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The effects of an invasive habitat modifier on the biotic interactions between two native herbivorous species and benthic habitat in a subtidal rocky reef ecosystem

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Abstract Range expanding species can have major impacts on marine ecosystems but experimental field based studies are often lacking. The urchin *Centrostephanus rodgersii* has recently undergone a southerly range expansion to the east coast of Tasmania, Australia. We manipulated densities of *C. rodgersii* and algal regrowth in urchin barrens habitat to test effects of the urchin on biotic interactions between two native herbivores, black-lip abalone (*Haliotis rubra*) and another urchin (*Heliocidaris erythrogramma*), and their benthic habitat. After 13 months, removals of only *C. rodgersii* resulted in overgrowth of barrens habitat by algae and sessile invertebrates. Densities of

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Institute for Marine and Antarctic Studies, University of Tasmania, Private Bag 129, Hobart, TAS 7001, Australia abalone increased (+92 %) only in patches from which C. rodgersii was removed and algal regrowth allowed. In contrast, densities of H. erythrogramma increased in all treatments (+45, +28, +25 %) in which C. rodgersii was removed, irrespective of the algal regrowth manipulations. These results suggest that C. rodgersii has a negative influence on the densities of abalone through competition for food and on densities of H. erythrogramma through competition for preferred habitat. Densities of abalone (+65 %) but not *H. erythrogramma* (+25 %), were lower in the patches from which C. rodgersii and canopy algae regrowth were removed relative to patches from which only C. rodgersii was removed (+92 and +28 %, respectively). These results suggest that C. rodgersii overgrazing of canopy-algae results in loss of structural complexity which could increase abalone susceptibility to predation, cause abalone to seek shelter in cryptic microhabitats and/or prevent their return to patches where canopy algae are absent. The ongoing spread of C. rodgersii and expansion of barrens habitat in eastern Tasmania will continue to negatively affect populations of these two native herbivores and their associated fisheries at a range of spatial scales. This example shows that habitat modifying species which become highly invasive can have disproportionate negative impacts on the structure and dynamics of the recipient community.

Keywords Range expansion · Habitat modifier · Biotic interactions · Urchins · Abalone

Introduction

Global climate change is leading to the poleward range expansion of many marine species (Parmesan and Yohe 2003; Hickling et al. 2006; Poloczanska et al. 2008). Range expanding species that create or modify habitat are predicted to have profound effects on ecosystem structure and function (Grosholz 2002; Harley et al. 2006). These ecosystems engineers can alter habitat complexity, environmental chemistry and other physical variables with major effects on the abundances, diversity and context-dependent interactions of native species (Sorte et al. 2010; Walther 2010). Despite their importance, field-based experimental studies on the impacts of many range expanding habitat modifiers on marine ecosystems are lacking (but see Bertness 1984; O'Connor and Crowe 2005; Hollebone and Hay 2008; Ling 2008; Firth et al. 2009).

The southeast coast of Australia has experienced an increase in the abundances of many warmer water species (Stuart-Smith et al. 2009; Johnson et al. 2011) as a result of greater poleward penetration of the East Australia Current (Ridgway 2007), which has been linked in part to Antarctic ozone depletion (Cai et al. 2005; Cai 2006). One of the most conspicuous species that has undergone southerly range expansion from New South Wales to the east coast of Tasmania as a result of greater penetration of the East Australian Current, is the long spined urchin Centrostephanus rodgersii (Johnson et al. 2005, 2011; Ling et al. 2009; Stuart-Smith et al. 2009). This urchin is a habitat modifier, well known for its ability to overgraze filamentous and foliose algae and sessile invertebrates, effecting a catastrophic shift to barrens habitat dominated by the urchin and characterised by bare rock (Johnson et al. 2005; Ling 2008) or, in its native habitat, encrusting red algae (Fletcher 1987).

Centrostephanus rodgersii was first detected off the north east coast of Tasmania in 1978 (Edgar 1997). Since then, the abundances and range of this species have increased, and extensive barrens habitat (>100 m diameter) has formed at some locations on the north east coast of Tasmania as a result of urchin overgrazing seaweeds (Johnson et al. 2005; Ling 2008). While extensive barrens are not yet a widespread feature of the east coast of Tasmania, development of incipient barrens patches (\leq 10 m diameter) in otherwise intact algal beds are characteristic of much of the coastline (Johnson et al. 2005; Ling 2008). The continued spread and increase in abundances of *C. rodgersii* poses a major threat to the structure and function of macroalgal beds (Edgar et al. 2004; Ling 2008) and the important commercial fisheries species they support (Johnson et al. 2005, 2011; Strain and Johnson 2009).

Urchins and abalone consume predominately understorey filamentous and foliose algae and share similar habitat and predators (Shepherd 1973; Tegner and Levin 1982; Day and Branch 2000; Naylor and Gerring 2001). Surveys along the south east coast of Australia have demonstrated a negative relationship between the abundances of C. rodgersii, and that of H. rubra and H. erythrogramma at a broad range of spatial scales (Johnson et al. 2005; Andrew and Underwood 1992). Previous studies have suggested that C. rodgersii has a negative impact on the abundances of H. rubra and H. erythrogramma through competition for food (Shepherd 1973; Andrew et al. 1998; Strain and Johnson 2009) and/or preferred habitat (Andrew and Underwood 1992) in intact algal beds. Overgrazing of canopy-algae by C. rodgersii could also have a negative effect on the abundances of H. rubra and H. erythrogramma through loss of structural complexity (Andrew 1993; Andrew et al. 1998; Edgar et al. 2004). However, experimental studies designed to separate these hypotheses and test the effects of C. rodgersii on the abundances of both H. rubra and H. erythrogramma in barrens are lacking. In this study we manipulated densities of C. rodgersii and the algal regrowth to determine the impacts of this urchin on the densities of H. rubra and H. erythrogramma through competition for food, preferred habitat and/or loss of canopyforming algae.

Materials and methods

Study area and experimental manipulations

This study was conducted at two randomly selected sites at the Lanterns $(43^{\circ}8'20''S, 148^{\circ}0'21''S)$ and $43^{\circ}8'19''S, 148^{\circ}0'21''S)$ on the east coast of Tasmania, Australia, between August 2005 and September 2006. Both sites have steeply sloping rocky substratum to a depth of >30 m and are moderately exposed.

At each site, flagging tape was used to mark the perimeter of 12 discrete Centrostephanus rodgersii barrens patches (mean width = $5.02 \text{ m}, \pm \text{SE} =$ 0.12 m, mean length = 5.14 m, \pm SE = 0.10 m), each supporting 18–22 resident C. rodgersii (mean = 19.833 individuals, $\pm SE = 1.726$ individuals) and of three other patches within the intact algal bed of similar size to the barrens patches but supporting seaweeds and no C. rodgersii. All barrens patches were randomly assigned to the following treatments: T1 = unmanipulated C. rodgersii barrens patches; T2 = removal ofC. rodgersii and all regrowth from patches; T3 = removal of C. rodgersii and regrowth of canopy-algae species from patches; and T4 = removal of C. rodgersii only from patches. The intact algal patches without C. rodgersii represented a control treatment (T5). There were n = 3 replicate patches of each

variables were not monitored within 0.1 m of the tape. At the beginning of the experiment, divers removed all *C. rodgersii* from T2, T3 and T4 patches using knives. Throughout the experiment, divers revisited the sites every 2 months to maintain the manipulations. At each visit, all reinvading *C. rodgersii* were removed (T2, T3 and T4), all algae and sessile invertebrate regrowth removed (T2) by scrubbing the substratum with a copper wire brush, and all regrowth of canopy-algae species (\geq 300 mm total length) removed by hand (T2 and T3), as appropriate for the treatment. These manipulations were undertaken for 13 months to allow at least one cycle of algal and sessile invertebrate regrowth (Ling 2008).

treatment. To avoid possible edge effects, the response

Benthic community

At both sites, the benthic community was assessed in a two-stage process, using a modification of the methods of Valentine and Johnson (2003). For each patch, the number of stipes and percentage cover of canopy-algae (\geq 300 mm in height) were assessed by eye in four randomly positioned 0.5 × 0.5 m quadrats. The fronds of these plants were then moved aside and the understorey community was assessed by photography using a digital Canon Powershot camera A95 with 2× Nikonos SB-102 strobes. A grid of 100 equidistant points was overlaid over the photographs and the taxa under each point identified to estimate community structure in terms of percentage cover. Understorey algae and sessile invertebrates that could not be

identified to species level were allocated to complexes or higher taxonomic groups (e.g. *Zonaria/Lobophora*, sponge, ascidian etc.). All visual assessments were conducted by the same diver.

Densities of urchins and abalone

At both sites, divers counted the total number of urchins (*C. rodgersii* and *Heliocidaris erythrogramma*) and abalone (*Haliotis rubra*) see below for sampling details. These counts were converted to total densities (m^{-2}) of *C. rodgersii*, *H. rubra* and *H. erythrogramma*. Patch area was calculated using the formula for an ellipsoid:

Patch area = $(\pi \times (\text{Length} \times \text{Width}))$.

Due to time constraints, the schedule of monitoring responses of the benthic community to the manipulations differed between the two sites however manipulations at both sites followed an identical schedule. At the first site, the benthic community was assessed immediately prior to the manipulations, 1 month later and then every 2 months. At the second site, the benthic community was assessed immediately prior to the manipulations and then after 7 and 13 months.

Analyses

The effects of the different treatments on benthic community structure after 13 months described as functional groups (density of stipes and percentage cover of canopy-algae, and percentage cover of bare rock, encrusting red algae, filamentous algae, foliose understorey algae, and sessile invertebrates) were analysed using univariate 3-way nested ANOVAs. In each case the model included the main effects of treatment (fixed, 5 levels = T1–T5) and site (random, 2 levels = A and B), and patches (random, 3 levels = 3 replicates) nested within the treatment and site interaction.

The overall effect of the treatments on the benthic community after 13 months was analysed using 3-way PERMANOVA (as per model described above). To depict community structure, we used non-metric multi-dimensional scaling (nMDS) plots. The PER-MANOVA and nMDS analyses were based on Bray-Curtis similarity matrices derived from percentage cover data after a square root transformation to reduce the influence of dominant species. All multivariate tests were undertaken using the statistical software Primer 6.0 with the PERMANOVA extension (Clarke and Warwick 2001; Anderson et al. 2008).

The effects of the different treatments on the densities of *H. rubra* and *H. erythrogramma* after 13 months were analysed using 2-way ANOVAs. The model included the main effects of treatment (fixed, 5 levels = T1-T5) and site (random, 2 levels = A and B) and their interaction.

The responses of the benthic community *H. rubra* and *H. erythrogramma* manipulations are depicted graphically for each assessment. However, the effects of the manipulations on the benthic community and the densities of *H. rubra* and *H. erythrogramma* were analysed at an a priori time of interest, i.e. after 13 months, to allow sufficient time for regrowth of both understory and overstorey algae in the barrens patches (see below Table 1).

Prior to all univariate tests, transformations to stabilize variances were determined from the relationship between group standard deviations and means. In all figures raw variables are depicted. Following the main analyses, one or two-tailed t-tests were made as planned comparisons at 13 months to assess the separate effects of competition for food, competition for preferred habitat, and loss of canopy-forming algae to facilitate interpreting overall effects of treatments on densities of *H. rubra* and *H. erythrogramma* (Table 1). For all tests α was adjusted using the procedure suggested by Todd and Keough (1994). All univariate tests and all univariate and multivariate graphical representations were undertaken using the statistical software R (www.R-project.org).

Results

A summary of trends through time for each treatment is given in Table 2. Detailed results are outlined below.

Benthic assemblage structure

At the initial assessment the benthic assemblage in all *Centrostephanus rodgersii* barrens patches was similar and distinctly different to that of the patches within the intact algal bed (Fig. 1). After 13 months, there were significant differences between the treatments and sites (treatment × site: $F_{4, 119} = 2.040$, P = 0.001), particularly in the treatments from which *C. rodgersii* and all regrowth were removed (T2). The planned comparisons showed clear separation in MDS space between the benthic assemblage structure in the unmanipulated *C. rodgersii* barrens patches (T1) and the barrens patches from which *C. rodgersii* and canopy-algae regrowth were removed (T3) (F =

 $10.518, P = 0.006, \alpha \text{ adjusted} = 0.0125$, and patches from which only C. rodgersii was removed (T4) $(F = 5.006, P = 0.010, \alpha \text{ adjusted} = 0.0125)$. There was also clear separation between the intact algal patches (T5) and the incipient barrens patches from which C. rodgersii and canopy-algae regrowth (T3) $(F = 13.848, P = 0.001, \alpha \text{ adjusted} = 0.0125)$ and from which C. rodgersii and all regrowth were removed (T2) (F = 3.557, P = 0.010, α adjusted = 0.0125) (Fig. 1). The barrens patches from which C. rodgersii and all regrowth (T2) was removed and the unmanipulated C. rodgersii barrens patches (T1) were also separated in MDS space, particularly at Site B, however these differences were not significant after adjusting for multiple testing (F = 3.557, P = 0.010, α adjusted = 0.0125) (Fig. 1). In contrast, after 13 months there were no detectable differences in community structure at a functional group level between incipient barrens patches from which only C. rodgersii was removed (T4) and the intact algal patches (T5) $(F = 1.000, P > 0.05, \alpha \text{ adjusted} = 0.0125)$ (Fig. 1).

Similarly, after 13 months of removing only C. rodgersii from barrens patches there were no detectable differences in the cover of foliose understorey algae and sessile invertebrates in the treatment patches (T4) relative to the control patches with no C. rodgersii in the intact algal bed (T5) (Table 3; Figs. 2, 3). There was however still a significantly higher density of stipes of canopy-algae but lower cover of canopy-algae, encrusting red algae and filamentous algae in the T4 patches compared with the intact algal patches (T5), indicating that the development of algae in these patches had not yet achieved the full characteristics of the seaweed community surrounding them (Table 3; Figs. 2, 3). Removals of C. rodgersii and all regrowth to simulate barrens patches (T2) resulted in a higher cover of filamentous algae compared with the unmanipulated C. rodgersii barrens patches (T1) (Table 3; Figs. 2, 3), reflecting that 2 monthly **Table 1** Details of planned comparisons, of the effects of manipulations on the benthic assemblage structure habitat and densities of *C. rodgersii, H. rubra* and *H. erythrogramma* in

treatment patches (mean patch size = 30.654 m^2 , $\pm SE = 0.630 \text{ m}^2$) after 13 months

Comparisons	Hypotheses	Expected response
Benthic communi	ty	
T2 versus T1	Removal of all regrowth will simulate <i>C. rodgersii</i> barrens	Similar benthic assemblage, density of stipes and cover of canopy-algae, cover of understorey
T2 versus T5	Removal of all regrowth will simulate <i>C. rodgersii</i> barrens	Significantly different benthic assemblage
T2 < T5	Removal of all regrowth will simulate <i>C. rodgersii</i> barrens	Significantly less density of stipes and cover of canopy-algae, cover of understorey
T3 versus T1	Removal of canopy algae regrowth will allow understorey regrowth	Significantly different benthic assemblage
T3 versus T5	Removals of canopy algae regrowth will allow understorey regrowth	Significantly different benthic assemblage
T3 versus T1	Removals of canopy algae regrowth will simulate the loss of structure on <i>C. rodgersii</i> barrens	No detectable differences density of stipes and cover of canopy-algae
T3 < T5	Removals of canopy algae regrowth will the simulate loss of structure on <i>C. rodgersii</i> barrens	Significantly less density of stipes and cover of canopy-algae
T3 > T1	Removals of canopy algae regrowth will allow understorey regrowth	Significantly higher cover of understorey
T3 versus T1	Removals of canopy algae regrowth will allow understorey regrowth	No detectable differences cover of understorey
T4 > T1	Removals of <i>C. rodgersii</i> will result in overgrowth of barrens	Significant differences benthic assemblage and Significantly higher density of stipes and cover of canopy-algae, cover of understorey
T4 versus T5	Removals of <i>C. rodgersii</i> will result in overgrowth of barrens	No detectable differences benthic assemblage, density of stipes and cover of canopy-algae, cover of understorey
H. rubra and H. e	erythrogramma density	
T3 > T1	C. rodgersii outcompetes H. rubra and H. erythrogramma for food and preferred habitat	Higher densities of <i>H. rubra</i> and <i>H. erythrogramma</i>
T2 > T1	C. rodgersii outcompetes H. rubra and H. erythrogramma for preferred habitat	Higher densities of <i>H. rubra</i> and <i>H. erythrogramma</i>
T3 > T2	<i>C. rodgerii</i> outcompetes <i>H. rubra</i> and <i>H. erythrogramma</i> for food rather than for preferred habitat	Higher densities of <i>H. rubra</i> and <i>H. erythrogramma</i>
T3 < T4	<i>H. rubra</i> and <i>H. erythrogramma</i> depend on structural complexity for protection from predators	Higher densities of <i>H. rubra</i> and <i>H. erythrogramma</i>

Treatments are, unmanipulated *C. rodgersii* barrens (T1), removal of *C. rodgersii* and all regrowth from patches (T2), removal of *C. rodgersii* and canopy-algae regrowth from patches (T3), removal of *C. rodgersii* only (T4), control no *C. rodgersii* in intact algal patches (T5)

visitations and associated manipulations were insufficient to prevent some development of filamentous algae in this treatment. However, after 13 months there were no detectable differences in the density of stipes and cover of canopy-algae, the cover of encrusting red algae, foliose understorey algae and sessile invertebrates between the T2 patches and unmanipulated incipient barrens patches (T1). Not surprisingly, there was a significantly lower density and cover of canopy-algae and cover of encrusting red algae, filamentous algae, understorey foliose algae and sessile invertebrates in the treatment patches (T2) relative to the intact algal patches (T5) (Table 3; Figs. 2, 3).

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Response variable	Treatments						
	T1	T2	Т3	T4	T5		
Density of C. rodgerst	ii						
Aug 05	0.800 (0.027)	0.910 (0.063)	0.900 (0.068)	0.680 (0)	0.080 (0.020)		
Sep 06	0.980 (0.053)	0 (0)	0 (0)	0.020 (0.010)	0 (0)		
% Change	+20	-100	-100	-94	-100		
Density of H. rubra							
Aug 05	0.120 (0.046)	0.090 (0.076)	0.020 (0.028)	0.220 (0.008)	1.220 (0.051)		
Sep 06	0.120 (0.044)	0.100 (0.044)	0.490 (0.036)	1.040 (0.027)	1.160 (0.047)		
% Change	0	+10	+92	+65	-2.520		
Density of H. erythrog	gramma						
Aug 05	0.250 (0.087)	0.490 (0.09)	0.450 (0.121)	0.590 (0.030)	0.280 (0.019)		
Sep 06	0.220 (0.06)	1.340 (0.225)	0.750 (0.052)	1.050 (0.121)	0.330 (0.024)		
% Change	-6	+45	+25	+28	+8		

Table 2 Mean densities (\pm SE) of Centrostephanus rodgersii, Haliotis rubra and Heliocidaris erythrogramma (m-2) prior tomanipulations (August 2005) and 13 months after the initial manipulations (September 2006) at the Lanterns, Tasmania, Australia

Treatments are, unmanipulated *C. rodgersii* barrens (T1), removal of *C. rodgersii* and all regrowth from patches (T2), removal of *C. rodgersii* and canopy-algae regrowth from patches (T3), removal of *C. rodgersii* only (T4), control no *C. rodgersii* in intact algal patches (T5)

Removals of C. rodgersii and canopy-algae regrowth from incipient barrens patches (T3) resulted in significantly higher cover of filamentous algae, understorey foliose algae and sessile invertebrates when compared with the unmanipulated barrens patches (T1) (Table 3; Figs. 2, 3). There were no detectable differences in the density and cover of canopy-algae and cover of encrusting red algae and sessile invertebrates between the treatment patches (T3) and the unmanipulated barrens patches (T1). There was a significantly lower density and cover of canopy-algae and cover of encrusting red algae but no detectable differences in the cover of filamentous algae, foliose understorey algae between the treatment patches (T3) and the intact algal patches (T5). In general, the trends were similar between sites however the cover of overstorey algae, encrusting red algae and filamentous algae, but not sessile invertebrates, in the intact algal patches was higher at site A relative to site B (Table 3; Figs. 2, 3). The cover of foliose algae and sessile invertebrates in the treatment from which C. rodgersii only was removed was also higher at site A than at site B (Table 3; Figs. 2, 3).

Density of urchins and abalone

Throughout the experiment the density of *C. rodgersii* was higher in the unmanipulated barrens patches (T1) than in the patches from which the urchins were

removed (T2, T3, T4) and the intact algal patches (T5) (Appendix 1 in ESM), indicating that manipulations were successful in maintaining removal patches at very low densities of *C. rodgersii*.

Prior to the manipulations, the densities of Haliotis rubra in the C. rodgersii incipient barrens patches (T1, T2, T3 and T4) were similar, and much lower than the densities of abalone in the intact algal patches (T5) (Fig. 4). After 13 months, there was a significantly higher density of *H. rubra* in the patches from which *C*. rodgersii and canopy-algae regrowth was removed (T3) relative to the unmanipulated C. rodgersii barrens patches (T1) (T = 4.583, P = 0.001, α adjusted = 0.025) (Fig. 4). There was a higher density of *H. rubra* in the patches from which only C. rodgersii was removed (T4) relative to the patches from which C. rodgersii and canopy-algae regrowth were removed (T3) $(T = 3.672, P = 0.004, \alpha \text{ adjusted} = 0.025).$ There was also a higher density of H. rubra in the patches which C. rodgersii and canopy-algae regrowth (T3) were removed relative to the patches from which C. rodgersii and all regrowth were removed (T2) $(T = 7.007, P < 0.001, \alpha \text{ adjusted} = 0.025)$ (Fig. 4). In contrast, there were no detectable differences in the densities of H. rubra between the patches from which C. rodgersii and all regrowth (T2) were removed and the unmanipulated C. rodgersii barrens patches (T1) $(T = 0.347, P > 0.05, \alpha \text{ adjusted} = 0.025)$ (Fig. 4).



Fig. 1 Ordinations (nMDS) of benthic community structure, showing the relationship between experimental treatments (n = 3 replicates) at **a** 0 month prior to manipulations (August 2005), and **b** 13 months after manipulations (September 2006), at two sites at the Lanterns, Tasmania, Australia. Before manipulations, at both sites the community assemblage in the control patches without *C. rodgersii* (T5) differed to all other treatments (*top panels*). After 13 months, clear differences in community structure were evident among several treatments while patches subject only to removal of *C. rodgersii* (T4)

At the initial assessment, the density of *Heliocidaris erythrogramma* in the barrens (T1, T2, T3, T4) and intact algal patches (T5) was similar (Fig. 5). After 13 months, the densities of *H. erythrogramma*

converged with the control plots (T5) (*bottom panels*). Treatments are, *upright triangles* unmanipulated *C. rodgersii* barrens (T1), *diamonds* removal of *C. rodgersii* and all regrowth from patches (T2), *downward triangles* removal of *C. rodgersii* and canopy-algae regrowth from patches (T3), *squares* removal of *C. rodgersii* only (T4), *circles* no *C. rodgersii* in intact algal patches (T5). The analysis is based on a Bray-Curtis matrix of square root transformed percentage cover data. Ellipses (95 % confidence interval) are drawn around all treatments for clarity

were significantly higher in the patches from which *C. rodgersii* and all regrowth were removed (T2) (T = 6.306, P < 0.001, α adjusted = 0.025) and the patches from which *C. rodgersii* and canopy-algae

 Table 3 Results of 3-way univariate ANOVAs and planned comparisons testing the effects of the manipulations on the density of stipes of canopy-algae, and the percentage cover of canopy

algae, bare rock, encrusting red algae, filamentous algae, understorey foliose algae and sessile invertebrates in treatment patches, at two sites at the Lanterns, Tasmania, Australia

Factors	df	MS	F	Р	Tests	Т	Р
Density of stipes of canopy-algae					T2 versus T1	0.351	>0.05
Treatment	4	0.340	6.384	0.049	T2 < T5	14.698	<0.001
Site	1	0.114	1.816	< 0.05	T3 versus T1	2.015	0.035
Treatment × Site	4	0.063	1.656	< 0.05	T3 < T5	-4.714	0.004
Patch (Treatment \times Site)	30	0.038			T4 > T1	-9.671	<0.001
Error	80				T4 versus T5	-4.288	<0.001
Cover of canopy-algae $[log(Y + 0.001)]$					T2 versus T1	0.311	>0.05
Treatment	4	599.660	5.818	0.049	T2 < T5	16.200	<0.001
Site	1	153.950	110.108	>0.001	T3 versus T1	1.666	>0.05
Treatment × Site	4	103.070	73.714	>0.001	T3 < T5	16.2	<0.001
Patch (Treatment \times Site)	30	1.400	0.311	>0.001	T4 > T1	-25.19	<0.001
Error	80	4.500			T4 versus T5	5.241	<0.001
Cover of encrusting red algae					T2 versus T1	0.859	>0.05
Treatment	4	4292.100	14.187	0.013	T2 < T5	4.878	<0.001
Site	1	64.500	0.346	< 0.05	T3 > T1	1.369	>0.05
Treatment \times Site	4	1210.200	1.621	< 0.05	T3 versus T5	-6.284	>0.05
Patch (Treatment \times Site)	30	5599.700	1.062	< 0.05	T4 > T1	-4.402	<0.001
Error	80	14066.700			T4 versus T5	-2.553	>0.05
Cover of filamentous algae					T2 versus T1	3.964	0.002
Treatment	4	205.030	9.585	< 0.05	T2 < T5	1.523	>0.05
Site	1	320.130	14.966	>0.001	T3 > T1	2.82	0.003
Treatment × Site	4	136.990	6.404	>0.001	T3 versus T5	-3.09	0.003
Patch (Treatment \times Site)	30	12.560	0.587	< 0.05	T4 > T1	-3.017	0.002
Error	80	21.390			T4 versus T5	2.760	0.008
Cover of foliose understorey algae $[log(Y + 0.001)]$					T2 versus T1	2.569	0.020
Treatment	4	57.970	0.98	< 0.05	T2 < T5	-8.477	<0.001
Site	1	205.870	36.684	>0.001	T3 > T1	3.87	<0.001
Treatment × Site	4	59.170	10.540	>0.001	T3 versus T5	0.358	>0.05
Patch (Treatment \times Site)	30	5.610	0.358	0.001	T4 > T1	-3.695	0.001
Error	80	15.700			T4 versus T5	0.358	>0.05
Cover of sessile invertebrates					T2 versus T1	2.144	0.04
Treatment	4	665.300	9.843	0.020	T2 < T5	3.000	0.002
Site	1	0.100	0.007	< 0.05	T3 > T1	3.503	0.001
Treatment × Site	4	67.900	0.835	< 0.05	T3 versus T5	-1.607	>0.05
Patch (Treatment × Site)	30	108.900	0.840	< 0.05	T4 > T1	-5.239	<0.001
Error	80	82.855			T4 versus T5	-0.829	>0.05

Treatments are, unmanipulated *C. rodgersii* barrens (T1), removal of *C. rodgersii* and all regrowth from patches (T2), removal of *C. rodgersii* and canopy-algae regrowth from patches (T3), removal of *C. rodgersii* only (T4), control no *C. rodgersii* in intact algal patches (T5). Significant *P* values are shown in bold face: P < 0.05 is significant for the main analysis and P < 0.0125 is significant for the t-tests (α adjusted after Todd and Keough 1994)

(T3) were removed (T = 7.368, P < 0.001, α adjusted = 0.025) relative to the unmanipulated *C. rodgersii* barrens (T1) (Fig. 5). In contrast, there

was no detectable difference in the density of *H. erythrogramma* in the patches from which *C. rodgersii* and all regrowth (T2) were removed and those from



Fig. 2 Mean (\pm SE) (i) density (m⁻²) and (ii) percentage cover of canopy-algae in treatment patches (n = 3 replicates) through time (months), at two sites at the Lanterns, Tasmania, Australia. Treatments are, *upright triangles* unmanipulated *C. rodgersii* barrens (T1), *diamonds* removal of *C. rodgersii* and all regrowth

from patches (T2), *downward triangles* removal of *C. rodgersii* and canopy-algae regrowth from patches (T3), *squares* removal of *C. rodgersii* only (T4), *circles* control no *C. rodgersii* in intact algal patches (T5). Note the different scale on the y-axes

which only *C. rodgersii* was removed (T4) (T = 2.333, P = 0.042, α adjusted = 0.025) (Fig. 5). There was also no detectable difference in the density of *H. erythrogramma* in the patches from which only *C. rodgersii* was removed (T4) relative to patches from which both *C. rodgersii* and canopy-algae regrowth were removed (T3) (T = 2.564, P = 0.038, α adjusted = 0.025) (Fig. 5).

Discussion

Effect of habitat modifying species on benthic structure and function

Habitat modifying species can have major impacts on marine ecosystem structure and function (Helmuth et al. 2006; Williams and Grosholz 2008). In many



Fig. 3 Mean (\pm SE) percentage cover of (i) encrusting red algae, (ii) filamentous algae, (iii) foliose algae, (iv) sessile invertebrates in treatment patches (n = 3 replicates) through time (months), at two sites at the Lanterns, Tasmania, Australia. Treatments are, *upright triangles* unmanipulated *C. rodgersii*

barrens (T1), *diamonds* removal of *C. rodgersii* and all regrowth from patches (T2), *downward triangles* removal of *C. rodgersii* and canopy-algae regrowth from patches (T3), *squares* removal of *C. rodgersii* only (T4), *circles* control no *C. rodgersii* in intact algal patches (T5). Note the different scale on the y-axes

ecosystems, urchins are well known habitat modifiers (Lawrence 1975; Chapman 1981; Chapman and Johnson 1990; Tegner and Dayton 2000). It is already well recognised that *Centrostephanus rodgersii* grazing has an important influence on the benthic community assemblage in south east Australia (Fletcher 1987; Johnson et al. 2005, 2011; Ling 2008). We demonstrate here that in its new habitat on

the east coast of Tasmania, this urchin is responsible for overgrazing the filamentous and foliose algal and sessile invertebrates and maintaining simplistic and homogeneous bare rock benthic habitat which is similar to the barrens described in its endemic range (Fletcher 1987) and broadly typically of urchin barrens habitat throughout the world (Pinnegar et al. 2000).



Fig. 4 Mean densities (\pm SE) of *H. rubra* (m⁻²) in treatment patches (n = 3 replicates) through time (months), at two sites at the Lanterns, Tasmania, Australia. Treatments are, *upright triangles* unmanipulated *C. rodgersii* barrens (T1), *diamonds* removal of *C. rodgersii* and all regrowth from patches (T2), *downward triangles* removal of *C. rodgersii* and canopy-algae regrowth from patches (T3), *squares* removal of *C. rodgersii* only (T4), *circles* control no *C. rodgersii* in intact algal patches (T5). Note the different scale on the y-axes

In our study, experimental removals of C. rodgersii from barrens patches resulted in bare rock being overgrown by filamentous algae (primarily red algae), foliose algae (red, juvenile canopy-forming and understorey foliose brown algae) and sessile invertebrates. However, after 13 months there were still differences in the benthic assemblage between experimental patches where only C. rodgersii was removed and control patches in intact algal beds. Similar to another study on the east coast of Tasmania (Ling 2008), removals of C. rodgersii from barrens patches resulted in a rapid return to an algal dominated state $(\geq 50 \%$ cover), however recovering patches were biased towards smaller and more abundant canopy forming algae and a lower cover of encrusting red algae, relative to the community of the long standing intact algal beds. Complete recovery of the algal community following the removal of C. rodgersii can take 2-3 years (Ling 2008), and is likely to be a



Fig. 5 Mean densities (\pm SE) of *H. erythrogramma* (m⁻²) in treatment patches (n = 3 replicates) through time (months), at the Lanterns, Tasmania, Australia. Treatments are, *upright triangles* unmanipulated *C. rodgersii* barrens (T1), *diamonds* removal of *C. rodgersii* and all regrowth from patches (T2), *downward triangles* removal of *C. rodgersii* and canopy-algae regrowth from patches (T3), *squares* removal of *C. rodgersii* only (T4), *circles* control no *C. rodgersii* in intact algal patches (T5). Note the different scale on the y-axes

function of the size of the cleared area and its proximity to established reproductive algal populations (Fletcher 1987). The recovered patches and intact algal beds provide increased 3 dimensional structural complexity (Graham 2004; Ling 2008), primary and secondary productivity (Chapman 1981; Duggins et al. 1989) and altered nutrient cycling and energy flows (Sauchyn and Scheibling 2009) relative to barrens habitat.

Effect of habitat modifying species on abundances of native species

Range expanding habitat modifiers can also alter biotic interactions among native species (Firth et al. 2009; Strain and Johnson 2009; Sorte et al. 2010; Walther 2010). Similar to research in large scale plots $(1,000 \text{ m}^{-2})$ in New South Wales (Andrew et al. 1998), we demonstrated that experimental removals of C. rodgersii from barrens patches (mean size 30.654 m^{-2}) resulted in overgrowth of bare rock by filamentous, foliose algae and sessile invertebrates and concomitant increases in the abundances of abalone Haliotis rubra. In other manipulations in Tasmania, the introduction of C. rodgersii into cages (9 m^{-2}) in intact algal beds resulted not only in declines in the percentage cover and standing biomass of foliose algae, but also reduced total weight, dry weight of stomach contents and survivorship of H. rubra relative to controls without C. rodgersii (Strain and Johnson 2009). The combined research strongly suggests that C. rodgersii is a more efficient and effective grazer of attached understorey algae than H. rubra (Andrew et al. 1998; Strain and Johnson 2009). We extend this research by demonstrating that although C. rodgersii is the superior competitor for attached algae in interactions with H. rubra, there is no evidence to suggest that these two herbivores also compete for preferred habitat.

Centrostephanus rodgersii overgrazing of canopyalgae could also potentially increase H. rubra susceptibility to predation (Andrew 1993; Andrew et al. 1998; Edgar et al. 2004) by reducing canopy algae cover. Our observations suggest there were fewer H. rubra, but more rock lobsters and fish predators in patches from which C. rodgersii and canopy-algae regrowth were removed relative to patches where C. rodgersii was removed but regrowth allowed. Since this manipulation does not decrease the availability of food to abalone (because they do not feed on established canopy-forming algae), this result could suggest that predation on abalone is higher in the absence of the canopy, abalone seek cryptic microhabitat in the interstices of the reef, and/or do not return to patches where canopy algae are absent. Our results are consistent with other research in Tasmania, in which removals of canopy-algae from large experimental plots (600 m⁻²) resulted in decreased abundances of H. rubra and an increase in the density of empty abalone shells (Edgar et al. 2004) relative to controls in intact algal beds. The combined results suggest increased predation on abalone in the absence of large canopy-forming algae. In the reverse manipulation, Andrew (1993) found significant increases in the densities of juvenile abalone (H. rubra and H. coccoradiata) after boulders covered in Ecklonia radiata were transplanted into urchin barrens habitat $(10,000 \text{ m}^{-2})$. These results suggest that the continued expansion of *C. rodgersii* on the east coast of Tasmania will have an ongoing negative impact on the abundances of *H. rubra* both through loss of food resources (Shepherd 1973; Strain and Johnson 2009) and structural complexity (Andrew 1993; Andrew et al. 1998; Edgar et al. 2004).

In contrast, to the response of abalone, removals of C. rodgersii from barrens patches, irrespective of our manipulations of regrowth, invariably resulted in an increase in the densities of H. erythrogramma relative to unmanipulated C. rodgersii barrens patches. However there were no detectable differences in the densities of the native urchin between the patches from which C. rodgersii and all regrowth were removed and the patches from which C. rodgersii was removed and regrowth allowed. These results strongly suggest that C. rodgersii outcompetes H. erythrogramma through competition for preferred habitat rather than for attached algal food resources. Studies on Diadematidae species have demonstrated that urchins aggressively defend their crevices by biting and pushing conspecifics and congeners (Williams 1977; McClanahan 1988; Shulman 1990) or by dislodging them from areas which are more favourable for catching drift algae, which is an important food resource for urchins inhabiting barrens habitat (Harrold and Reed 1985; Vanderklift and Kendrick 2005; Vanderklift and Wernberg 2008). A similar interaction could be operating in Tasmania although we have never observed it, perhaps because C. rodgersii is highly nocturnal (Flukes et al. 2012). Further manipulations of the availability of crevices and drift algae are required to elucidate the detailed effects of C. rodgersii on H. erythrogramma.

Our experiment also provides insight into the interactions between the two dominant native macroherbivores on the southeast coast of Australia, H. rubra and H. erythrogramma. Interestingly, throughout the 13 months experiment, in the intact algal patches without C. rodgersii, there was a low density of H. erythrogramma but a high density of H. rubra. These results are consistent with broad-scale surveys along the east coast of Tasmania and elsewhere in southeast Australia which have showed that abundances of H. erythrogramma and H. rubra are negatively correlated at a range of spatial scales (Shepherd 1973; Johnson et al. 2005). The nature and effects of interactions between the two native herbivores are poorly understood. However, our results provide some support for the hypothesis that these two native herbivores could also compete for food and/or preferred habitat (Shepherd 1973; Johnson et al. 2005).

Alternatively, intact algal beds on moderately exposed coastlines could be unsuitable to support high abundances of H. erythrogramma. Studies on the east coast of Tasmania and elsewhere in southern Australia, have demonstrated that H. erythrogramma are relatively immobile and remain in crevices in intact algal beds in wave exposed habitat (Connolly 1986; Keesing 2006). In these areas, the whiplash like action of canopy-algae is thought to limit the attachment and destructive grazing behaviour of this urchin (Konar 2000; Ling et al. 2010). Certainly, surveys have demonstrated that H. erythrogramma occurs in abundance only in sheltered and, at worst, moderately exposed sites (Johnson et al. 2005). Overall, our results suggest that the initial establishment of C. rodgersii on the east coast of Tasmania and associated overgrazing of canopy-algae might initially benefit populations of H. erythrogramma but, as the densities of the non-native urchin increase, the native urchin will be locally displaced.

Studying invasions

Range expanding species can have complex biotic interactions (competition and/or predation) with native species (Sorte et al. 2010; Walther 2010). A useful extension to this study would be to quantify the rates of predation on H. erythrogramma and H. rubra in C. rodgersii barrens patches, and compare this to predation on these species at comparable sites with intact algal beds. There is already strong evidence that both fishes and rock lobsters have an important influence on the behavior and survival of H. erythrogramma and H. rubra in intact algal beds in marine protected areas in Tasmania, Australia (Pederson and Johnson 2006; Pederson et al. 2008) but the effects of these predators in incipient barrens patches remain unclear. Irrespective, we demonstrate here that C. rodgersii appears to have a stronger effect on the densities of H. rubra and H. erythrogramma through competition for food and preferred habitat rather by overgrazing of canopy-algae per se.

It is predicted that numerous marine species will shift their range polewards in response to global climate change (Parmesan and Yohe 2003). While many of these species will have little or no effect on marine ecosystems (Sorte et al. 2010; Walther 2010), highly invasive range expanding species are often generalist grazers (e.g. sea urchins, gastropods and sea stars) which modify habitat by consuming a wide variety, and large amounts, of prey (Helmuth et al. 2006; Williams and Grosholz 2008; Sorte et al. 2010; Walther 2010). Thus, scientists and managers should focus on developing strategies to control the abundances of newly established generalist grazers to limit their impacts on biodiversity and commercial fisheries. Continued research into the impacts of range expanding species on the biotic interactions between native species and their environment is important for predicting and understanding and managing the effects of invaders on marine community dynamics and ecosystems function and structure (Bertness 1984; O'Connor and Crowe 2005; Ling 2008; Firth et al. 2009).

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