

# Rapid recovery of native habitat-builders following physical disturbance on pier pilings offsets colonization of cryptogenic and non-indigenous species in a Chilean port

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Jean-Charles Leclerc, Antonio Brante, Frédérique Viard. Rapid recovery of native habitatbuilders following physical disturbance on pier pilings offsets colonization of cryptogenic and non-indigenous species in a Chilean port. Marine Environmental Research, 2021, 163, 10.1016/j.marenvres.2020.105231. hal-03321963

## HAL Id: hal-03321963 https://hal.science/hal-03321963

Submitted on 18 Aug 2021

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1	Rapid recovery of native habitat-builders following physical disturbance on pier pilings
2	offsets colonization of cryptogenic and non-indigenous species in a Chilean port.
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#### 13 Abstract

Examining the effects of disturbances within marine urban communities can shed light 14 on their assembly rules and invasion processes. The effects of physical disturbance, through the 15 removal of dominant native habitat-builders, were investigated in the recolonization of 16 disturbed patches and colonization of plates on pier pilings, in a Chilean port. On pilings, 17 disturbance substantially affected community structure after 3 months, although it slowly 18 19 converged across treatments after 10 months. On plates, cryptogenic and non-indigenous species richness increased with removal severity, which was not observed in natives. 20 Opportunistic taxa took advantage of colonizing at an early successional stage, illustrating a 21 22 competition-colonization trade-off, although indirect effects might be at play (e.g. trophic competition or selective predation). Recovery of the habitat-builders then occurred at the 23 expense of cryptogenic and non-indigenous taxa. Whether natives could continue winning 24 against increasing propagule and colonization pressures in marine urban habitats deserves 25 further attention. The interactions between disturbance and biological invasions herein 26 27 experimentally shown in situ contribute to our understanding of multiple changes imposed by marine urbanization in a growing propagule transport network. 28

#### 29 Keywords

30 Biofouling, Sessile invertebrates, Succession, Invasions, Cosmopolitanism, Urban ecology,

31 Artificial structures, SE Pacific

33 Introduction

Disturbances take multiple forms and regimes, and significantly affect population 34 dynamics and community assembly (Connell and Slatyer 1977; Sousa 1979; Shea and Chesson 35 2002; Turner 2010; Zelnik et al. 2019). Human activities influence the type, the frequency and 36 intensity of disturbances, as well as the way they interact with each other in space and time. 37 These disturbances can be particularly diverse and intense in urban areas (e.g., habitat 38 modification, maintenance work, pollution, resource exploitation, and biological invasions), 39 which are predominantly located along coastlines (Geraldi et al. 2019; Todd et al. 2019). At 40 multiple spatial scales, marine urbanization may profoundly affect community assembly and 41 the functioning of coastal marine and estuarine ecosystems (Dafforn et al. 2009; Duarte et al. 42 2013; Malerba et al. 2019; Todd et al. 2019). 43

Marine urban habitats, such as ports and marinas, may have long constituted key fertile 44 grounds for taxonomic and functional homogenization on a global scale. Indeed, they constitute 45 highly similar (and disturbed) environments, embedded in a transportation network of 46 propagules, and, in turn, form invasion hubs (Carlton 1996; Floerl et al. 2009; Johnston et al. 47 2017; Darling and Carlton 2018). The multiplication of artificial structures have notably been 48 shown to promote the intensification of gelatinous plankton blooms (Duarte et al. 2013; 49 Vodopivec et al. 2017). In a way, this follows two twin hypotheses of invasion biology (cf. 50 51 review by Enders et al. 2020), namely "human commensalism" (Jeschke and Strayer 2006) or "habitat legacy" (Pyšek et al. 2015), wherein opportunist and stress-tolerant taxa are likely to 52 invade networks of disturbed habitats, with consistent habitat affinities between source and 53 recipient regions. Similarly, other stress-resistant fouling taxa are widely distributed and 54 conspicuous in ports, regardless of their non-indigenous versus native species status locally 55 (Clark and Johnston 2009; Dafforn et al. 2009; Canning-Clode et al. 2011; Ferrario et al. 2020). 56

All taxa inhabiting these marine urban habitats are candidates for transport and may possess 57 58 traits allowing them to overcome associated stresses (anoxia, pollutants); it is therefore of little surprise to find a high and growing prevalence of non-indigenous species (NIS) in these 59 habitats. Although disturbance sources are diverse in marine urban systems (see Todd et al. 60 2019 for review), it is worth noting that even the initial deployment of novel artificial structures 61 constitutes a pulse physical disturbance (e.g. Connell and Slatver 1977 and references therein), 62 initiating a novel (secondary) succession sequence for this pervasive kind of marine 63 community. In these systems, community responses to disturbances can provide important 64 insights into assembly rules, especially in the context of ever-increasing biological 65 66 introductions (Shea and Chesson 2002; Pearson et al. 2018a).

In the last two decades, the SE Pacific coastline has been profoundly affected by ocean 67 sprawl (i.e. the proliferation of artificial structures at sea, Duarte et al. 2013; Bishop et al. 2017). 68 Artificial structures have been expanding either to protect inhabited shores from storms and 69 tsunamis or to promote economic development, through shipping, mining and aquaculture etc. 70 71 (Aguilera 2018). Coastal urban communities of the SE Pacific do not only differ in composition and structure as compared to neighboring communities of natural habitats. They also suffer 72 from diverse stressors, such as the accumulation of contaminants (plastics, pollutants, effluents) 73 74 (Aguilera et al. 2019), the loss of structuring interactions (e.g. predation) across the modified seascape (Dumont et al. 2011; Leclerc et al. 2020b), the introduction and spread of NIS (Castilla 75 and Neill 2009), and both legal and illegal fishing of a wide array of resources, including fish, 76 benthic predators, kelp and sessile invertebrates (Siegfried et al. 1994; Aguilera et al. 2019). 77 78 Despite the pressures undergone by marine urban communities, abundant populations of native 79 species have also been found to be associated with some of these artificial habitats. Depending how they use resources (e.g. space pre-emption, predation), native species may help reduce the 80 establishment and spread of NIS (Dumont et al. 2011; Leclerc et al. 2020a; 2020b). A 81

manipulative field experiment conducted in three ports of central Chile recently supported this 82 83 scenario by bringing evidences that benthic predators (mostly invertebrates) could – as far as their diversity is maintained – efficiently regulate NIS richness and abundances, hence favoring 84 competitive exclusion by native fouling species over succession (Leclerc et al. 2020a). 85 Interestingly, these native fouling species constitute a three-dimensional habitat for the very 86 same predators (Leclerc et al. 2020a), therefore habitat formation may reinforce biotic 87 resistance in these communities. These fouling species (giant barnacles, ascidians, mussels) are 88 also commonly fished and it is still unknown to what degree this unregulated harvesting (and 89 subsequent disturbances for associated communities) could affect such a positive feedback. 90

91 In the present study, we determined the effects of the removal of dominant native habitat-builders, with varying disturbance severity, on community assembly. We used a 92 manipulative experiment to examine the recolonization of established assemblages within 93 manipulated patches (varying surface areas were available across treatments) as well as the 94 colonization of settlement plates (standardized for available surface areas), on pier pilings. 95 96 Given the proximity of shipping traffic and prior ecological knowledge of our study site (see methods), we hypothesized (H1) that the severity of disturbance would induce contrasting 97 colonization responses, with greater prevalence of opportunistic cryptogenic and non-98 99 indigenous species when larger patches were available. We further hypothesized (H2) that such treatment effects would be greater on plates (controlled for substrate availability) compared to 100 pilings due to indirect effects, such as the loss of biogenic habitats used by local predators 101 (Leclerc et al. 2020a). Post disturbance assemblages were surveyed at both early and late stages 102 103 of succession to examine community composition resilience (cf. e.g. Hillebrand et al. 2018), 104 which was hypothesized (H3) to vary negatively with the severity of the disturbance.

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#### 106 Materials and methods

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### 108 Study site and associated community

This study was performed in the port of San Vicente Bay (36.7591°S, 73.1551°W), 109 central-southern Chile. In this Bay, environmental conditions are fairly stable across seasons, 110 with temperatures – averaged over 20 m depth CTD profile at its mouth (in 1996) – ranging 111 from 10°C in the winter to 13°C in the summer (Rudolph et al. 2002), although this range tends 112 to be greater in shallow waters (from 10 to 15°C measured at 4 m over 2016-2018 surveys, 113 Leclerc et al. 2018 and JCL pers. obs.). Heavy metal concentrations are generally greater than 114 in adjacent pristine locations, with an average of 16.0, 18.2 and 46.1 mg kg<sup>-1</sup> found in the 115 116 sediment for copper, lead and zinc, respectively (Leclerc et al. 2018). At the edges of San Vicente Bay, water quality (organic matter, turbidity, dissolved oxygen) may be highly variable 117 over the year, notably due to changes in the wind regime and organic matter effluents from 118 rivers and local fishing industries (Mudge and Seguel 1999; Rudolph et al. 2002). This Bay (ca. 119  $3 \times 4$  km) also constitutes one of the main hotspots of maritime traffic of the conurbation of 120 121 Concepción. In 2016, a total of 427 foreign commercial ships and 217 national commercial ships made stopovers, while a total of 162 fishing crafts moored in San Vicente Bay (Leclerc 122 et al. 2020b). 123

The center of San Vicente Bay is crossed by a private, ca. 2 km long steel dock used as 124 a gas pipeline connecting carriers with the local fuel industry (Abastible ©). Previous studies 125 in the area (Leclerc et al. 2018; Leclerc et al. 2020a; 2020b) have shown massive fouling by 126 large native barnacles (Austromegabalanus psicattus) and ascidians (Pyura chilensis) on the 127 pilings of this dock (Fig. 1C). According to repeated rapid assessment surveys on established 128 communities, these ascidians and barnacles remain common to super-abundant (SACFOR 129 scale), respectively, all yearlong (Leclerc et al. 2018). Patterns of dominance by the same 130 habitat-builders are actually common on artificial structures throughout the temperate SE 131

Pacific, with consistent observations up to 800 km northwards and 700 km southwards 132 (Valdivia et al. 2005; Manríquez et al. 2014; Leclerc et al. 2020b). The relatively long life 133 duration of these species may significantly contribute to these patterns, although cohort 134 monitoring beyond the ages of 18-20 months are, to the best of our knowledge, lacking 135 (Manríquez and Castilla 2005; López et al. 2008; Leclerc et al. 2020a). Multiple surveys of 136 settlement plates in our study site over the period 2016-2018 indicated a fair convergence (with 137 no signs of attrition) after 18 months, regardless of the timing of their deployment (winter, 138 summer), towards the dominance of these two species along with colonial organisms growing 139 either on the primary surface of the piling or as epibionts of these habitat builders (e.g. the 140 141 hydroid Amphisbetia operculata and the bryozoan Alcyonidioides mytilii, see also Table S1) (Leclerc et al. 2020a). The seasonality of these communities is rather marked in smaller 142 associated species (epibionts or not), showing massive recruitment and/or vegetative growth 143 periods (e.g. the barnacle Balanus laevis, the hydroids Obelia dichotoma and Halecium sp. as 144 well as filamentous turfs such as Ectocarpales or Antithamnionella sp., Table S1 and Leclerc et 145 146 al. 2020a). Most cryptogenic and non-indigenous species are poorly-represented in the mature fouling communities of these pilings throughout the year, unless they are released from the 147 influence of predators, either through exclusion cages (Leclerc et al. 2020a) or because of the 148 presence of inaccessible cavities, such as those formed by the arrangement of experimental 149 panels and plates (Leclerc et al. 2018, and JCL pers. obs.). 150

151

152 Study design and data collection

The manipulative experiment aimed at simulating the 'severity' of a physical disturbance (i.e. the effect of a disturbance event on the community), which is closely related to the disturbance 'intensity' (the energy associated with this event, per area per time, sensu Turner 2010). The experiment took place in December 2017 and consisted of the control (0%),

partial (50%), and complete (100%) removal of the dominant habitat-builders, from the surface 157 to the seafloor (circa 8 m depth) of distinct pilings along the dock described above (Fig. 1A, 158 B). In the partial removal treatment, dominant organisms were scraped from the pilings along 159 160 45° inclined parallel strips of ca. 20 cm widths, separated by the same width. The manipulation was replicated in two areas (hereafter blocks) separated by ca. 200 m, within which one piling 161 was randomly assigned to one of each treatment. Within each block, treatment pilings were 162 separated by ca. 20-40 m (Fig. 1A). The removal of sessile taxa was conducted by commercial 163 divers (Prosub Ltd) using a large customized spatula (10 cm width) welded to a 150 cm steel 164 stick (Fig. 1B) that they commonly employ for maintenance works in the submerged 165 166 infrastructures of the region. It is noteworthy that these scraping tools are also similar to those employed by fishermen to harvest the edible fouling taxa, which include the giant barnacle and 167 ascidian habitat-builders (up to 10 cm width) that were targeted in our removal experiments. 168 169 We thus simulated the realistic physical disturbance 'intensity' experienced by these communities through seafood harvesting. Such hard fieldwork implied, however, some 170 171 imprecision in the intended disturbance effect, i.e. the 'severity' (Turner 2010). In fact, several species associated with the three dimensional habitat formed by these dominant species were 172 visibly damaged or removed from the piling surface following manipulation, therefore our 173 results may rather be interpreted as the outcome of a pulse physical disturbance on the whole 174 community, similar to those related with ocean sprawl. 175

Biodiversity and community structure were determined at ca. 4 m depth for each
treatment over time, using a combination of *in situ* assessment and the deployment of settlement
plates using scuba diving.

*In situ* assessments were non-destructive and relied on percentage cover estimations of sessile biota within a flexible  $0.1 \text{ m}^2$  quadrat (n = 3 per piling; 2 pilings per treatment) divided into 25 sub-quadrats. This was done by adding all of the sub-quadrats; within each an abundance

score from 1 to 4 was assigned to each taxon (Dethier et al. 1993). These samplings were 182 183 conducted prior to- (procedural control at T0) as well as 3 and 10 months post-removals. As commonly considered for these types of communities (especially in the region), 3 and 10 month 184 periods are appropriate to address, respectively, early and later stages of succession or recovery 185 post- disturbances (Dafforn et al. 2009; Canning-Clode et al. 2011; Leclerc et al. 2020b). We 186 expected colonization of pilings to be the result of both growth of remnant biota (e.g., colonial 187 188 ascidians, hydrozoans, and bryozoans) and larval settlements from local and external sources. This sources include adjacent pilings of the dock (in either treatment or untouched, Fig. 1A). 189 Considering the diversity of life history strategies in these fouling communities, the relative 190 191 likelihood of larval dispersal (approximated by pelagic larval duration, Shanks 2009)) across adjacent pilings would vary across major groups and be high in barnacles (e.g. 13-23 days 192 measured at 20°C in Austromegabalanus nigrescens (Egan and Anderson 1987; Pappalardo et 193 194 al. 2016)) and molluscs (e.g. 15-16 days in Crepipatella fecunda (Chaparro et al. 2005)), intermediate for tunicates (e.g. 12-24 h for Pyura chilensis (Cea 1970 in ; Segovia et al. 2020)) 195 196 and low in hydrozoans and bryozoans (minutes to hours in shallow species (Shanks 2009)).

When the removal was completed, a series of plates  $(150 \times 150 \text{ mm})$  were also deployed 197 vertically, using large flexible panels ( $600 \times 600$  mm) as main frames, tied up to each piling 198 (i.e. two panels per treatment) using ropes (Figure S1). Prior to deployment, the surface below 199 each panel was also cleared using a crowbar. The panels were made of plastic fence  $(25 \times 25)$ 200 mm mesh) – on which a total of 8 plates were attached with cable ties – and secured with PVC 201 tubes and garden hoses (25 mm in diameter) to maintain some distance between the plates and 202 the piling surface (Figure S1). For these reasons, we expected the colonization of plates by 203 sessile biota to be made possible mainly by larval settlements (from local and external pools), 204 rather than vegetative growth of remnant biota (or 'residuals', sensu Turner 2010) from the 205 adjacent piling surface. Plates were made of black fluted polypropylene, mimicking most of the 206

coated dock or ship surfaces within the region. Although substrate type indeed affects early stages of colonization (e.g. Smith et al. 2014; Vaz-Pinto et al. 2014; Pinochet et al. 2020) and exacerbates differences between pilings and plates, we were only interested in differences and similarities across removal treatments for each substrate, separately. After 3 and 10 months, 4 plates at random per piling (i.e. 8 per treatment) were retrieved underwater, placed in polypropylene rubble bags (< 0.5 mm mesh), and then stored in a tank filled with sea water until processing in the laboratory (located along the SW shore of San Vicente Bay).</p>

In the laboratory, plates were treated as detailed in Leclerc et al. (2020b). Briefly, sessile 214 taxa (mostly fauna, see Results) were identified under a dissecting microscope and their 215 216 abundances were assessed using cover. Within a  $120 \times 120$  mm working area (avoiding plate edges), species cover was estimated under 100 random intersection points. Any species only 217 identified out of these intersection points was given a cover of 0.5%. Using the same procedure, 218 219 cover was also determined for bare surface, grazing marks and organism remains (e.g. empty tubes or barnacle plates). On either plates or non-destructive quadrats, species layering was 220 221 taken into account; therefore, the total cover frequently exceeded 100%.

Following a framework of criteria analogous to Chapman and Carlton (1991), the identified specimens were categorized as 'native', 'non-indigenous', 'cryptogenic' or 'unassigned'. The cryptogenic species, from uncertain origin (sensu Carlton, 1996), found in this study displayed a cosmopolitan distribution, most likely resulting from historical anthropogenic dispersal (Darling and Carlton 2018) and most likely non-indigenous to the study area (Carlton and Hodder 1995; Leclerc et al. 2018; 2020b). In contrast, the unassigned category was reserved for specimens for which the taxonomic resolution and/or available information was insufficient.

229

230 Statistical analyses

For both in situ and plate surveys, univariate patterns in species richness and abundance 231 232 were examined with a three-way design using PERMANOVAs with 4999 permutations. Factors included 'treatment' (fixed, 3 levels: control, 50% removal and 100% removal), 'time' 233 (fixed) and 'block' (random). The number of levels of 'time' differed between in situ and plate 234 surveys. By using standardized plates in the latter, time before-removal (T0) had no variation 235 associated with any response variable. By including the blocking term in the model, part of the 236 237 total variance was attributed to differences between our two working areas within the study site, thereby reducing the residual unexplained variation (Quinn and Keough 2002). In order to 238 further improve the statistical power of the model, non-significant interactions at  $\alpha = 0.25$  were 239 240 excluded and the model was run again (Underwood 1997). Univariate analyses were based on Euclidean distance matrices (analogous to the ANOVA) generated from either raw or 241 transformed data. The homogeneity in univariate dispersion was checked for the main factor of 242 243 interest, that is 'treatment', using PERMDISP (Anderson et al. 2008). When no transformation (square-root, fourth root, log, arcsine) allowed homoscedasticity to be achieved in univariate 244 245 data (herein the case of biomass), analyses were run on untransformed data (Underwood 1997). PERMANOVAs were followed by pairwise comparisons and P-values were estimated using a 246 Monte Carlo procedure. For the plate survey, univariate analyses were performed either on all 247 taxa combined (including unassigned taxa), native species or cryptogenic and non-indigenous 248 species combined. 249

Multivariate patterns in community structure were analyzed using the same PERMANOVA designs described above. All multivariate analyses were based on Bray-Curtis similarity matrices generated from fourth root-transformed data, in order to downplay the importance of the most abundant species and homogenize multivariate dispersion (which was checked using PERMDISP). PERMANOVA results were supported by ordination (Anderson et al. 2008; Anderson et al. 2017) using principal coordinate (PCO) analyses based on the same similarity matrices. In addition, the main taxa or abiotic variables (e.g., bare surface, grazing marks, dead
biota) explaining differences among treatments were identified according to their correlation to
PCO axes. From this first ordination, distances among centroids for each combination of
treatment × time were determined in a second stage and their temporal trajectories were
visualized using non-metric multidimensional scaling (nMDS). All analyses were performed
using PRIMER 7 (Anderson et al. 2008).

262

#### 263 **Results**

A total of 38 taxa, including 6 non-indigenous and 9 cryptogenic taxa were identified across *in situ* and plate surveys over time (cf. Table S1 for complete list and authorities). Prior to the experiment, neither the percentage of cover of fouling habitat-builders on pilings (average cover of live individuals:  $75\% \pm 14$ , SD; Fig. 1C) nor community structure varied across the pilings (pairwise comparisons in Table 1, Fig. 2). As expected, the experimental removal of habitat-builders (*Austromegabalanus psicattus* and *Pyura chilensis*) on pier pilings had substantial effects (Fig. 1C, Table 1), which are detailed below.

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272 Effect of the removal on pier piling communities.

The effect of removal (i.e. severity of disturbance, (Turner 2010)) on these habitatbuilders was detectable (and significant, Table 1) after 3 months, wherein habitat-builders covered ca. 30 % and < 5 % of the piling surface in the 50% and 100% removal treatments, respectively (Fig. 2D). Although the actual values must be compared to the procedure control prior to the treatments (average cover of 75%), we note a discrepancy observed after 3 months between expected (e.g. 37.5% in the 50% removal treatment) and observed (30%) covers, which may be due to post-manipulation die-off following neighbor removals. Considering Turner's framework (2010), a given energy of a disturbance event per area (i.e. the disturbance intensity)
can indeed conceptually induce a larger effect (i.e. its severity) than expected.

Along with decreasing cover (and contribution) of habitat-builders, there was an 282 283 increasing contribution of various animals to the community structure (cf. first axis of PCO, Fig. 2A), including native (Balanus laevis), cryptogenic (Alcvoniodioides mytilii) and non-284 indigenous taxa (Bougainvillia muscus, Bugulina flabelatta), as well as various unassigned 285 286 algae, including turf (Antithamnionella sp. and Ectocarpales) and foliose forms Ulva sp. Most of these taxa were already present on pilings prior to treatments. Nonetheless, after 10 months, 287 the contribution of these taxa decreased as habitat-builders recuperated their cover to 30 and 288 289 50% of the piling surface in the complete and partial removal treatments, respectively. These values were statistically similar across manipulated assemblages, though still lower than 290 controls (pairwise comparisons in Table 1). After 10 months (discriminated from 3 month-291 samples along axis 2), the community structure converged among treatments (see both PCO 292 and nMDS in Fig. 2A), although significant differences were still present between the complete 293 294 removal treatment and the other two (Table 1, Fig. 2A). Abundant hydroid colonies of Halecium sp. (unassigned status) were found in the complete removal treatments, as illustrated along the 295 second axis of the PCO. 296

297 Effect of the removal on community development on plates

Major shifts in community structure were observed between 3 and 10 months postdeployment of the plates (ca. 52% explained variation according to PCO, Fig. 2B, significant time effect in Table 2), but these shifts were not influenced by the experimental manipulations (removal) of the fouling assemblages on pilings, according to the PERMANOVA (i.e. no interaction 'time × treatment', Table 2). Independent of age, the community structure differed statistically across all treatments (Table 2), with an increase in the contribution of one native ascidian (*P. chilensis*) and two cryptogenic taxa (the bryozoan *Amathia* cf. *gracilis* and the hydroid *Plumularia* cf. *setacea*) to community structure when the partial removal was
compared to control treatments (Fig. 2B). This was further pronounced in the complete removal
treatment.

308 In the complete removal treatment, biomass increased significantly (about twice as heavy) compared to other treatments after 10 months (Table 2, Fig. 3A). Interestingly, neither 309 the cover nor the richness of native taxa differed across treatments at any time post-deployment 310 311 (Table 3, Fig. 3B-E). This result contrasts with cryptogenic and non-indigenous taxa, which displayed an increasing mean richness with removal severity, although cover remained 312 unaffected (Table 3). This pattern of increased colonization was particularly strong after 3 313 314 months, when the average richness of these taxa was greater than after 10 months (Table 3, Fig. 3F, G) In total, 1, 2 and 4 non-indigenous and 5, 7 and 8 cryptogenic taxa were recorded in 315 control, partial and complete removal treatments, respectively (Fig. 3B). 316

317

#### 318 **Discussion**

The removal of the dominant habitat-builders affected community assembly in the 319 recolonization of established assemblages and the colonization of novel substrata on pier 320 321 pilings. As hypothesized (H1), the patches made available by the substrate clearing offered colonization opportunities for novel assemblages, including cryptogenic and non-indigenous 322 species (NIS) otherwise less abundant, rare or absent locally (i.e. at the scale of the focal piling), 323 with contrasting contributions to community structure across treatments. Interestingly, even 324 while initially offered the same free surface for colonization, the richness (but not the 325 abundance) of cryptogenics and NIS on settlement plates increased significantly with the 326 severity of the disturbance experienced by the surrounding assemblages. As hypothesized (H2), 327 the contrasting response on pilings and plates could indicate indirect effects decoupled from 328

spatial competition at the patch scale. As for natives, neither their richness nor their abundance 329 330 varied across treatments in the colonization of plates. Although none of the cryptogenics and NIS are new in this highly frequented, principal regional port, contrasting responses to 331 treatments among natives versus cryptogenics and NIS could reflect differences in life history 332 traits and distinct legacies of these two species pools. Finally, regarding H3, although habitat-333 builders did not recover their initial (control) occupied surface in any of the treatments, 334 335 community structure on pilings generally converged across treatments after 10 months; differences were only detected between complete removal and other treatments. As these native 336 dominants bounced back rapidly, cryptogenic and non-indigenous taxa decreased in their 337 338 contribution on pilings.

#### *1. Direct effect of physical disturbance: decreasing spatial competition*

One way in which physical disturbance directly influences community assembly is by 340 providing open substratum for remote and local colonizers (e.g. Levin and Paine 1974; Connell 341 342 and Slatyer 1977; Sousa 1979; Shea and Chesson 2002). When propagule and colonization 343 pressures are high in the long-term, local disturbances can increase the probability of the establishment of more abundant and diverse NIS (Davis et al. 2000; Clark and Johnston 2009; 344 Blackburn et al. 2020), including their possible impacts (being intimatelly linked with their 345 proliferation, Geraldi et al. 2019). Here, the piling-scale removal of the dominant native space-346 occupiers (Austromegabalanus psicattus and Pyura chilensis) - both of which are harvested 347 along the SE Pacific – has generally led to a greater contribution of diverse native, cryptogenic 348 and non-indigenous species (as well as unassigned turf algae) to the community structure (as 349 hypothesized, H1), but only at an early post-disturbance stage (i.e. hereafter 3 months). 350

The observed early colonizers are either stress resistant (to the physical disturbance), opportunistic or both. Withstanding stress in this context implies that live remnant colonies (or 'residuals', sensu Turner 2010) are capable of rapid vegetative growth, permitting the

colonization of vacant patches (Sousa 1979; Valdivia et al. 2005; Hiebert et al. 2019). In spite 354 355 of the limited larval dispersal of shallow water hydroids (over a few meters in a few minutes, Shanks 2009), such spreading capabilities are notably found in vines and runners (through 356 357 stolonal expansion, Greene and Schoener 1982; Schuchert 2007), such as Bougainvillia muscus and *Clytia linearis* (see disturbed pilings in Fig. 2). Opportunism can involve a greater diversity 358 of traits (e.g., reproductive output), but it generally implies a low rank in the competitive 359 hierarchy (i.e., along an inter-specific competition-colonization trade-off, Greene and Schoener 360 1982; Chesson 2000; Valdivia et al. 2005; Wernberg et al. 2020). For instance, although 361 barnacle larvae may disperse over kilometers in several weeks (Shanks 2009; Pappalardo et al. 362 363 2016), small species such as Balanus laevis (see Fig. 2) are unable to secure the primary substratum when competing with larger ones, such as A. psicattus (Leclerc et al. 2020a, this 364 study). The prevalence of cryptogenic and non-indigenous species fitting into these categories 365 366 may illustrate the legacy of the disturbance regimes typical of artificial habitats. These species, most of which could be considered neocosmopolitan (sensu Darling and Carlton 2018), have 367 been frequently and rapidly observed on settlement plates deployed at various ports in the 368 region (Leclerc et al. 2018; Leclerc et al. 2020b). Over their natural history, they have likely 369 colonized diverse artificial habitats (e.g. ship hulls, debris, port infrastructures), selecting for 370 traits conducive of the fouling network that these habitats compose. Although this hypothesis 371 is beyond the scope of this study, we note that this phenomenon is akin to trait biases observed 372 within terrestrial plants and the legacy of ruderal habitats (Catford et al. 2012; Hufbauer et al. 373 2012; Pearson et al. 2018b). The expansion of novel artificial structures (ocean sprawl), almost 374 independent of further co-varying stresses (Todd et al. 2019) is likely to worsen future invasions 375 across novel or intensifying pathways (Dafforn et al. 2009; Aguilera 2018; Geraldi et al. 2019; 376 Sardain et al. 2019). 377

While the contrasting contribution of cryptogenic and non-indigenous species to community structure across treatments on pilings was probably driven by varying amounts of open surface, the same hypothesis hardly stands for the patterns observed on settlement plates, which had standard open surfaces (see H2). Beyond the competition for free surface, removal of the dominant habitat-builders of our system probably influenced community assembly by additional and more indirect mechanisms discussed in the following sections.

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#### 2. Indirect effect of physical disturbance: decreasing competition for other resources

Although open space is generally considered as the main limiting resource for sessile 385 invertebrates in fouling assemblages (Osman and Whitlatch 1995a; Stachowicz et al. 2002), 386 there is mounting evidence that competition for trophic resources can also influence community 387 assembly in such systems, even at local scales (Buss 1979; Svensson and Marshall 2015). Using 388 in situ experiments, Svensson and Marshall (2015) demonstrated that the nature, diversity and 389 abundance of planktonic resources can influence the richness and structure of sessile 390 391 assemblages, at least at an early stage (1 month). In our experiment, we cannot rule out that the 392 varying amount of sessile fauna left in place across treatments led to contrasting trophic competition between these organisms and the new recruits, even those which colonized 393 settlement plates with standardized free surfaces. Although certainly not exclusive (e.g., 394 predation on larvae by neighbour adults, Young and Gotelli 1988; see also next section), 395 competition for food resources could actually constitute a solid hypothesis explaining the 396 significant difference across treatments observed in biomass on fully covered plates after 10 397 months (Ctrl = 50% removal < 100% removal, Fig. 3A). Escaping such competition, recruiting 398 organisms may indeed have been promoted for growth between sampling periods in the full 399 400 removal treatment, as compared to their counterparts in control and partial removal treatments. The ability of species to secure phytoplankton resources could thus mediate competition for 401 space (Buss 1979). Considering such a possibility, assessing individual growth and post-402

recruitment survival on a series of focal species could be valuable in future research to
disentangle diverse interacting processes at a micro-scale, such as propagule pressure,
competition for food and/or space, and predation.

406 *3. Indirect effect of physical disturbance: partially releasing incomers from predation* 

Habitat-builders often provide shelter to the predators of their own competitors (and 407 predators); therefore, their removal can have side effects. Previous work carried out in three 408 ports of the region indicated that benthic predators had an overall stronger effect on most non-409 indigenous species than on native species at an early stage of development on settlement plates 410 (Leclerc et al. 2020a). Results varied across sites and predator types (separated by size) but, in 411 general, predator exclusion led to an increase in cryptogenic and non-indigenous richness, 412 categories of which 18% and 36% taxa, respectively, were observed only under predator 413 exclusion. These results mirror the differences herein observed for richness among treatments 414 (Figs. 3B, 3F). Although benthic predators were not quantified, local loss of some of them, due 415 416 to the removal of habitat-builders, is very likely (Hernández et al. 2001; Sepúlveda et al. 2003). 417 This was also supported by our observations of macro-predators (> 2 cm) in situ, seemingly associated with complex biogenic habitats made up by well-developed fouling communities 418 (Fig. 1D, JCL, pers. obs., Leclerc et al. 2018) (see also Coleman and Connell 2001). Such a loss 419 in predator diversity may have contributed to the observed differences in community structure 420 421 across treatments. Along with other studies (Dumont et al. 2011; see also Dias et al. 2020), the experiments mentioned above also demonstrated that an important predation on most of the 422 non-indigenous and cryptogenic ascidians gave a competitive advantage to their native, 423 dominant and less predated counterpart, namely Pyura chilensis, one of our habitat-builders. 424 425 This effect, measured by percentage of cover reduction after 4 months in a settlement panel experiment, was, however, not consistently detected by the exclusion of macro-predators 426 427 (Leclerc et al. 2020a). According to further exclusion treatments, the authors found that micro-

predators (ranging between 2 mm and 2 cm) largely contributed to the biotic resistance 428 429 described above, by controlling post-settlement survival (see also Osman and Whitlatch 1995b; Oricchio et al. 2016). Considering the turn-over, dispersal abilities and rapid colonization of 430 many micro-invertebrates across diverse habitats (dispersing 10s m within a few days to weeks 431 (Waage-Nielsen et al. 2003; Martins et al. 2017), many micro-predators may have rapidly 432 recolonized disturbed pilings from adjacent ones, throughout the recovery of sessile 433 434 assemblages, even at an early stage of biogenic habitat formation (see also Leclerc and Viard 2018). As in other marine systems (e.g. Waage-Nielsen et al. 2003; Freestone et al. 2011), 435 diverse and abundant assemblages of micro-invertebrates may have strong influences on the 436 437 structure and dynamics of associated communities, as well as their responses to anthropogenic influences, which would be worth further considering in future studies. 438

#### 439 4. Implications, perspectives and concluding remarks

Regardless of the precise mechanism explaining differences in biomass on plates across 440 441 treatments, and although we did not follow a strict functional approach, our results most likely 442 illustrate that the distribution of traits conferring high efficiency in exploiting and maintaining resources in the native community may be critical not only for its biotic resistance (Davis et al. 443 2000; Bulleri et al. 2008), but also for its resilience after a physical disturbance (see H3). In our 444 system, the native habitat-builders largely recovered from removal within a few months. 445 Opportunist taxa, such as those characterizing most non-indigenous fouling species established 446 in our region, are indeed poorly efficient in securing resources and inhibiting overgrowth by 447 native dominants (Valdivia et al. 2005; Cifuentes et al. 2010). Although experimental outcomes 448 at an early stage of succession would certainly vary with the timing and regime of the 449 450 disturbance (e.g. Chesson 2000), the convergence in community structure shown here has been observed across multiple succession trials in this very site (Leclerc et al. 2020a) and in nearby 451 regions (Valdivia et al. 2005; Cifuentes et al. 2010). 452

Nonetheless, this rather deterministic hypothesis overlooks the variations in propagule and 453 454 colonization pressures characterizing port areas. The observed outcome at later stages of succession are indeed undermined by an "ecological roulette" (sensu Carlton and Geller 1993), 455 456 related to the diversity of traits in the non-indigenous species pool and niche similarities with natives, in recipient communities (Davis et al. 2000; Catford et al. 2019; Zelnik et al. 2019). In 457 the framework of a competition-colonization trade-off, for instance, disturbed communities 458 459 could be greatly susceptible to profound and perennial changes in native community structure depending on standoff, self-organization, and competitive traits in successful settlers (Sebens 460 1986; Catford et al. 2018; Bertolini et al. 2019; Catford et al. 2019). Strong competitive traits 461 462 are common in widespread sessile marine invaders, such as seagrasses, seaweeds, oysters and mussels, which are known to affect ecosystems through habitat engineering, resource use and 463 nutrient cycling (Paine 1974; Thomsen et al. 2014; Bailey et al. in press). As elsewhere, 464 465 successful introduction and establishment of such highly competitive space-occupiers have been common throughout the SE Pacific (Castilla and Neill 2009; Villaseñor-Parada et al. 466 2018). Many of them are still being transported due to little regulated human activities (e.g. 467 Cardenas et al. 2020), which are showing alarming signs of expansion across the planet 468 (Seebens et al. 2018; Sardain et al. 2019). Altogether, our results reveal, on the one hand, that 469 the dominant native fouling species of the study region display a series of traits and competitive 470 abilities conferring their population with rapid recovery post-disturbance and the potential 471 sustainability of well-managed localized fisheries. On the other hand, the currently unregulated-472 harvesting practices certainly deserve further attention, most notably in a context of marine 473 urbanization and expanding shipping networks. 474

475

476 Acknowledgements. We are particularly grateful to N Valdivia for helpful discussions and
477 critical reading of this manuscript. The experimental manipulation has been made possible

underwater by the logistical support of the Valderrama family at Prosub Ltd (*http://prosub.cl/*) 478 479 and by access to the port thanks to N Cofré. We would like to warmly thank J Martínez and J Cruz for sailing assistance and to M Altamirano, R Reed and O Marin for diving assistance. 480 We would also like to acknowledge D Gordon for help with Exochella identification. JCL 481 further thanks the Centre IDEAL (Investigación Dinámica de Ecosistemas Marinos de Altas 482 Latitudes, Universidad Austral de Chile) for its kind hospitality regarding the writing of the 483 manuscript. JCL and AB were supported by FONDECYT (N°3160172 and N°1170598, 484 respectively). 485

486 **Credit Author Statement.** JCL and FV conceived and designed the study. JCL, AB and FV 487 obtained the initial funding. JCL led the sample collection and processing. JCL analyzed the 488 data and led the writing, with substantial contributions from FV to the first draft. All authors 489 approved the manuscript submission.

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#### 747 Tables

Table 1: Results of PERMANOVA tests for differences in habitat-builders cover (HB, %) and community structure on pier pilings among levels of the main fixed (experimental treatment, time) and random (block) factors, as well as their interactions. Non-significant interaction terms at  $\alpha = 0.25$  were removed (as indicated by "-") and the model was run again. Significant *P*values are in bold. Pairwise tests are summarized, with T0 indicating the moment of removal; T3 and T10 indicate 3 and 10 months post-removal, respectively; and Ctrl, 50%r, 100%r indicate control, partial and complete removals, respectively.

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		HB cover (%)	)	Community structure			
Transformation	-	None		Fourth-roo	Fourth-root		
Dispersion		$F_{8,45} = 2.17 P$	= 0.111	$F_{8.45} = 1.84 P = 0.157$			
Main test							
				Pseudo-			
Source	df	Pseudo-F	Р	F	Р		
Treatment (Co)	2	45.39	< 0.001	1.59	0.203		
Time (Ti)	2	10.64	0.122	7.25	0.066		
Block (Bl)	1	0.16	0.684	2.37	0.038		
Co x Ti	4	12.45	< 0.001	2.84	< 0.001		
Co x Bl	2	-	-	1.96	0.037		
Ti x Bl	2	2.15	0.128	1.62	0.096		
Pairwise							
comparisons:							
T0:		Ctrl = 50%r = 100%r		Ctrl = 50%r = 100%r			
T3:	$Ctrl > 50\%r > 100\%r$ $Ctrl \neq 50\%r \neq 100\%r$						
T10: $Ctrl > 5$			100%r	Ctrl = 50%	$Ctrl = 50\%r \neq 100\%r$		

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- Table 2: Results of PERMANOVA tests for differences in biomass (g) and community structure
  on settlement plates among levels of the main fixed (experimental treatment, time) and random
  (block) factors, as well as their interactions.. Significant *P*-values are in bold. Abbreviations
  are the same as in Table 1.

	]	Biomass (g)	Community structure			
Transformation	]	None		Fourth-root		
Dispersion		$F_{5,45} = 6.11 P =$	<i>F</i> <sub>5,42</sub> = 7.04 <i>P</i> < 0.001			
Main test						
Source	df	Pseudo-F	P	Pseudo-F	Р	
Treatment (Co)	2	4.20	0.024	4.42	< 0.001	
Time (Ti)	1	39.30	< 0.001	13.35	0.011	
Block (Bl)	1	1.81	0.190	2.25	0.041	
Co x Ti	2	3.78	0.029	1.41	0.171	
Ti x Bl	1	-	-	4.08	< 0.001	
Pairwise						
comparisons:						
	r	$\Gamma 3: \operatorname{Ctrl} = 50\% r$	= 100%r	$Ctrl \neq 50\%r \neq 100\%r$		
	r	$\Gamma 10: \operatorname{Ctrl} = 50\%$				

765 Table 3: Results of PERMANOVA tests for differences in richness (S) and cover (%) of native, cryptogenic and non-indigenous species on

settlement plates among levels of the main fixed (experimental treatment, time) and random (block) factors, as well as their interactions.

767 Significant *P*-values are in bold. Abbreviations are the same as in Table 1.

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		Native				NIS + Cryptogenics			
		Richness							
		S		Cover (%)		Richness S		Cover (%)	
Transformation		None		None		None		None	
Dispersion		$F_{5,42} = 1.68 P = 0.203$		$F_{5,42} = 2.9 P = 0.107$		$F_{5,42} = 1.63 P = 0.268$		$F_{5,42} = 0.43 P = 0.937$	
Main test									
Source	df	Pseudo-F	Р	Pseudo-F	Р	Pseudo-F	Р	Pseudo-F	Р
Treatment (Co)	2	1.48	0.400	1.74	0.191	13.03	< 0.001	2.49	0.099
Time (Ti)	2	0.97	0.333	6.27	0.236	18.61	< 0.001	2.93	0.340
Block (Bl)	1	1.61	0.213	0.41	0.516	0.02	0.890	2.37	0.127
Co x Ti	2	0.35	0.747	0.27	0.787	2.01	0.146	2.28	0.116
Co x Bl	2	1.67	0.194	-	-	-	-	-	-
Ti x Pl	1	-	-	5.19	0.029	-	-	7.07	0.013
Co x Ti x Bl	2	2.94	0.073	3.25	0.053	-	-	-	-
Pairwise									
comparisons:					Ctrl < 50%r	< 100%r			

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#### 771 Figures

Figure 1: Experimental design and habitat-builders cover, before and after removal treatment, 772 upon pier pilings at the study location. Within each block (A), experimental treatments 773 774 consisted of the removal (partial ca. 50%, 100% vs. control, B) of native fouling habitat-builders (Austromegabalanus psicattus, Pyura chilensis), commonly inhabited by various macro- (e.g. 775 Romaleon setosum, Homalaspis plana) and micro-predators (e.g. Hypsoblennius sordidus 776 777 inhabiting empty barnacle shells, and Palaemon sp. crawling among epibiotia) that are highlighted here by the white arrows in the pictures (C, photos are courtesy of Mauricio 778 Altamirano). For each sampling occasion (before, after removal), letters indicate significant 779 780 differences in the cover of habitat-builders among treatments (D, n = 6). Around the median (horizontal line), the box plots are bounded by the quartiles, the 95% confidence intervals 781 (whiskers) and the outsiders. 782

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Figure 2: Community structure (principal coordinate analysis, PCO) across treatments on 784 settlement plates (A) and on surrounding assemblages (piling, B) before and after experimental 785 removal of habitat-builders (3 and 10 months). Vector plots of variables correlated with the 786 PCO axes are indicated, with r > 0.4 and 0.6 for pier piling and plate assemblages, respectively. 787 Taxa are in bold with the color code indicating the status: unassigned (black), native (blue), 788 cryptogenic (orange) and non-indigenous (red) species. Temporal trajectories of the centroids 789 790 (determined from PCO) axes for each combination of treatment × time are ordinated using nonmetric multidimensional scaling (right insets). 791

Figure 3: Biomass (g), richness (S) and cover (%) on settlement plates after 3 and 10 months
following experimental removal. The total biomass is given (A), while richness (B, D, F) and

cover (C, E, G) values are given for all, native, NIS and cryptogenic taxa. Richness values are either given as cumulated over (bar charts in B, C) or averaged (box plots in D, F) across replicate plates per combination of treatment  $\times$  time (n = 8 plates). Around the median (horizontal line), the box plots are bounded by the quartiles, the 95% confidence intervals (whiskers) and the outsiders.

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Figure 1



Figure 2



