia, nitrate, nitrite, and phosphate in the laboratory. Six replicates of all variables were collected, except for T0, where only one replicate was taken at each site.

At T0 only, six replicated 20 × 20 cm samples were randomly scraped at each site to quantify the abundances of the four target species. Due to diving permitting constraints, sampling was conducted from a small boat, selecting days with low tidal conditions (average tidal amplitude is ~ 80 cm), to reach and sample the low intertidal seawall habitats (~0.1-0.3 m above Mean Low Waterlevel). Each sample was preserved in plastic bags and rapidly processed in the laboratory. Abundance was defined as

the number of live specimens belonging to one of the four target species, identified under a stereomicroscope (Garci et al., 2007).

The scraped surfaces at each site were used to deploy six replicated Artificial Substrate Units (ASU), consisting of polypropylene kitchen scouring pads serving as bivalve spat-collectors (see Porri et al., 2006). Each ASU was retrieved and replaced every 2 weeks from March (T1) to July (T7), for a total of $n = 210$ samples (6 replicates per 5 sites per 7 times). Upon retrieval, each ASU was placed into a sterile plastic bag and processed in the laboratory following the protocol described by Porri et al. (2006). Bivalve settlers (~ 360 μm; Porri et al., 2006) were sorted out and counted under a stereomicroscope (Nikon SMZ 15000), and further identified and quantified by molecular approaches. On replicates containing 1 to 4 settlers, individuals were barcoded for species identification, while on replicates with more than 4 settlers, we used a metabarcoding approach. In both cases, the cytochrome oxidase subunit I (COI) gene was targeted, and the number of reads was considered as proxy for settler abundance. The detailed protocol for molecular analyses is described in the [Supplementary Material](#).

2.4 Statistical analysis

All statistical analyses were carried out in 'R' (R Core Team, 2020) and figures obtained via 'ggplot2' (Wickham, 2016). All logistic regressions were applied and interpreted according to Fox and Weisberg (2018).

Each environmental variable was plotted for each site and time (See [Supplementary Material](#), [Supplementary Figures 1–10A](#)). To investigate potential environmental gradients across sites,

the environmental variables were further scaled and analyzed for each time using Principal Component Analysis (PCA) via the 'vegan' package (Oksanen et al., 2018; Supplementary Figure 11). Some PCAs required flipping to align the site distribution among different time steps. The centroid PC1 value was then extracted for each site at each time, and the overall gradient was represented through this PC1 value as a function of the site. Further details can be found in the Supplementary Material. Subsequently, a logistic regression was employed to test for the effect of site (ordered discrete factor) on the PC1 value.

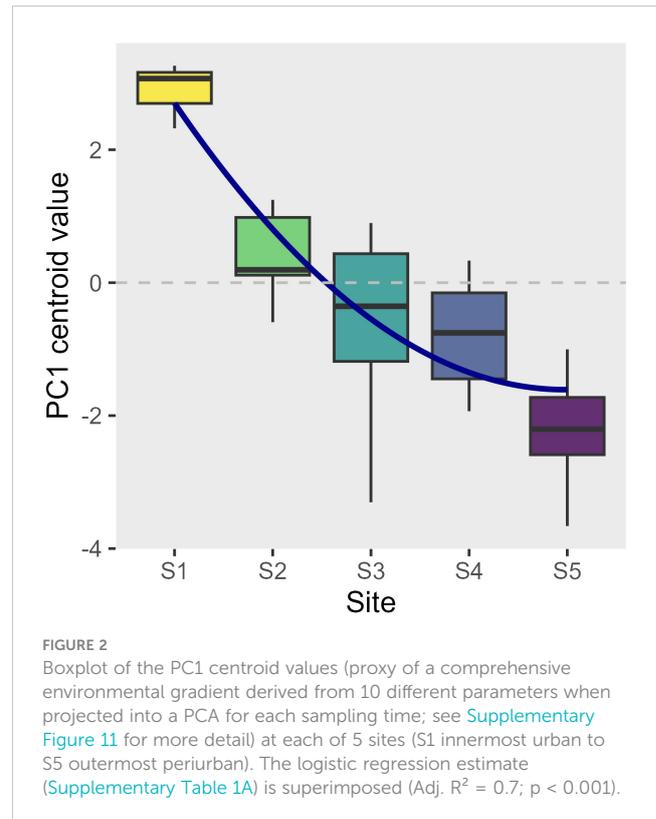
The abundances (number of adult individuals) of *M. galloprovincialis* (Native), *X. securis* (NIS), *O. edulis* (Native), and *M. gigas* (NIS) per sample were box-plotted for each site at T0, and logistic regressions (with site as ordered discrete factor) were used to identify their distributions among sites. If the overall model for any species was significant, the fitted model with all terms of the logistic regression (i.e., linear L, quadratic Q, cubic C, 4th degree) was superimposed onto the box plot distribution of the species.

Similarly, the numbers of newly settled larvae were plotted and analyzed as for the adults. As virtually no oyster settlers were found (see Results), these analyses were run only for mussel settlers. Since settlers were sampled seven times, we used a logistic linear mixed-effects model (LMM) via REML from the 'lme4' package (Bates et al., 2015) to account for time as random factor.

3 Results

Environmental factors varied across sites and times (Supplementary Figures 1–11). In general, the urban innermost site S1 was characterized by the highest temperature (1 to 3 degrees higher compared to the most external site S5, depending on the time of the year), salinity, and turbidity, which decreased moving towards the outermost periurban site, while nutrient levels tended to peak at the most industrial sites S3 and S4. When all the environmental data were combined, the PCAs for each time showed that overall, a temporally stable environmental gradient structure was present, with a high differentiation across sites from S1 to S5 (Supplementary Figure 11). The PC1, based on the 10 environmental variables, explained between 31.2% and 50.2% of the data distribution, and the boxplot of the PC1 centroid value showed an ordered structure of the sites (Figure 2). The logistic regression, testing for the effect of site on the PC1 centroid value, revealed a significant linear decrease (L: t-val. = -9.06; Pr (> |t|) < 0.001; Figure 2) with deceleration moving towards the outermost site (Q: t-val. = 2.714; Pr (> |t|) = 0.01; Figure 2). The overall model significance was high (Adj. R² = 0.7; F-value = 23.4; 35 DF; p < 0.001).

Oysters were sparse at all study sites, with *O. edulis* virtually absent in most samples, and *M. gigas* sparsely occurring at only few plots at sites S2, S3, and S4 (Supplementary Figure 12). Like adult abundances, oyster settler abundances were unexpectedly low/absent based on molecular data. During the 7 sampling times, a total of only two settlers of *M. gigas* and no settlers of *O. edulis* were



found on the ASUs, which prevented further analyses for these species.

The adults of the two mussel species (*M. galloprovincialis* and *X. securis*) significantly varied across sites, showing opposite patterns along the gradient (Adj. R² > 0.66; F-value > 15.27; 25 DF; p < 0.001; Figures 3A, B; Table 1). A significant increase of the native *M. galloprovincialis* was observed moving from the innermost waterfront to the outermost periurban areas (L: t-val. = 9.5; Pr (> |t|) < 0.001; Figure 3A; Table 1) accelerating (Q: t-val. = 6.47; Pr (> |t|) < 0.001; Figure 3A; Table 1) towards the outermost periurban site. In this model, the cubic term was significant (C: t-val. = 3.26; Pr (> |t|) = 0.003; Figure 3; Table 1) indicating the presence of a threshold at which the acceleration starts centered between S3 and S4. For the non-indigenous *X. securis*, the distribution of adults was opposite to that of *M. galloprovincialis*, with a significant linear decrease from the innermost waterfront to the outermost periurban areas (L: t-val. = -7.2; Pr (> |t|) < 0.001; Figure 3B; Table 1).

The distribution of *M. galloprovincialis* settlers over 7 sampling times paralleled the adult distribution, while there was a notable mismatch between adult and settler abundance for *X. securis*. Like for adults, *M. galloprovincialis* settlers had an increasing (L: t-val. = 4.78; Pr (> |t|) < 0.001; Figure 3C; Table 1) and accelerating (Q: t-val. = 2.99; Pr (> |t|) = 0.003; Figure 3C; Table 1) abundance towards the outermost periurban site. For *X. securis*, on the other hand, only the quadratic polynomial had significant effect (Q: t-val. = -2.83; Pr (> |t|) = 0.005; Figure 3D; Table 1) reflecting maximum abundances in the middle portion of the gradient and very low abundances at the periurban site. This notably deviates from the adult abundances,

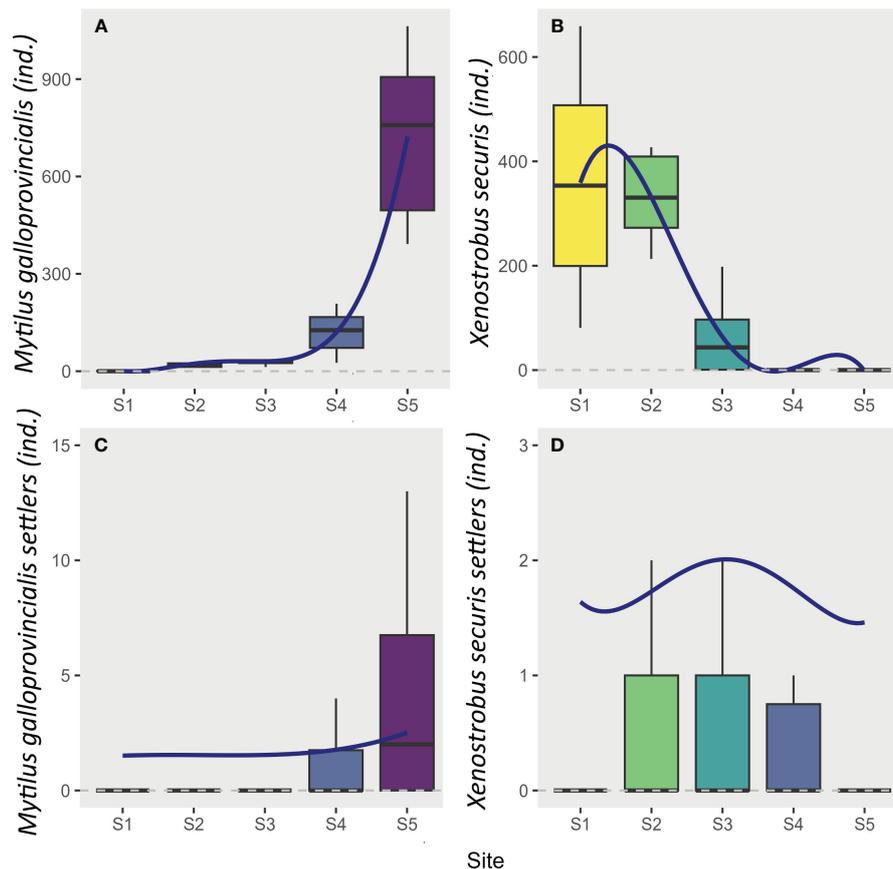


FIGURE 3

Boxplots of the abundances (number of individuals per sample) of (A) adults of *Mytilus galloprovincialis* (Native), (B) adults of *Xenostrobus securis* (NIS), (C) settlers of *Mytilus galloprovincialis* (Native), and (D) settlers of *Xenostrobus securis* (NIS) at each site. The settler abundances are an average over 7 sampling times, from March to July 2017. Logistic regression estimates (see Table 1) are superimposed when a model was significant.

which were highest in the two innermost sites. This mismatch is especially striking for the innermost waterfront site, which had the highest adult abundances, but second lowest settler abundances (after the periurban site).

4 Discussion

In the present study, we investigated whether a gradient of increasing urbanization structured the distribution and abundances of adult populations of four target bivalves and their settlers. The periurban-urban gradient observed at the harbor of Ravenna was reflected in a temporally stable, significant gradient in relevant environmental parameters, including temperature, salinity, turbidity, and nutrient concentration, with consistent dissimilarities among the outermost and inner sites (Figure 2). This occurred despite strong temporal fluctuations in the individual parameters (Supplementary Figures 1–10). Only two of the four tested adult bivalve species were significantly influenced by this gradient, however, the distribution profiles align with our hypothesis: the NIS was more abundant in the inner parts of the canal, while the native species dominated the entrance. For settlers, only one species (*M. galloprovincialis*) exhibited a distribution

profile that overlapped with adult distribution, indicating preferential settlement in habitats suitable for recruitment. *Xenostrobus securis* displayed a notable mismatch between adult and settler abundances, potentially linked to more indiscriminate recruitment owing to its resistance to disturbance.

Environmental conditions in the canal changed between March and July, as expected for both natural and urban areas (Lacroix et al., 2017). However, in accordance with hypotheses focusing on the importance of hydrodynamics and water exchange with the open sea (Floerl and Inglis, 2003; Schiff et al., 2007; Menniti et al., 2020), we observed a gradient structure in environmental conditions. This gradient was found to be temporally stable for individual parameters such as temperature ($T_{S1} > T_{S5}$; Supplementary Figure 1), but also for the multivariate distribution of combined environmental conditions, with consistently high dissimilarity between S1 and S5 on the PC1 axis (Figure 2; Supplementary Figure 11). This gradient structure was however not entirely linear, as a slight deceleration trend for differences between sites was observed between the middle and the outermost site (Figure 2; Table 1), indicating that environmental conditions exhibit steeper spatial variations at the innermost part of the canal.

In Ravenna harbor, oysters showed surprisingly low abundance compared to similar marine urban areas in the northern Adriatic

TABLE 1 Summary of the logistic regressions and the linear mixed effect models testing for the effect of site (ordered discrete factors) on different response variables of interest.

PC1 value of Centroids in environmental variables PCA			
Predictors	Estimates	t-value	Pr(> t)
(Intercept)	-0.00	0	1.000
Site [linear]	3.41	9.057	< 0.001
Site [quad.]	-1.02	-2.714	0.010
Site [cubic]	0.75	2.002	0.054
Site [4th deg.]	0.17	0.442	0.661
Observations	40	F-stat.	23.4
R ²	0,728	p-value	< 0.001
R ² adjusted	0,697		

<i>Mytilus galloprovincialis</i> adults				<i>Mytilus galloprovincialis</i> settlers			
Predictors	Estimates	t-value	Pr(> t)	Predictors	Estimates	t-value	Pr(> t)
(Intercept)	179.43	7.821	< 0.001	(Intercept)	1.78	2.247	0.026
Site [linear]	487.41	9.501	< 0.001	Site [linear]	4.25	4.779	< 0.001
Site [quad.]	331.76	6.467	< 0.001	Site [quad.]	2.66	2.994	0.003
Site [cubic]	167.28	3.261	0.003	Site [cubic]	1.03	1.161	0.247
Site [4th deg.]	39.60	0.772	0.447	Site [4th deg.]	0.04	0.042	0.967
Observations	30	F-stat.	35,85	Observations	210	Random Effects	
R ²	0.851	p-value	< 0.001	Marginal R ²	0.126	σ ²	33.16
R ² adjusted	0.828			Condit. R ²	0.205	τ _{00 Date}	3.27
						ICC	0.09
						N _{Date}	7

<i>Xenostrobus securis</i> adults				<i>Xenostrobus securis</i> settlers			
Predictors	Estimates	t-value	Pr(> t)	Predictors	Estimates	t-value	Pr(> t)
(Intercept)	151.73	7.389	< 0.001	(Intercept)	1.72	1.559	0.121
Site [linear]	-330.77	-7.203	< 0.001	Site [linear]	-0.62	-0.845	0.399
Site [quad.]	67.88	1.478	0.152	Site [quad.]	-2.09	-2.83	0.005
Site [cubic]	93.92	2.045	0.051	Site [cubic]	-0.44	-0.6	0.549
Site [4th deg.]	-71.28	-1.552	0.133	Site [4th deg.]	0.86	1.158	0.248
Observations	30	F-stat.	15.17	Observations	210	Random Effects	
R ²	0.708	p-value	< 0.001	Marginal R ²	0.036	σ ²	22.99
R ² adjusted	0.661			Condit. R ²	0.279	τ _{00 Date}	7.74
						ICC	0.25
						N _{Date}	7

For *Mytilus galloprovincialis* settlers and *Xenostrobus securis* settlers, time was computed as a random factor to account for different sampling dates. Significant p-values in bold.

(see photographic comparison with an urban substrate from the Venice Lagoon: [Supplementary Figure 12](#)). In total, only 35 live adult oysters were sampled, with the majority (27) being *M. gigas*. Oyster settlers were virtually absent during the study, despite the study period coinciding with their spawning period (Bataller et al., 1999; Massapina et al., 1999). Both oyster species have been documented in estuaries, harbors, and marinas (Stagličić et al., 2020), including in the study area and nearby lagoons (Airoldi et al.,

2016), and should be able to resist the environmental conditions in the area. Transient disturbances could have nevertheless led to mass mortality of oysters, possibly replaced by faster-growing mussels, *X. securis*, in the innermost sites, and *M. galloprovincialis* in the outermost sites respectively.

Expectedly, adult distributions reflected their status (NIS vs Native), with *M. galloprovincialis* (Native) increasing towards the periurban part of the canal (S5), and *X. securis* (NIS) decreasing.

This dichotomous distribution aligns with observations made by Montes et al. (2021) in a Spanish harbor, where a combined effect of spatially organized features (*i.e.*, environmental gradients) and seasonal dispersal barriers were hypothesized as underlying drivers. In our study, for *M. galloprovincialis*, a linear trend was accompanied by an acceleration setting at a threshold (Quadratic term; Table 1), centered between S3 and S4. Between S3 and the canal's entrance, the abundance of adults increased at a high rate. The distribution pattern of its settlers closely mirrored this trend. Given that the length of the canal falls well within the typical dispersal distance of *M. galloprovincialis* (Gomes et al., 2016), this may indicate a preferential settlement as it tightly matched the adult distribution. *Mytilus galloprovincialis* adults and larvae are known to be controlled by their tolerance to environmental parameters, including temperature, salinity (Olabarria et al., 2016; Vasquez et al., 2022), and some sources of pollution (Serrano et al., 1995; Boukadida et al., 2016). Environmental conditions between S1 and S3 may not be favorable to *M. galloprovincialis* and may constitute a limiting factor for their settlement and survival. The distribution profile of *M. galloprovincialis* adults and settlers was similar to the PC1 value profile used as a proxy to outline the environmental gradient, with S4 and S5 markedly different from S3, S2 and S1 (Table 1; Figure 1). It is likely that *M. galloprovincialis* distribution was mostly influenced by its inability to thrive in the environmental conditions of inner urban areas and its preference for more suitable environments for settlement.

Decreasing from the innermost waterfront to the periurban site, the abundance of *X. securis* adults was consistently high at the two innermost sites, while it was almost absent at the outermost ones (Figure 2; Table 1). Unlike *M. galloprovincialis*, where the distribution of settlers closely mirrored that of the adults, *X. securis* displayed a significant discrepancy between the abundance of settlers and adults. Settlers of *X. securis* were consistently spread out in the canal, with the highest abundance found in the middle areas, where adult populations were already decreasing. The mismatch between adult and settler abundances could indicate that for this species, post-settlement processes could drive the adult population distribution patterns as settlers colonize a broader range than that to what the adult population was limited. One such post-settlement process could be predation from boring snails, crabs, and fish, which depending on the environmental conditions and characteristics of the mussel aggregates, can reduce *X. securis* survival (Gestoso et al., 2014), and consequently adult populations. Another post-settlement process affecting *X. securis* distribution could be competition with the native species *M. galloprovincialis* at the outermost part of the canal. Despite *in aquaria* experiments revealing a high resistance to a wide range of temperatures, pH, and salinities (Garci et al., 2007; Gestoso et al., 2016; Olabarria et al., 2016) *in situ*, *X. securis* seems restricted to the upper estuarine sites with low salinities (Garci et al., 2007; Montes et al., 2021). The general high resistance of NIS to multiple and variable environmental stressors (Lenz et al., 2011), which also applies to *X. securis* (Gestoso et al., 2016; Olabarria et al., 2016; Astudillo et al., 2017) could explain its high abundance at the inner canal sites. However, stress resistances come with costly metabolic

trade-offs (Pook et al., 2009; Gauff et al., 2022). This metabolic challenge could lead to decreased competitiveness in environments with lower disturbance and may explain why numerous NIS thrive in high disturbance habitats like anthropic habitats or estuaries but are far less common in nearby natural sites (Airoldi et al., 2015; Gauff, 2021). Other species could also compete with *X. securis*, but keystone bivalves were selected for this study because they are the most abundant species in the study area, thus it seems unlikely that other species compete too strongly with *X. securis*.

Overall, our data support the hypothesis that NIS are more prevalent in disturbed habitats (Piola and Johnston, 2008; Megina et al., 2016; Rondeau et al., 2022), potentially due to unfavorable post-settlement competitive matchups against similar native species in less disturbed environments (Montes et al., 2021; Gauff et al., 2022), and/or due to specialization into resistant life strategy traits implying certain metabolic trade-offs (Gauff, 2021; Gauff et al., 2022). Future work on larval stages and settlers of a greater range of species is worth pursuing to test the generality of this hypothesis and to identify the underlying drivers, by disentangling biotic interactions from abiotic parameters like temperature, salinity, and chlorophyll-*a*, which considerably varied in our experiment and could have impacted both species (Gestoso et al., 2014; Olabarria et al., 2016; Montes et al., 2021). This should include tests on competition between native and NIS with similar niches, as well as other post-settlement processes like predation or seasonal post-settlement mortality (Montes et al., 2021), which could also lead to the discrepancies observed here.

Data availability statement

The datasets are available within the [Supplementary Material](#) and scripts for this study will be provided upon reasonable request.

Ethics statement

Ethical approval was not required for the study involving animals in accordance with the local legislation and institutional requirements because not needed for these specimens.

Author contributions

RG: Data curation, Formal Analysis, Visualization, Writing – original draft, Writing – review & editing. FM: Data curation, Investigation, Methodology, Writing – original draft, Writing – review & editing. PM: Conceptualization, Methodology, Supervision, Writing – original draft, Writing – review & editing. FP: Conceptualization, Methodology, Writing – review & editing. FC: Conceptualization, Funding acquisition, Methodology, Project administration, Supervision, Writing – review & editing. LA: Conceptualization, Funding acquisition, Methodology, Project administration, Supervision, Writing – review & editing.

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Conflict of interest

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Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fmars.2024.1401552/full#supplementary-material>

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