

# Rapid recovery of fauna following simulated ice rafting in a Nova Scotian seagrass bed

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**ABSTRACT:** The shallow portion of the seagrass, *Zostera marina* L., bed in Pomquet Harbour, Nova Scotia, Canada, is subject to patchy disturbance by ice rafting, in which areas of seagrass are frozen into the underside of ice, which breaks up into large pieces and floats away. To examine the potential effect of small-scale removal of patches of eelgrass on the seagrass fauna, ice rafting was simulated by clearing 1.2 × 0.4 m patches in the spring. Eelgrass recovery on the experimentally cleared patches was relatively slow, and 4 mo later the biomass was still significantly lower on the cleared patches than in the surrounding seagrass bed. In contrast, recovery of macroalgae was much more rapid, biomass approximating control levels by 1.5 mo after disturbance. Clearing also had a relatively short-lived effect on most invertebrate species, numbers of all species on the cleared patches being at or above those in the surrounding seagrass bed 3 to 4 mo after clearing. The recovery of epifauna was significantly linked to macroalgal biomass on cleared patches, indicating that the rapid return of macroalgae to cleared patches enabled this community to recover rapidly after disturbance despite the slower seagrass recovery.

## INTRODUCTION

Disturbance is widely acknowledged to have a fundamental effect in many communities (White & Pickett 1985). Most studies have focussed on communities of sessile organisms, in which disturbance often creates open areas by the removal of established individuals, permitting colonization by other species. This pattern has been observed in a wide variety of systems including marine intertidal macroalgae (Sousa 1985, Heine 1989), salt marshes (Bertness & Ellison 1987), grasslands (Hobbs & Mooney 1985), and temperate forests (Runkle 1985), though sometimes the previous occupant recovers first and no change in species composition is observed (Dethier 1984, Kennelly 1987). Small-scale disturbances remove only a few individuals at a time, producing a heterogeneous mosaic of patch types, the character of each reflecting its history of disturbance and its course of recolonization (Sousa 1979, Runkle 1985). The effect of small-scale disturbances on mobile animals, either through direct detrimental effects on populations or indirect effects via altera-

tion in vegetation habitat, has received less attention. The latter effect is the focus of this study.

Macrophytes in vegetated aquatic systems provide much of the physical infrastructure upon which the system is based; indicative of this close relationship is the observation that invertebrate abundance and diversity can often be related to macrophyte biomass (Heck & Orth 1980, Stoner 1980, Bronmark 1985, Friday 1987). In these systems, studies of animal responses to varying vegetation composition have focussed on selection based on feeding (Nicotri 1980, Norton & Benson 1983, Hay et al. 1988) or on suitability as habitable space. Many animals are dependent upon the presence of refuges both from physical effects, such as water currents (Orth 1977, McRoy & Helfferich 1980), and from predators (Crowder & Cooper 1982, Summerson & Peterson 1984, Joyce & Weisberg 1986).

Disturbances not only alter the amount of vegetation, but also its patchiness. There are few studies of the response of mobile animals to gaps in vegetation cover in aquatic systems, but many terrestrial species forage selectively in gaps (Mills 1986, Crome & Richards 1988). There is some indication that animals in seagrass beds migrate to open areas both within and outside the

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bed (Holt et al. 1983, Summerson & Peterson 1984) and that bare areas may function as predator feeding loci in seagrass beds (Heck & Orth 1980) and salt marshes (Zimmerman & Minello 1984).

Seagrass systems are subject to a variety of natural small-scale disturbances including ice scouring (Robertson & Mann 1984, Schneider 1990), wave-generated erosion (Patriquin 1975) and herbivore feeding (Orth 1977, Williams 1988), but little attention has been paid to effects on mobile animals. It has been widely demonstrated that faunal densities and species diversity are much higher in seagrass beds than in adjacent unvegetated areas (Stoner 1980, Orth et al. 1984, Summerson & Peterson 1984). This relationship has been attributed to the effects of seagrasses on physical characteristics, such as current speed and sediment stability (Orth 1977, Peterson et al. 1984), or modification of biotic interactions such as predation (Nelson 1979, Heck & Thoman 1981, Summerson & Peterson 1984). Invertebrate assemblages also vary between areas of different seagrass species (Lewis 1984, Virnstein & Howard 1987a), between seagrass plants and bare areas (Lewis & Stoner 1983), and between seagrass and macroalgae (Lewis 1987, Virnstein & Howard 1987b, Schneider & Mann 1991a). This variation among microhabitats has been linked to differences in predation, food availability, and other species-specific habitat requirements (Heck & Orth 1980, Coen et al. 1981, Leber 1985, Schneider & Mann 1991b). Given this effect on a small habitat scale, disturbance-induced changes in vegetation cover should have a significant effect on invertebrate distribution.

An eelgrass bed in Pomquet Harbour, Nova Scotia, Canada, was chosen for study of disturbance by ice. The harbour is covered with ice up to 0.7 m thick from January to mid-March. The rise and fall of the tide has the effect of breaking the ice into sections, or 'pans' each having a surface area of one to a few m<sup>2</sup>. In shallow water these pans freeze to the eelgrass and sediment surface at low tide, but when the tide rises they may lift a patch of eelgrass, complete with rhizomes and roots, out of the sediment. In spring, when the ice began to break up, it was found that a number of pans were moving away with eelgrass frozen into their undersides. This is known as 'ice rafting'.

Examination of the eelgrass bed revealed bare patches from which all eelgrass, including rhizomes and roots, had been removed. In spring, the patches were most abundant in a zone 2 m wide parallel with the inshore edge of the eelgrass bed (which was at mean high tide level). In a zone 3 m further into the eelgrass bed (mean depth 0.5 m at mid-tide) the frequency of bare patches decreased, and in the next 3 m wide zone (mean depth 0.8 m) the incidence of bare patches decreased to zero.

Confirmation that the bare patches were caused by ice rafting in winter was obtained by aerial photography from a light plane in spring and fall of 1987 and 1988. In the most shallow edge zone the proportion lacking eelgrass cover was 55% in spring 1987, decreasing to 35% in fall and increasing to 67% the following spring. In the next zone the figures were 29, 2 and 28%. In the deepest of the 3 zones the figures were 7, 0 and 2%.

In an attempt to understand the dynamics of this process, and especially the effect of patch formation on the flora and fauna, ice rafting was mimicked by creating artificial patches of a standard size and comparing them at intervals with control patches with full eelgrass cover.

## METHODS AND MATERIALS

**Study site.** The study site was situated in an eelgrass, *Zostera marina*, bed on a muddy sand sediment in the sheltered inner basin of Pomquet Harbour, Nova Scotia (45°50' N, 61°90' W). The eelgrass bed contains large quantities of macroalgae, especially *Chondrus crispus* and *Gracilaria tikvahiae*, year round, and *Polysiphonia urceolata* and *Sphaerotrichia divaricata* in the summer. The first 2 species usually intermingle in freely drifting balls, while the latter 2 are typically epiphytic on larger macrophytes. When *Sphaerotrichia* occurs in large quantities it may break off and drift about as unattached clumps. A complete site description can be found in Schneider & Mann (1991a).

**Selection of artificial patch size.** Since natural patches differed in size, shape and location, replicated, standardized artificial gaps in the eelgrass cover were created. It was desirable to produce artificial cleared patches within the size range normally produced by ice scour. A survey of natural bare patches was conducted in early April 1986 along a 155 m transect within a 5 m wide strip along the inshore edge of the eelgrass bed. The longest axis of unvegetated patches and the maximum width perpendicular to this axis were found to average 1.2 m (range 0.3 to 2.3 m) and 0.6 m (range 0.1 to 1.0 m). Because these represented maximum dimensions, a rectangular patch of 1.2 × 0.6 m would have a greater area than the average of the natural patches. Therefore slightly smaller rectangular patches of 1.2 × 0.4 m were experimentally created.

**Experimental protocol.** The experiments were conducted in a plot which extended 40 m along the shore, with its inner edge 5 m and its outer edge 11 m from the shore. It was divided longitudinally into a shallow zone extending 5 to 8 m from shore, with a mean depth of 0.8 m, and a deep zone 8 to 11 m from shore, with a mean depth of 1.0 m. The shallower of these 2 areas

corresponded to the depths where scouring still occurred, but not extensively, while no ice-rafted patches were observed in the deeper zone. The overall  $40 \times 6$  m plot was divided into 160 sampling units of  $1.5 \times 1$  m. Of these 45 were randomly chosen to be manipulated and 45 as controls, with equal numbers in the shallow and deep zones. The sampling time of each was randomly assigned to one of 3 dates. Following the recommendation of Hurlbert (1984), adequate inter-spersion of treatments along the long axis of the plot was assured by constraining randomization such that equal numbers of replicates were collected from each half of the plot.

Experimental clearing was conducted during the month after ice breakup, i.e. in May 1986. The margins of the patches to be cleared were marked with a rope, the eelgrass rhizomes entering the patch were cut with a knife, and then eelgrass shoots, macroalgae and eelgrass rhizomes were removed with a hand rake. This procedure removed most epifauna along with the above-ground vegetation and many infauna with the eelgrass rhizomes. As this situation was thought to mimic natural ice rafting of vegetation, no attempt was made to dig out any remaining infauna. The 4 corners of the cleared patches were marked with stakes so that they could be found at later dates.

Samples were collected from the centers of cleared patches and control units. Infaunal samples were taken with a 10 cm diameter clear plastic corer inserted 15 cm into the sediment in June and August 1986. Epifaunal samples were taken with a sampler based on the design of Virnstein & Howard (1987a), which consisted of 2 halves of a  $30 \times 30$  cm box, 10 cm wide in total, connected by a piano hinge at one end (Fig. 1). The sampler frame was constructed of 5 cm wide strips of aluminum and the large sides of each half were covered with 1.0 mm plastic mesh. Samples were collected while wading or snorkeling. The use of a face mark permitted observation of the sample collection procedure to ensure that no vegetation slipped out of the sampler. The sampler was placed with the hinged end perpendicular to the bottom, opened 30 cm and moved forward 30 cm. Care was taken to keep the sampler off the sediment and, after the sampler was closed, all external vegetation was clipped off. In this manner, the invertebrates within a single sample were directly associated with the vegetation enclosed, and uniform, quantitative samples were collected each time. The sampler enclosed a total volume of  $0.012 \text{ m}^3$ , associated with  $0.039 \text{ m}^2$  of bottom. Because the comparisons of interest were between numbers in control and manipulated samples, all animal numbers and emergent macrophyte biomass were reported on a per sample basis. Epifaunal samples were taken in July and September 1986 and again in July 1987

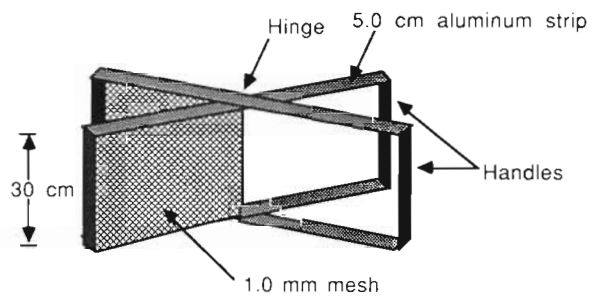


Fig. 1 Sampler used to collect epifauna and vegetation

Infaunal samples were washed through 1.0 mm sieves and the epifaunal samples through 0.5 mm sieves (the smaller mesh size reduced damage to delicate amphipods caused by washing). Vegetation and animals longer than 1 mm retained on the screens were frozen until sorted. Changes in biomass of the major vegetation types subsequent to the initial clearing were estimated from the amount of vegetation collected in the epifaunal sampler. Vegetation was sorted into 3 categories: (i) eelgrass, (ii) coarsely branched algae (*Chondrus crispus* and *Gracilaria tikvahiae*), and (iii) finely branched algae (mainly *Sphaerotrichia divaricata*), and dried to constant weight at  $80^\circ\text{C}$ .

**Statistical analysis.** To minimize the effects of seasonal changes in the abundance of organisms, the analysis was concentrated on a comparison of manipulated and control plots on each sampling date. For each data set (vegetation biomass, epifaunal numbers, infaunal numbers) analysis of variance, ANOVA, was performed using treatment (cleared vs control), depth zone (shallow vs deep), and time (since experimental clearing) as factors. All data were transformed  $\ln(Y + 1)$  and Bartlett's test for homogeneity of variances showed that this procedure successfully standardized variances. ANOVA revealed which factors produced significant effects in the experiment as a whole. A posteriori pairwise Scheffé contrasts between cleared and control patches from the same depth zone and date were used to assess significant differences when ANOVA showed significant interactions with treatment main effects.

To determine the direct relation of animal numbers on experimentally cleared patches to vegetation biomass within the same patches, regression analyses were conducted on samples from the cleared patches (all dates were pooled). For both the infaunal and epifaunal samples, the dependent variable was animal number  $[\ln(Y + 1)]$  transformed to stabilize variances and the independent variable was vegetation biomass (dry wt).

To determine whether species too rare to be examined individually were affected by experimental clearing, patterns of variation in the total invertebrate

community were described using principal components analysis, PCA. Data were transformed  $\ln(Y+1)$  as recommended by Pielou (1984) and plotted separately for cleared and control samples and for shallow and deep zones, according to their scores along the first 2 axes of the PCA. Separation among groups was estimated by visual inspection and mean factor scores were compared by ANOVA. The contribution of individual species to the patterns was assessed by examination of the species' loading along the factors of interest. Since animals were collected by both epifaunal and infaunal (core) samplers, the 2 sets of data were analyzed separately. Thus, 12 species are included in the epifaunal analysis and 7 in the infaunal. Three species normally classed with the infauna, *Gemma gemma*, *Mytilus edulis* and *Nereis succinea*, were also included in the epifaunal analysis (in the PCA only) because they are abundant in the vegetation and debris at the sediment-water interface and occurred abundantly in the samples collected by the epifaunal sampler.

A potential problem of PCA in many ecological applications is that its linear model assumes that species abundances change linearly along underlying gradients. Distortions are produced if this assumption is violated (Pielou 1984, Austin 1985). We had shown previously (Schneider & Mann 1991a) that the principal source of variation in animal abundance at a given time was a linear response to vegetation biomass, so the requirements of the method were met.

For comparison of faunal diversity on cleared and control areas 2 measures were calculated: the total number of species (species richness) and a modified Shannon-Wiener index (Peet 1974)

$$H' = \exp \left( - \sum_{i=1}^s p_i \ln p_i \right)$$

where  $s$  is the total number of species and  $p_i$  is the proportion of total individuals belonging to the  $i$ th species.

## RESULTS

### Vegetation recovery on manipulated patches

Recovery of eelgrass after experimental clearing was relatively slow (Fig. 2). It appeared to be entirely by lateral growth of plants at the edges, for no seedlings were found. After 1.5 mo the average biomass of eelgrass had reached only 3 to 4 % of the control values, and at the end of the summer considerable differences were still present. Even in July of the following year, eelgrass biomass on the cleared patches in the shallow zone was only one third of control values. This did not represent 14 mo of continuous regrowth as all plants shed large numbers of leaves in late fall and began to grow again in spring.

Analysis of variance (Table 1) showed that the differences between treated and control patches were highly

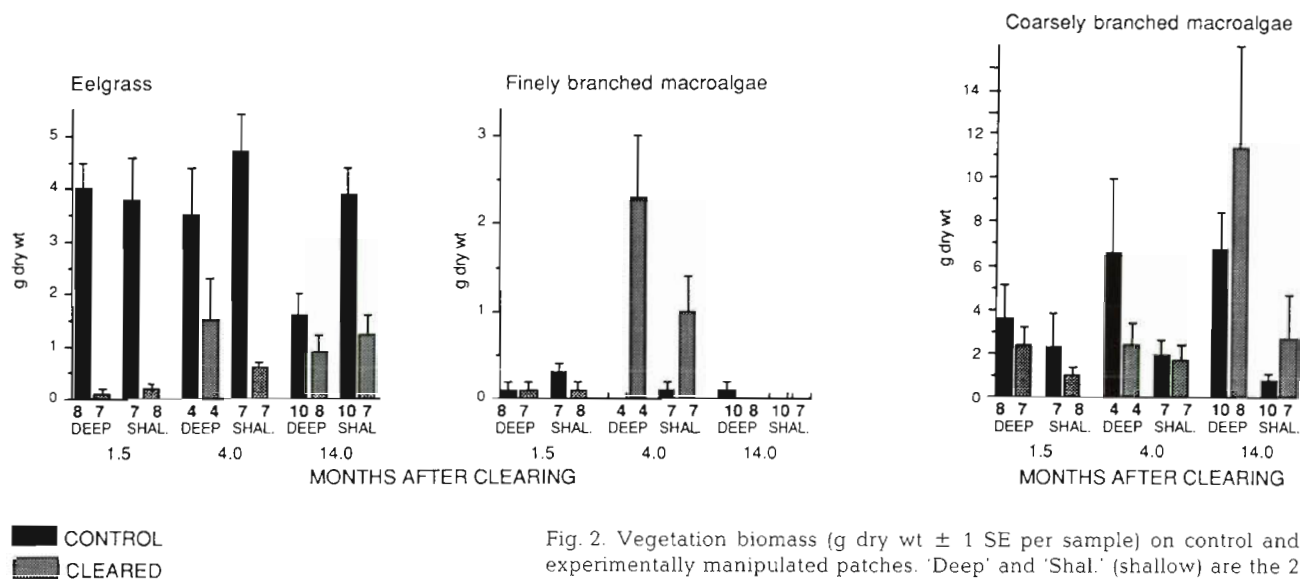


Fig. 2. Vegetation biomass (g dry wt  $\pm$  1 SE per sample) on control and experimentally manipulated patches. 'Deep' and 'Shal.' (shallow) are the 2 depth zones. Number of replicates is shown beneath each column



Table 1 *F*-values of ANOVA comparing vegetation biomass between control and experimentally cleared patches. ANOVA was performed on  $\ln(Y+1)$  transformed macrophyte dry weights using the following factors: 'Time' = time of sampling; 'Depth' = position in either shallow or deep zone; and 'Trt' = treatment, i.e. experimentally cleared vs control. Total  $n = 87$ ; \*\*  $p < 0.01$ ; \*  $0.01 < p < 0.05$

|                         | Time   | Depth  | Trt     | Time × Depth | Time × Trt | Depth × Trt | Time × Depth × Trt |
|-------------------------|--------|--------|---------|--------------|------------|-------------|--------------------|
| Degrees of freedom      | 2      | 1      | 1       | 2            | 2          | 1           | 2                  |
| Eelgrass                | 2.84   | 1.91   | 118.2** | 3.1*         | 9.9**      | 3.4         | 2.0                |
| Coarsely branched algae | 0.5    | 13.2** | 0.3     | 1.7          | 0.5        | 0.2         | 0.1                |
| Finely branched algae   | 24.3** | 2.8    | 23.1**  | 3.0*         | 32.8**     | 4.5*        | 4.0*               |

Table 2 *F*-values of a posteriori Scheffé contrasts comparing vegetation biomass on control and experimentally cleared patches within depth zones at single sampling times. Values were  $\ln(Y+1)$  transformed macrophyte dry weights. \*\*  $p < 0.01$ ; \*  $0.01 < p < 0.05$

|                         | July 1986 |         | September 1986 |         | July 1987 |         |
|-------------------------|-----------|---------|----------------|---------|-----------|---------|
|                         | Deep      | Shallow | Deep           | Shallow | Deep      | Shallow |
| Eelgrass                | 51.8**    | 42.2**  | 6.2*           | 33.2**  | 1.4       | 17.5**  |
| Coarsely branched algae | 0.1       | 0.3     | 0.5            | 0.1     | 0.1       | 0.3     |
| Finely branched algae   | 0.1       | 1.9     | 53.8**         | 21.5**  | 0.6       | 0.1     |

significant. Pairwise contrasts showed that eelgrass biomass differed significantly between control and cleared patches at all depths and times except for the deep zone in July 1987 (Table 2). The time × treatment interaction was also significant and this is in part due to the significant changes in eelgrass biomass on both deep and shallow treated plots between July and September 1986 (a posteriori pairwise contrasts,  $p < 0.01$ ).

In contrast, recovery of coarsely branched algae on the cleared patches was relatively rapid (Fig. 2). Analysis of variance (Table 1) showed no significant differences between manipulated and control patches at any sampling date.

Finely branched macroalgae were present in very small amounts at most sampling times, though there was a marked bloom on the cleared patches in September 1986 (Fig. 2). Analysis of variance (Table 1) showed a significant main treatment effect and significant time × treatment interaction. Pairwise contrasts showed that finely branched macroalgal biomass differed significantly between control and treated plots only in September 1986 (Table 2). The difference occurred in both deep and shallow zones.

#### Comparison of animal abundances on control and manipulated patches

The abundances of total epifauna (all crustaceans and gastropods collected in the epifaunal sampler),

total infauna (all bivalves and annelids collected in the infaunal sampler), and 6 individually abundant invertebrate species on the cleared and control patches are shown in Figs. 3 & 4.

Total epifaunal numbers were significantly decreased by clearing only on the first sampling date, 1.5 mo after clearing (significant time × treatment interaction, Table 3a; significant contrasts for July 1986 in both depth zones, Table 4a). Likewise, 2 of the individual epifaunal species examined, the tube-dwelling amphipod *Ampithoe* (primarily *A. longimana* though some *A. rubricata* were present) and the herbivorous snail *Bittium alternatum*, were significantly less abundant on manipulated than on control patches in the same depth zone 1.5 mo after clearing (significant treatment main effect and/or significant time × treatment interaction, Table 3a; significant contrasts for July 1986 in both depth zones, Table 4a). Four months after clearing no significant differences were observed. In contrast, the average abundance of the free-swimming amphipod *Gammarus mucronatus* tended to be very similar in manipulated and control plots, and analysis of variance showed no significant treatment effects.

Infaunal abundances were less affected by experimental clearing than the epifauna. For total infaunal numbers, neither the treatment main effect nor interactions involving treatment were significant (Table 3b). The polychaete *Nereis succinea* was significantly less abundant on cleared patches (in the shallow zone) on

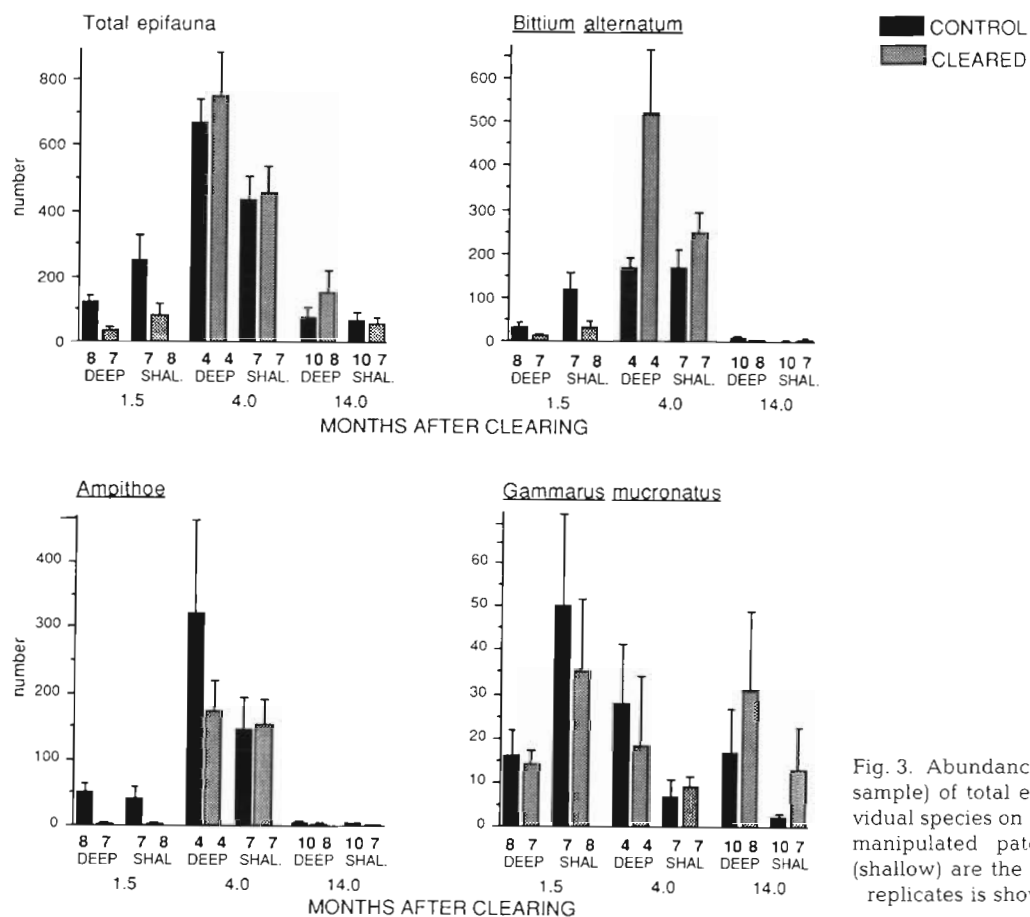


Fig. 3. Abundances (numbers  $\pm$  1 SE per sample) of total epifauna and selected individual species on control and experimentally manipulated patches. 'Deep' and 'Shal.' (shallow) are the 2 depth zones. Number of replicates is shown beneath each column

the first sampling date (significant treatment main effect and time  $\times$  treatment interaction, Table 3b; significant contrast for July 1986 in shallow zone, Table 4b). *Tellina agilis* displayed no significant differences between control and experimental patches.

*Gemma gemma* differed from all other species in that the average numbers in the cleared patches were significantly greater than in the controls (significant treatment effects, Table 3b). Numbers in the shallow zone were also consistently higher than in the deep (significant depth effect, Table 3b). In terms of individual pairwise comparisons between cleared and control patches, only the difference 3.0 mo after clearing in the shallow zone was significant (Table 4b).

#### Relationship of animal abundances to vegetation biomass on experimentally cleared patches

Regression analysis showed that total epifaunal numbers on experimentally cleared patches were signifi-

cantly related to the biomass of finely branched algae. There was a weak relationship with coarsely branched algae and no relation to eelgrass biomass (Table 5a). Examination of individual epifaunal species showed a strong relationship between the abundances of both *Bittium alternatum* and *Ampithoe* spp. and the biomass of finely branched macroalgae, and between the abundance of *Gammarus mucronatus* and coarsely branched macroalgae. No species was positively related to eelgrass biomass (*Gammarus* showed a negative relationship).

In contrast, abundances of infauna were not significantly positively related to vegetation biomass within the same sample (Table 5b). In fact, total infaunal numbers were negatively related to eelgrass root biomass in core samples, probably because of the numerical dominance of *Gemma gemma*, which tended to be negatively related to root biomass ( $0.05 < p < 0.10$ ). The other 2 infaunal species, *Nereis succinea* and *Tellina agilis*, were not significantly related to vegetation biomass within the samples.

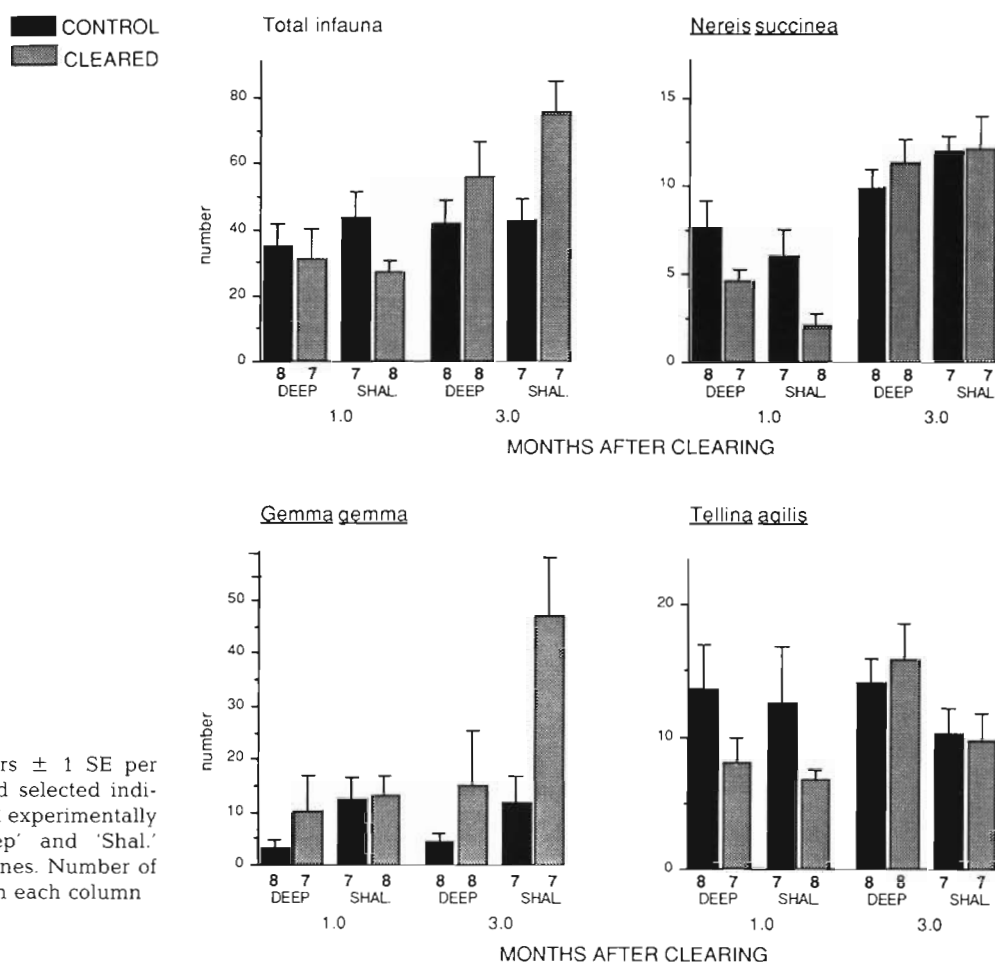


Fig. 4. Abundances (numbers  $\pm$  1 SE per sample) of total infauna and selected individual species on control and experimentally manipulated patches. 'Deep' and 'Shal.' (shallow) are the 2 depth zones. Number of replicates is shown beneath each column

Table 3. *F*-values obtained in an ANOVA of total epifauna, total infauna and selected individual species on control and experimentally cleared patches. ANOVA was performed on  $\ln(Y+1)$  transformed animal numbers using the following factors: 'Time' = time of sampling; 'Depth' = position in either shallow or deep zone; and 'Trt' = treatment, i.e. experimentally cleared vs control. \*\*  $p < 0.01$ ; \*  $0.01 < p < 0.05$

| Species                      | Time    | Depth  | Trt    | Time $\times$ Depth | Time $\times$ Trt | Depth $\times$ Trt | Time $\times$ Depth $\times$ Trt |
|------------------------------|---------|--------|--------|---------------------|-------------------|--------------------|----------------------------------|
| <b>(a) Epifauna (n = 87)</b> |         |        |        |                     |                   |                    |                                  |
| Degrees of freedom           | 2       | 1      | 1      | 2                   | 2                 | 1                  | 2                                |
| Total epifauna               | 34.6**  | 0.0    | 2.5    | 2.5                 | 3.0*              | 0.2                | 0.4                              |
| <i>Ampithoe</i> spp.         | 111.4** | 1.6    | 23.8** | 0.8                 | 14.4**            | 0.3                | 0.9                              |
| <i>Gammarus mucronatus</i>   | 7.7**   | 0.4    | 0.0    | 1.9                 | 0.7               | 0.2                | 0.7                              |
| <i>Bittium alternatum</i>    | 122.0** | 0.0    | 2.2    | 6.0**               | 8.3**             | 0.1                | 1.4                              |
| <b>(b) Infauna (n = 60)</b>  |         |        |        |                     |                   |                    |                                  |
| Degrees of freedom           | 1       | 1      | 1      | 1                   | 1                 | 1                  | 1                                |
| Total infauna                | 15.3**  | 2.0    | 0.2    | 0.1                 | 9.3               | 0.0                | 1.3                              |
| <i>Gemma gemma</i>           | 1.0     | 19.9** | 4.3*   | 0.2                 | 1.1               | 0.7                | 3.1                              |
| <i>Tellina agilis</i>        | 3.2     | 2.4    | 1.8    | 1.5                 | 1.2               | 0.1                | 0.0                              |
| <i>Nereis succinea</i>       | 54.3**  | 2.1    | 6.2*   | 6.8*                | 7.5**             | 1.8                | 0.5                              |

Table 4. *F*-values of a posteriori Scheffé contrasts comparing animal numbers on control and experimentally cleared patches within depth zones at single sampling times. Values were  $\ln(Y+1)$  transformed. \*\*  $p < 0.01$ ; \*  $0.01 < p < 0.05$

| <b>(a) Epifauna</b>        |           |         |                |         |           |         |
|----------------------------|-----------|---------|----------------|---------|-----------|---------|
|                            | July 1986 |         | September 1986 |         | July 1987 |         |
|                            | Deep      | Shallow | Deep           | Shallow | Deep      | Shallow |
| Total epifauna             | 5.2*      | 3.9*    | 0.0            | 0.0     | 0.5       | 0.6     |
| <i>Ampithoe</i> spp.       | 35.6**    | 21.1**  | 0.4            | 0.1     | 0.1       | 1.8     |
| <i>Gammarus mucronatus</i> | 0.1       | 0.3     | 0.9            | 0.4     | 0.6       | 0.6     |
| <i>Bittium alternatum</i>  | 4.9*      | 13.3**  | 2.3            | 1.0     | 2.0       | 0.1     |
| <b>(b) Infauna</b>         |           |         |                |         |           |         |
|                            | June 1986 |         | August 1986    |         |           |         |
|                            | Deep      | Shallow | Deep           | Shallow |           |         |
| Total infauna              | 0.7       | 3.3     | 1.3            | 5.2*    |           |         |
| <i>Gemma gemma</i>         | 0.9       | 0.0     | 0.1            | 7.7**   |           |         |
| <i>Tellina agilis</i>      | 1.4       | 1.5     | 0.0            | 0.1     |           |         |
| <i>Nereis succinea</i>     | 2.5       | 13.3**  | 0.2            | 0.0     |           |         |

#### Comparison of patterns of faunal similarity among samples as determined by principal components analysis

##### Species in epifaunal sampler

Principal components analysis distinctly separated control and manipulated samples 1.5 mo after clearing and average scores of these 2 groups for both Factors 1 and 2 were significantly different (Fig. 5a, significant treatment effect for July 1986, Table 6a). The samples from cleared plots occupy a distinct area characterized by low scores on Factor 1 and high scores on Factor 2. Four months after clearing, this separation cannot be made; the points are intermingled though average values of Factor 2 were still significantly different (Fig. 6b, significant treatment effect for September 1986, Table 6a). Examination of the species loadings for the first sampling date (July 1986, Table 7a) shows that for Factor 1 all species except one had a positive loading between 0.36 and 0.86, and the single negative value was small, contributing little to the scores. From this we see that negative scores on Factor 1, which characterized the cleared plots after 1.5 mo, arose from the low abundance of a wide range of species rather than from a change in the balance of species. The separation along the second factor appeared to be due to greater relative proportion of certain species, such as *Gemma gemma* and *Corophium insidiosum*, in the cleared patches. For the second sampling time, the slight separation of manipulated and unmanipulated samples along the second factor was due to higher weighting of some species, such as *Bittium alternatum* and *Ampithoe* spp., which in preceding sections were demonstrated to be associated with the finely branched algae that bloomed on the cleared patches at this time.

Table 5. Results of simple regression analysis relating  $\ln(Y+1)$  transformed animal numbers to dry weight of single vegetation types. Data were from cleared plots only and data from all sampling dates were pooled. \*\*\*  $p < 0.001$ ; \*\*  $0.001 < p < 0.01$ ; \*  $0.01 < p < 0.05$ ; +  $0.05 < p < 0.10$

|                              | Slope | R <sup>2</sup> | F-value |
|------------------------------|-------|----------------|---------|
| <b>(a) Epifauna (n = 40)</b> |       |                |         |
| Total epifauna               |       |                |         |
| Eelgrass                     | 0.09  | 0.00           | 0.1     |
| Coarsely branched algae      | 0.08  | 0.08           | 3.2*    |
| Finely branched algae        | 1.03  | 0.39           | 24.4*** |
| <i>Ampithoe</i> spp.         |       |                |         |
| Eelgrass                     | 0.42  | 0.04           | 1.6     |
| Coarsely branched algae      | -0.01 | 0.00           | 0.04    |
| Finely branched algae        | 1.49  | 0.53           | 43.8*** |
| <i>Gammarus mucronatus</i>   |       |                |         |
| Eelgrass                     | -0.70 | 0.17           | 8.1**   |
| Coarsely branched algae      | 0.13  | 0.24           | 11.9*** |
| Finely branched algae        | 0.02  | 0.00           | 0.0     |
| <i>Bittium alternatum</i>    |       |                |         |
| Eelgrass                     | 0.16  | 0.00           | 0.2     |
| Coarsely branched algae      | -0.03 | 0.00           | 0.2     |
| Finely branched algae        | 1.63  | 0.48           | 35.1*** |
| <b>(b) Infauna (n = 30)</b>  |       |                |         |
| Total infauna                |       |                |         |
| Eelgrass roots               | -1.50 | 0.29           | 11.2**  |
| Coarsely branched algae      | 0.14  | 0.00           | 0.1     |
| <i>Gemma gemma</i>           |       |                |         |
| Eelgrass roots               | -2.16 | 0.10           | 3.2+    |
| Coarsely branched algae      | -0.02 | 0.00           | 0.0     |
| <i>Tellina agilis</i>        |       |                |         |
| Eelgrass roots               | -0.21 | 0.01           | 0.2     |
| Coarsely branched algae      | 0.08  | 0.00           | 0.0     |
| <i>Nereis succinea</i>       |       |                |         |
| Eelgrass roots               | -0.61 | 0.03           | 0.9     |
| Coarsely branched algae      | 0.50  | 0.04           | 1.0     |



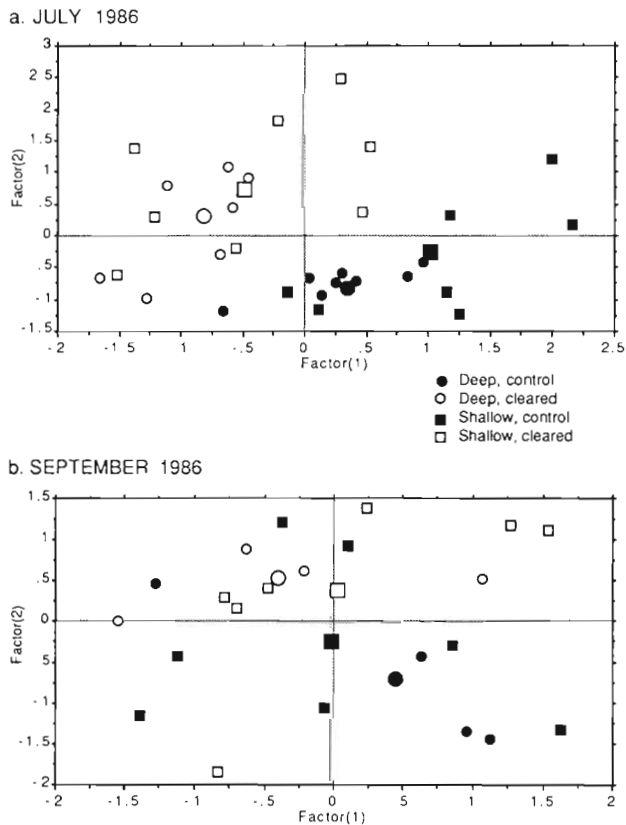


Fig. 5. Principal component scores of epifaunal samples for 2 sampling dates in 1986. The type of sample (cleared or control collected in the shallow or deep zones) is indicated. Means are represented by over-size symbols. Species loadings for factor axes are in Table 7a

For both dates, average scores along one of the 2 factors differed significantly among depths (Table 6a) but scores for depth along the factors in question were too intermingled to warrant further consideration (Fig. 5a, b).

#### Infauna

One month after experimental clearing, the samples from manipulated plots had a significantly lower average score on the first factor than those from control plots, though there was not a clear separation (Fig. 6a, June 1986, Table 6b). Factor loadings showed the difference between the 2 sets of samples was due to a greater relative abundance of *Gemma gemma* and *Spio* sp. on manipulated patches and a lower abundance of the remaining species except *Tellina agilis* (June 1986, Table 7b). This is indicated by the negative loadings of *Gemma* and *Spio* along Factor 1 (producing greater negativity of scores of the cleared plots) and positive

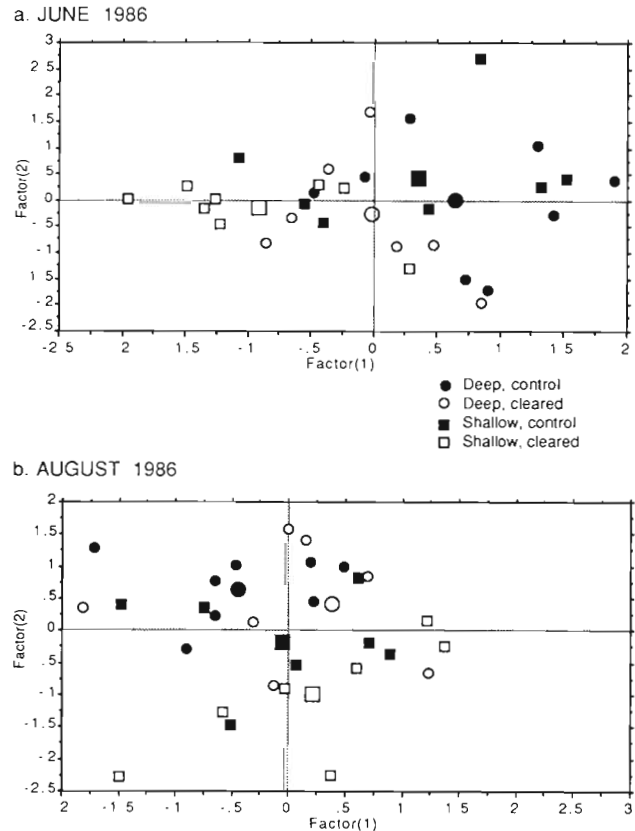


Fig. 6. Principal component scores of infaunal samples for 2 sampling dates in 1986. The type of sample (cleared or control collected in the shallow or deep zones) is indicated. Means are represented by over-size symbols. Species loadings for factor axes are in Table 7b

loading of most of the remaining species (producing the greater positivity of scores of the control plots). *Tellina*, which had a weighting near zero, did not contribute to the observed separation between experimental and control plots. Although the average scores were also significantly different between depth zones (June 1986, Table 6b), the scores of individual samples were too intermingled to warrant further consideration (Fig. 6a).

Three months after clearing there was no clear separation between manipulated and control samples though the average scores of control and treatment samples were significantly different along Factor 2 (August 1986, Table 6b). Separation between the deep and shallow zones, regardless of treatment, was marked along this factor and average scores were significantly different. This effect arose due to relatively higher numbers of certain species, particularly *Gemma gemma* and lower numbers of others such as *Tellina agilis* in the manipulated patches and/or the shallow zone (August 1986, Table 7b).

Table 6. *F*-values of ANOVA comparing principal component factor scores between control and experimentally cleared patches. ANOVA was performed using the following factors: 'Depth' = position in either shallow or deep zone; and 'Trt' = treatment, i.e. experimentally cleared vs control. \*\*  $p < 0.01$ ; \*  $0.01 < p < 0.05$

|                       | Depth  | Trt    | Depth × Trt |
|-----------------------|--------|--------|-------------|
| <b>(a) Epifauna</b>   |        |        |             |
| <u>July 1986</u>      |        |        |             |
| Degrees of freedom    | 1      | 1      | 2           |
| Factor (1)            | 6.3*   | 29.6** | 0.5         |
| Factor (2)            | 3.2    | 12.6** | 0.3         |
| <u>September 1986</u> |        |        |             |
| Degrees of freedom    | 1      | 1      | 2           |
| Factor (1)            | 0.0    | 0.4    | 0.7         |
| Factor (2)            | 0.1    | 5.1*   | 0.3         |
| <b>(b) Infauna</b>    |        |        |             |
| <u>June 1986</u>      |        |        |             |
| Degrees of freedom    | 1      | 1      | 2           |
| Factor (1)            | 5.3*   | 12.4** | 0.61        |
| Factor (2)            | 0.9    | 1.9    | 0.1         |
| <u>August 1986</u>    |        |        |             |
| Degrees of freedom    | 1      | 1      | 2           |
| Factor (1)            | 0.1    | 1.9    | 0.4         |
| Factor (2)            | 15.2** | 4.7*   | 1.0         |

### Comparison of diversity and richness on manipulated and control patches

Average epifaunal diversity and richness are shown in Fig. 7a. The numbers indicate that in every case the average diversity was lower on manipulated patches, but analysis of variance (Table 8a) showed that overall the effect of manipulation was not significant. Similarly with richness, no significant effects were observed (Table 8a).

For infauna, the average diversity was significantly lower in manipulated than in control samples (Fig. 7b, Table 8b). Clearing also significantly reduced species richness in deep and shallow sites but this effect changed with time (significant time × treatment interaction, Table 8b), disappearing by the 3.0 mo sampling date.

### DISCUSSION

Experimental clearing showed that eelgrass was relatively slow to colonize bare patches, having a considerably lower average biomass after 4 mo of the growing season, and even a year later having a reduced biomass compared with controls. It was there-

fore surprising to find that the invertebrate fauna showed much less response to clearing, since eelgrass is known to be of fundamental importance to the faunal community (e.g. Heck & Orth 1980, Stoner 1980). Total infaunal numbers were not decreased by clearing, and only one of 3 species examined individually was lower than in controls 1.0 mo after clearing. One clam, *Gemma gemma*, became more abundant on cleared patches. Epifaunal numbers showed greater response, as could be expected for organisms directly associated with vegetation. Total epifaunal numbers, and abundances of 2 of 3 individual species examined, were significantly lower on cleared patches 1.5 mo after clearing, but these had fully recovered by 4 mo after clearing.

For the epifauna, the key to this recovery appears to have been the colonization of bare patches by macroalgae. The colonizing algal species were the same as those that occurred in the surrounding undisturbed

Table 7. Amount of variation explained by first 2 factors of the PCA and standardized factor loadings showing the relative weights assigned to species

|                                 |  | Sampling time |       |                |       |
|---------------------------------|--|---------------|-------|----------------|-------|
|                                 |  | July 1986     |       | September 1986 |       |
| <b>(a) Epifauna</b>             |  |               |       |                |       |
| Factor number                   |  | 1             | 2     | 1              | 2     |
| % variation explained by factor |  | 38            | 18    | 30             | 24    |
| <i>Ampithoe</i> spp.            |  | 0.80          | -0.34 | 0.17           | 0.61  |
| <i>Bittium alternatum</i>       |  | 0.75          | -0.05 | 0.21           | 0.63  |
| <i>Corophium insidiosum</i>     |  | 0.62          | 0.58  | 0.88           | -0.12 |
| <i>Gammarus mucronatus</i>      |  | 0.66          | 0.37  | 0.84           | -0.08 |
| <i>Gemma gemma</i>              |  | -0.11         | 0.85  | 0.52           | 0.33  |
| <i>Idotea baltica</i>           |  | 0.65          | -0.39 | -0.13          | 0.17  |
| <i>Idotea phosphorea</i>        |  | 0.44          | -0.36 | 0.45           | 0.59  |
| <i>Jaera marina</i>             |  | 0.43          | -0.21 | 0.28           | -0.79 |
| <i>Leptochelia rapax</i>        |  | 0.63          | 0.54  | 0.43           | 0.69  |
| <i>Mitrella lunata</i>          |  | 0.86          | -0.23 | 0.67           | 0.13  |
| <i>Mytilus edulis</i>           |  | 0.67          | 0.02  | 0.57           | -0.62 |
| <i>Nereis succinea</i>          |  | 0.37          | 0.38  | 0.71           | -0.39 |
| <b>(b) Infauna</b>              |  |               |       |                |       |
|                                 |  | Sampling time |       |                |       |
|                                 |  | June 1986     |       | August 1986    |       |
| Factor number                   |  | 1             | 2     | 1              | 2     |
| % variation explained by factor |  | 32            | 21    | 22             | 21    |
| <i>Gemma gemma</i>              |  | -0.33         | 0.47  | 0.41           | -0.68 |
| <i>Heteroclitus</i> sp.         |  | 0.43          | -0.53 | 0.66           | -0.19 |
| <i>Mytilus edulis</i>           |  | 0.71          | -0.30 | -0.38          | 0.45  |
| <i>Nereis succinea</i>          |  | 0.85          | 0.24  | -0.53          | -0.15 |
| <i>Petricola pholadiformis</i>  |  | 0.66          | 0.41  | -0.45          | -0.19 |
| <i>Spio</i> sp.                 |  | -0.46         | 0.42  | 0.65           | 0.17  |
| <i>Tellina agilis</i>           |  | 0.15          | 0.79  | 0.38           | 0.75  |

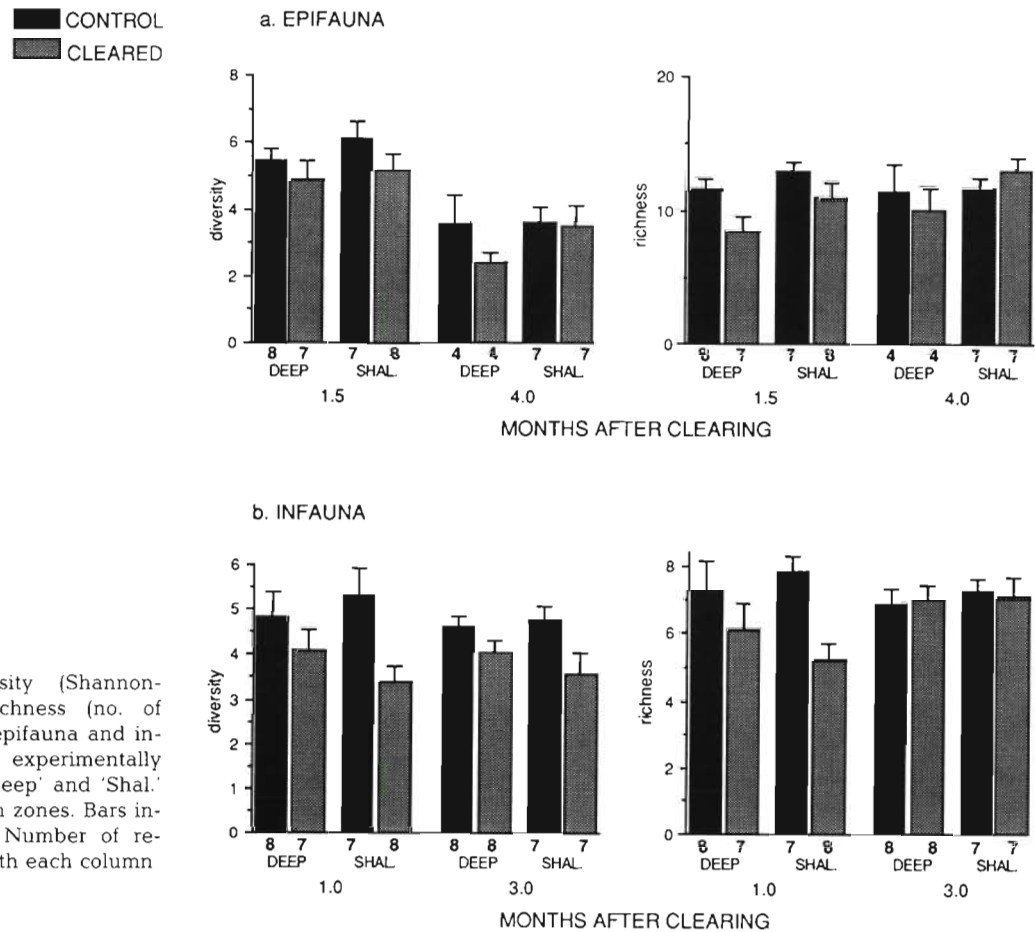


Fig. 7. Average diversity (Shannon-Wiener Index) and richness (no. of species) per sample of epifauna and infauna on control and experimentally manipulated patches. 'Deep' and 'Shal.' (shallow) are the 2 depth zones. Bars indicate standard errors. Number of replicates is shown beneath each column

Table 8. *F*-values of ANOVA comparing diversity and richness on control and experimentally cleared patches. The factors were: 'Time' = time of sampling; 'Depth' = position in shallow or deep zone; and 'Trt' = treatment, ie. experimentally cleared vs control. \*\*  $p < 0.01$ ; \*  $0.01 < p < 0.05$

| Variable                    | Time   | Depth | Trt    | Time × Depth | Time × Trt | Depth × Trt | Time × Depth × Trt |
|-----------------------------|--------|-------|--------|--------------|------------|-------------|--------------------|
| <b>(a) Epifauna</b>         |        |       |        |              |            |             |                    |
| Degrees of freedom          | 1      | 1     | 1      | 1            | 1          | 1           | 1                  |
| Diversity                   | 35.7** | 1.9   | 3.6    | 0.0          | 0.0        | 0.1         | 1.0                |
| Richness                    | 0.4    | 4.4*  | 2.7    | 0.1          | 2.3        | 1.6         | 0.2                |
| <b>(b) Infauna (n = 60)</b> |        |       |        |              |            |             |                    |
| Degrees of freedom          | 1      | 1     | 1      | 1            | 1          | 1           | 1                  |
| Diversity                   | 0.3    | 0.2   | 14.0** | 0.0          | 0.4        | 2.3         | 0.3                |
| Richness                    | 1.2    | 0.0   | 5.2*   | 0.3          | 5.1*       | 1.2         | 0.6                |

eelgrass bed, in contrast to hard-bottom communities (e.g. Dethier 1984, Sousa 1985, Turner 1985) where colonists are often different from the neighbouring species. On the cleared patches, total epifaunal numbers, and abundances of each of the individual species

examined, were found to be significantly related to either one of the 2 major algal groups which quickly returned to the cleared patches. In contrast, no positive relation to eelgrass biomass on the cleared patches was observed.

Previous work has shown that epifaunal-macrophyte associations in this seagrass bed varied in intensity and the type of macrophyte favored (Schneider & Mann 1991a). The recovery of specific invertebrate species following disturbance was dependent on the recovery of suitable macrophyte species. For example, the amphipod *Gammarus* was strongly associated with coarsely branched algae in the undisturbed eelgrass bed (Schneider & Mann 1991a), explaining its rapid reappearance on experimentally cleared areas in conjunction with the influx of coarsely branched algae. In contrast, the tube-dwelling amphipod *Ampithoe* was associated with eelgrass or, when seasonally abundant, *Sphaerotrichia* (Schneider & Mann 1991a). It recovered to control numbers on the cleared patches in September, 1986, because of the pronounced *Sphaerotrichia* bloom. Therefore, assessment of faunal responses to disturbance-induced changes in vegetation cover must incorporate not only the possibility of differential recovery by various macrophyte types, but also species-specific responses of invertebrates to macrophytes.

The more varied infaunal response to clearing reflects the absence of a direct relationship to vegetation as seen for the epifauna. The clam *Gemma gemma*, the one species which became more abundant in cleared patches, is characteristic of open areas and unstable sediments (Saunders et al. 1962), and thus appeared to be responding to the provision of suitable habitat on the cleared patches. Numbers of the remaining infaunal species were either unaffected or depressed 1.0 mo after clearing, but in any case had returned to control levels by the summer's end. The removal of eelgrass was expected to have a major impact because its root-rhizome mat has been previously demonstrated to stabilize the sediment and protect the infauna from predators (Peterson 1982). Casual observation indicated that the removal of the root-rhizome mat on such a small scale did not noticeably affect sediment stability. In areas where large predatory crabs occur, the root-rhizome mat is essential for the protection of many infaunal species (Blundon & Kennedy 1982), but no such species occur in this system (Schneider & Mann 1991b).

The experimental manipulations performed in this study demonstrated that the creation of small artificial gaps had a relatively short-lived effect in the depth zones investigated. Experimental clearing by raking accurately mimicked the total removal of above- and below-ground plant parts observed on naturally ice rafted patches. Two important differences between experimental clearing and natural ice scour are the season of clearing and the greater variation in the size and position of naturally cleared patches. Natural ice scour occurs during winter, while patches were experi-

mentally cleared in early spring. Sediment erosion may be greater following natural ice scour during the winter than experimental clearing in spring, but casual observation did not reveal major erosion on either patch type. The difference in timing of natural and artificial patch creation probably did not have a direct impact on animal recolonization because animals in this eelgrass bed are quiescent during the winter and become active and reproduce during late spring and summer, numbers of recruits being highest in late summer and early fall (Schneider & Mann 1991a).

Some natural ice-scoured patches were much larger than experimentally cleared ones (which represented the average dimensions of natural patches). Although not documented, it was noticed that the size ranges of animals colonizing the bare patches were about the same as those in the control samples. Migration of adults from surrounding vegetation thus appeared to be an important source of colonists and it has elsewhere been demonstrated that epifauna, even sedentary forms such as tubicolous amphipods, undertake frequent migrations (De Witt 1987) and many infaunal species including clams are also mobile (Sortin 1989). If animals move only short distances at one time, then recolonization by immigration of the central portions of larger ice-scoured areas would be slower than that observed for the small experimentally cleared patches.

In the natural seagrass bed, disturbance by ice rafting occurs in the shallow part of the experimental area but not in the deep part. Instances where cleared patches are more similar to control samples from the shallow than the deep zone could reflect the effects of natural ice scour in the shallow zone. *Gemma gemma* was more abundant in control samples from the shallow than the deep zone, and markedly more numerous on experimentally cleared patches. The absence of more widespread similarities between shallow control and experimentally cleared samples might have been because none of the control samples from the shallow zone coincided with patches which had been naturally cleared in the preceding winter. This suggests that natural ice scour does not have a long term impact on the faunal assemblage, a view supported by the observation that the faunal assemblage on the experimentally cleared patches was virtually indistinguishable from control samples by the summer's end.

In conclusion, the recovery of epifaunal and infaunal invertebrates after disturbance was more rapid than expected from eelgrass regrowth. The rapid re-establishment of macroalgae provided a suitable habitat for the epifauna. The removal of eelgrass on such a small scale does not appear to have a long-term impact on the infauna in this eelgrass community.



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