Habitat selection and adult-larvae interactions in settling larvae of soft-shell clam *Mya arenaria*

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**ABSTRACT:** Like many benthic taxa, the soft-shell clam *Mya arenaria* L. is patchily distributed in nature. Near Halifax, Canada (Eastern Passage), *M. arenaria* and several other species (*Polydora cornuta*, *Pygospio elegans*, *Gemma gemma*, *Hydrobia sp.*) occur in higher densities at a sheltered site than at an exposed site ~300 m away with similar grain size but different sediment organic texture. Total faunal densities at the exposed site were comparatively low. To evaluate whether larval settlement plays a role in establishing *M. arenaria* patterns, laboratory flume experiments were conducted with competent *M. arenaria* larvae. Natural cores from the sheltered and exposed sites with resident infauna intact, as well as cores from the same sites that had been defaunated by freezing, were inserted flush with the flume bottom. Highest settlement was observed in faunated cores from the sheltered site where *M. arenaria* are more common. Significantly lower settlement was observed in other treatments, including defaunated cores from the sheltered site. For corresponding treatments, settlement from the exposed site was less than that at the sheltered site. Of the abundant taxa in intact flume cores, *Gemma gemma* densities were a significant, albeit weak predictor of *M. arenaria* settlement. We propose that *G. gemma* influence interface sediment characteristics which, in turn, result in differential larval response. Settlement patterns contradict a previous study (Emerson & Grant 1991; *Limnol Oceanogr* 36:1288–1300) that reported an absence of recently settled *M. arenaria* spat at the sheltered site, despite high numbers at the nearby exposed site. The contrast in results likely reflects our focus on settlement and their focus on spat abundance, and these results considered in tandem suggest that both pre- and post-settlement processes likely determine soft-shell clam distributions at Eastern Passage.

**KEY WORDS:** *Mya arenaria* - Clam - Intertidal - Settlement - Flume - Larvae - *Polydora cornuta* - *Gemma gemma* - *Pygospio elegans*

**INTRODUCTION**

One of the most basic problems in benthic ecology is understanding why variability in spatial and temporal patterns exists in soft-sediment communities (Butman 1987, Olafsson et al. 1994, Snelgrove & Butman 1994). One explanation for this variability is that active choice for a specific sedimentary environment, in the form of habitat selection by settling larvae, may increase the likelihood of placing offspring into that habitat, although differences in larval supply through passive transport (Butman 1987) as well as post-settlement processes (Wilson 1991) offer equally viable hypotheses.

may also be important. Post-settlement predation has also been demonstrated to be a major structuring force in soft-sediment environments (Peterson 1979, 1982).

Flume experiments suggest that species with some degree of habitat specificity as adults have larvae that preferentially settle in defaunated sediments typical of their adult environment (Butman et al. 1988, Butman & Grassle 1992, Grassle et al. 1992a,b, Sneltgrove et al. 1993, 1998). These experiments have deliberately utilised defaunated sediments to test how specific variables such as organic content (e.g. Butman et al. 1988, Butman & Grassle 1992, Grassle et al. 1992a,b), nearbed flow (Jonsson et al. 1991, Sneltgrove et al. 1993), individual species interactions (Andre et al. 1993), and species-specific compounds (e.g. Woodin et al. 1993) may impact initial larval settlement. Thus, a logical progression in the use of laboratory flumes in understanding marine benthos is to use natural communities in experimental manipulations. Field approaches to this question are often complicated by complex and difficult to control variables such as storms (Emerson & Grant 1991), transient predators (Hall et al. 1991), and variability in larval supply (Luckenbach 1984). Laboratory flume experiments allow reproduction of representative bottom flow but under the controlled conditions of a laboratory setting. Near-bottom flow is relevant to many benthic processes (Nowell & Jumars 1984), including larval settlement.

The soft-shell clam *Mya arenaria* L. is broadly distributed throughout shallow northern boreal waters. Populations tend to be highest in muddy sand areas (Gosner 1978, Günther 1992) and can often be patchy on multiple scales. One such example is Eastern Passage, Nova Scotia, Canada, where population dynamics of *M. arenaria* differ markedly between a sheltered and an exposed intertidal site located only hundreds of meters apart. Prior work indicates that despite the differences in wave exposure, the bulk grain size (~250 μm) and silt-clay content (~0.5%) of sediments are similar and several species occur at both sites (Emerson & Grant 1991). Nonetheless, the sites differ in several respects. Faunal densities at the exposed site are comparatively low, and the most abundant species at the sites, including *M. arenaria*, occur in higher densities at the sheltered site. *Gemma gemma*, *Hydrobia* sp., and * Arenicola marina* occur only at the sheltered site and *Spisula* sp. occurs only at the exposed site (Emerson & Grant 1991). In addition, an organic mineral aggregate fluff layer is observed only at the sheltered site, along with a reduced carbon:nitrogen ratio suggestive of more abundant microalgae (Grant & Emerson 1994). *M. arenaria* spat densities and transport have also been found to be substantially higher at the exposed site during some times of the year (Emerson & Grant 1991).

The goal of the present study is to use laboratory flume simulations to determine how differences in sediment composition and natural infaunal communities may influence habitat selection by settling *Mya arenaria* larvae and thus contribute to *M. arenaria* population dynamics.

**MATERIALS AND METHODS**

**Eastern Passage fauna and flume sediment treatments.** Eastern Passage is located in outer Halifax Harbour, Nova Scotia (Fig. 1), and the Sheltered Site and Exposed Site that are the focus of this study are located approximately 300 m apart. In order to compare faunas at the 2 sites, 6.5 cm diameter cores were pushed to 10 cm depth during collections on July 19, 1996 (4 replicates site−1). The choice of core size represented a compromise between being sufficiently small to allow enumeration of the most abundant taxa, and being sufficiently large to sample bigger organisms. Undoubtedly, larger individuals of *Mya arenaria*, * Arenicola marina* and *Spisula* sp. were not effectively enumerated by this sampling. Spot checks of cores collected on August 4 and August 12, 1996, when the second and third sets of flume experiments were conducted, indicated no obvious changes in fauna densities at the exposed site during that time which might have confounded interpretation of settlement patterns in the flume. Cores were preserved in 10% buffered formalin and transferred to 80% ethanol within 2 wk of sampling. Samples were processed over a 106 μm sieve to ensure that recently settled macrofauna would also be effectively enumerated. In most cases, fauna could be identified to species.

Sediments used in the flume experiments were collected from the Sheltered and Exposed Sites, and included cores that were either left intact (hereafter referred to as ‘Infauna’ cores) or defaunated by freezing (hereafter referred to as ‘No Fauna’ cores). For No Fauna cores, bulk sediments were collected from both sites in July, 1996, at low tide by scraping the upper 2 to 3 cm of sediments into large plastic storage containers, rinsing several times with fresh water and allowing fine particles to settle before decanting. Large shell fragments and dead organisms were removed. Sediments were frozen until the day prior to experiments and sediment from the same collection was used in all flume experiments. On the day of an experiment, No Fauna sediments were stirred vigorously and the sediments allowed to settle before coring with 4.36 cm diameter cores.

Preparation of Infauna cores was delayed until an experiment was imminent. For a given set of experi-
ments, six 11.4 cm diameter infaunal cores were collected from the Exposed and Sheltered Sites at Eastern Passage and transported to Dalhousie, Halifax, Nova Scotia, in coolers with ice packs. At Dalhousie's aquatron facility, cores were maintained in filtered, running seawater until used in experiments within the next 72 h. Infauna cores from each of the 2 sites were subsampled immediately prior to a flume run by pushing a 4.36 cm diameter core into the larger core and pulling away the larger core making sure that seawater was maintained above the sediment and that the sediment surface was well below the top of the smaller core. An O-ring sealed plug was then inserted in the bottom of the core to prevent sediment loss. The purpose of the smaller cores was 2-fold. First, edge effects associated with the outer periphery of the larger cores would be eliminated; we reasoned that the periphery of cores would have the most anomalous biology and geochemistry and that these effects would be exacerbated during storage of cores between flume runs; recoring would introduce a new edge effect but of lesser magnitude. The smaller cores also allowed cross-stream placement of 4 cores with minimal flume wall effects (see below); larger cores would have necessitated fewer treatments. All sediments and cores were collected between mid-July and mid-August, and were therefore chosen to correspond approximately to annual peak *Mya arenaria* settlement at Eastern Passage as documented by Emerson & Grant (1991).

**Conditioning adults.** Eastern Passage, though not obviously polluted, is sufficiently close to Halifax Harbour that a more pristine site was felt to be more suitable for obtaining adults for conditioning, and in April, 1996, ~60 adult *Mya arenaria* (individuals 5 to 7 cm in length) were collected from Lawrencetown Beach, a site ~12 km northeast of Eastern Passage.

Adults were maintained in running seawater at approximately 17°C. During a 2 h period each day flow was shut off and concentrated algae from Coast Oyster Company and from live laboratory cultures (*Isochrysis* sp., *Thalassiosira pseudonana* and/or *Chaetoceros muelleri*) were downwelled through the sediments where adults were maintained (initial concentration of 100,000 cells ml$^{-1}$ in 1000 l). *Adults were conditioned* for ~3 mo and spawned by placing at least 15 individuals overnight in filtered, aerated seawater warmed to ~22°C. The following morning, larvae were collected on a 20 µm sieve, distributed into 20 l containers and maintained at concentrations of less than 10 ind. ml$^{-1}$. Larvae were fed a mixture of *Isochrysis* sp. and *T. pseudonana* at a combined concentration of ~50,000 cells ml$^{-1}$, and seawater was changed every second day during the 10 to 13 d it took between fertilization and a given experiment.

**Competency.** Timing of experiments was reliant on a batch (i.e. individuals from a single spawning) of cultured larvae being physiologically competent to settle. The primary criterion for evaluating larval competency was observation of several hundred *individuals* from cultures to see whether pediveligers were present. Once ~5% of individuals appeared to have attained a pediveliger stage, a series of flume experiments was initiated within the next 12 h. In addition, once pediveligers were present, still-water competency tests were conducted concurrently with flume experiments. Competency tests were similar to those described else-
where for bivalves (e.g. Grassle et al. 1992a, Snelgrove et al. 1993, 1998) and are not reported here except to note that they supported the assumption that at least some of the larvae in a given batch were competent during all experiments.

The flume. All flume experiments were conducted in the recirculating flume at Dalhousie described in Roegner et al. (1995). Briefly, the flume consists of a 7.32 m long, 50 cm wide, and 50 cm deep raceway with a 37 cm diameter return pipe that runs beneath the flume and an impeller that regulates flow speed via a variable speed DC motor. In the test section of the flume (529 cm downstream of the entrance) 4 holes were cut in the flume floor to allow insertion of cores (inner diameter = 4.36 cm) containing the sediment treatments. Cores were distributed across the flume, perpendicular to the raceway, and confined to the central 25 cm in the cross-stream direction. This placement reduced the likelihood of wall effects associated with treatments located too close to the flume walls (see Nowell & Jumars 1987). Cores were inserted flush with the flume floor and held in place with nylon collars fitted with internal O-rings secured to the underside of the flume that created a watertight seal.

Experimental protocol. After cores with live infauna had been collected in the field, the return pipe in the flume was filled with seawater pumped in from a comparatively clean area of Halifax Harbour and filtered to 5 μm. Water was added to the pipe until no more water could be added without flooding the flume raceway. Each core containing 1 of the 4 treatments was inserted from the underside of the flume until the rim of the plastic core barrel was flush with the flume bottom. Additional filtered seawater was then added to the flume until the water depth in the runway was 10 cm. Sediment in the cores was raised until flush with the flume bottom using a modified turnbuckle to push up the O-ring fitted plastic plug and the sediment above it. Flow was then turned on to achieve a $u_0$ of $0.7 \text{ cm s}^{-1}$, as calculated earlier from velocity profiles collected with a hot wire anemometer (DanTech) using the log-profile technique (e.g. Jumars & Nowell 1984). Flow was characterized separately from runs with larvae present to avoid any interaction between the hot wire probe and larvae; flows were reproduced for runs with larvae by duplicating water depth and pump speed. This shear velocity is within the typical range observed at Eastern Passage (Emerson & Grant 1991), and although higher shear velocities do occur (particularly in the fall), we deliberately chose a flow regime below the critical shear velocity of the sediment ($1.0 \text{ cm s}^{-1}$, Emerson & Grant 1991). In addition to the complication that would arise if sediments were eroding during experiments, the semi-diurnal tides over the sandflats are insufficient in themselves to move sediments and require wind input greater than is typical for the site in July and August (Emerson 1991). The impeller rotation rate required to achieve this flow speed was surprisingly slow, so that damage to swimming larvae was expected to be negligible. Approximately 40000 larvae were added at the upstream end of the flume by pouring them from a beaker that was moved slowly back and forth in the cross-stream direction. The windowless room was lit by fluorescent bulbs well above the flume; during flume experiments lights were kept on at all times, whereas during culturing they were maintained on a 12 h light-dark cycle.

Each experimental run lasted 12 h, after which a 100 μm mesh net that fit tightly across the full width and depth of the flume was placed in the flume for 5 min to filter out any remaining suspended larvae. Once filtering was completed, the flow was turned off, weighted caps were placed over the sediment cores, and the flume was drained down and sponged until the raceway was dry. Sediments were then carefully removed from the top 2 cm of each core and preserved in 10% buffered formalin; within 2 wk samples were transferred to 80% ethanol and Rose Bengal for later enumeration of settled Mya arenaria and all macrofauna. Core barrels were then removed from the flume, rinsed, and fresh cores were prepared and inserted into the flume as described above, except that cross-stream positions were shifted. A new aliquot of 40000 larvae from the same larval batch was then added and a second experimental run initiated. This procedure was repeated 2 more times so that each treatment occupied a different position in each of the 4 successive runs for a given larval batch in a Latin squares design. The same protocol was repeated with 3 different batches of larvae for a total of 12 experimental runs (3 batches × 4 runs batch$^{-1}$).

Data analysis. Flume settlement data were analyzed in a single mixed model nested analysis of variance (ANOVA) in which $y = \text{constant} + \text{Batch} + \text{Column} + \text{Treatment} + \text{Run(Batch)} + \text{Treatment} \times \text{Batch} + \text{Column} \times \text{Batch} + \text{error}$; $y$ refers to number of settled Mya arenaria larvae, Batch refers to larval batch, Column refers to cross-stream position in the flume, Treatment refers to the sediment treatments described above, and Run refers to sequential experiments conducted with larvae of the same batch. M. arenaria densities were $\log(x + 1)$ transformed to homogenize variances. Not all interaction terms could be tested in a single analysis because of insufficient degrees of freedom, and only those considered to be most important were included.

In order to determine whether resident infauna impacted larval settlement, linear regression analysis was performed separately for the 7 taxa that were abundant in Sheltered Infauna cores. Multiple regres-
sion was not used because of problems with co-linearity of different taxa. Thus, the general model was \( y = m \) (infaunal density) + constant, where \( y \) refers to number of settled Mya arenaria larvae, \( m \) refers to slope, constant refers to the \( y \) intercept and infaunal density refers to densities of Gemma gemma, Hydrobia sp., Pygospio elegans, Polydora cornuta, Oligochaeta spp., or adult M. arenaria in Sheltered Faunated cores. Other core treatments were not pooled for regression analysis because of the significant Treatment effect observed in the analysis described above (see ‘Discussion’), and the complete absence of infauna in No Fauna cores. Densities of infauna in Exposed Infauna cores were so low that regression analysis for this treatment would not have been meaningful.

**RESULTS**

Cores collected from the Sheltered and Exposed Sites revealed that, although bulk sediment characteristics are similar, the sediment structure at the sediment surface was quite different (Grant & Emerson 1994). The Sheltered Site had a fluffy looking sediment that clearly had very fine material concentrated at the sediment-water interface and many aggregates attached to grains. At the Exposed Site, a fluffy grain layer was not apparent, and there was little evidence of biological activity. In addition to differences in flow regime, faunal cores indicated substantial faunal differences between the 2 sites that may contribute to the observed differences in biological structure. The polychaetes Polydora cornuta, Pygospio elegans, and Capitella spp., Oligochaeta spp., and the molluscs Gemma gemma, Hydrobia sp., and Mya arenaria were all more abundant at the Sheltered Site; total faunal densities at the Exposed Site were very low in comparison with those at the Sheltered Site (Fig. 2). All of the dominant taxa at the Exposed Site were also present at the Sheltered Site but in very low abundance. In addition to adults, a few small individuals of M. arenaria were noted in some cores, but all were sufficiently large that they could not have been confused with M. arenaria larvae that settled during flume experiments.

Competency tests for Mya arenaria larvae produced less than definitive results in terms of proportion of competent larvae and selectivity. However, at least a few individuals settled and metamorphosed during each of the competency tests, indicating that some portion of the culture was competent to settle during all experimental runs.

Conditions for all flume experiments are summarized in Table 1. Temperature was maintained within a fairly narrow range of 16.2 to 19.8°C. Larval age for flume experiments ranged from 11 to 13 d, depending on specific Run and Batch. Core age varied from 2 to 62 h. Neither larval age nor core age had any consistent effect on patterns of settlement, in that Sheltered Infauna cores had highest settlement in all 12 experiments. Responses to other treatments appeared to vary more with different batches than with core age or larval age. Estimated recoveries in experimental runs ranged from 2.2 to 22.5% of the ~40,000 larvae added.

ANOVA of flume settlement experiments indicated a significant Treatment effect as well as a significant Run (nested in Batch) effect (Table 2) with the general result that sheltered cores had higher settlement. A Tukey’s multiple comparisons test indicated that significantly higher settlement occurred in Sheltered Infauna cores compared to all other treatments and that Sheltered No Fauna cores had significantly higher settlement than Exposed No Fauna cores. No other significant differences were observed. The significant Run effect can be attributed to changes in total settlement that occurred over different runs of a given Batch (Fig. 3). In the first set of flume experiments, total settlement increased as larvae became older. In the second and third experiments, settlement peaked during the second run (Table 1). Thus, there is no consistent trend regarding core or larval age and peak settlement.

![Fig. 2. Dominant macrofauna at Eastern Passage study sites. Bars denote means and lines denote 1 SE (n = 4). Note that different axes are used for the 2 sites](image-url)
Table 1. Experimental conditions during each of the flume runs

<table>
<thead>
<tr>
<th>Batch</th>
<th>Expt run</th>
<th>Start date</th>
<th>Duration (h)</th>
<th>Larval age (d)</th>
<th>Core age (h)</th>
<th>Water temp. (°C)</th>
<th>Approx. number of larvae added</th>
<th>Percent larvae recovered</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>A</td>
<td>20 Jul 1996</td>
<td>12</td>
<td>11</td>
<td>18</td>
<td>16.2</td>
<td>40000</td>
<td>2.8</td>
</tr>
<tr>
<td></td>
<td>B</td>
<td>23 Jul 1996</td>
<td>12</td>
<td>12</td>
<td>34</td>
<td>16.8</td>
<td>40000</td>
<td>11.7</td>
</tr>
<tr>
<td></td>
<td>C</td>
<td>23 Jul 1996</td>
<td>12</td>
<td>12.5</td>
<td>48</td>
<td>17.0</td>
<td>40000</td>
<td>12.4</td>
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<tr>
<td></td>
<td>D</td>
<td>24 Jul 1996</td>
<td>12</td>
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<td>13.1</td>
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<td>2</td>
<td>A</td>
<td>05 Aug 1996</td>
<td>12</td>
<td>11</td>
<td>12</td>
<td>17.9</td>
<td>40000</td>
<td>2.2</td>
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<td></td>
<td>B</td>
<td>06 Aug 1996</td>
<td>12</td>
<td>11.5</td>
<td>26</td>
<td>18.6</td>
<td>40000</td>
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<td>40000</td>
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<td>12</td>
<td>12.5</td>
<td>48</td>
<td>18.9</td>
<td>40000</td>
<td>9.3</td>
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Table 2. (a) ANOVA results with all batches included. Data were log(x+1) transformed to homogenize variances. Significant results are shown in bold. (b) Means were compared with a Tukey's multiple comparisons test. Lines connect means that were not significantly different.

(a) ANOVA

<table>
<thead>
<tr>
<th>Effect</th>
<th>df</th>
<th>SS</th>
<th>F</th>
<th>p</th>
<th>r²</th>
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<td>Treatment</td>
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<td>63.12</td>
<td>30 100</td>
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<td>Column</td>
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<td>1.651</td>
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<td>Batch</td>
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<td>Treatment x Batch</td>
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<td>4.196</td>
<td>2.259</td>
<td>0.064</td>
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</tr>
<tr>
<td>Column x Batch</td>
<td>6</td>
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<td>0.715</td>
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<tr>
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<td>21.642</td>
<td>7.766</td>
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<tr>
<td>Error</td>
<td>18</td>
<td>5.573</td>
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</table>

(b) Tukey's multiple comparisons test

<table>
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<tr>
<th>Treatment</th>
<th>Sheltered Infauna</th>
<th>Sheltered No Fauna</th>
<th>Exposed Infauna</th>
<th>Exposed No Fauna</th>
<th>Mean</th>
</tr>
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<tr>
<td></td>
<td>3577.2</td>
<td>653.2</td>
<td>396.6</td>
<td>182.9</td>
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</table>

Table 3. Regression analyses of dominant taxa from Sheltered Infauna cores as predictors of Mya arenaria settlement. Because a significant relationship was observed only for Gemma gemma, slopes and intercepts are not reported for other taxa. The regression equation for G. gemma is given in Fig. 4.

<table>
<thead>
<tr>
<th>Taxon</th>
<th>n</th>
<th>SS regression</th>
<th>SS residual</th>
<th>F</th>
<th>p</th>
<th>r²</th>
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<tbody>
<tr>
<td>Polydora cornuta</td>
<td>12</td>
<td>2082039.66</td>
<td>4.68272x10⁷</td>
<td>0.45</td>
<td>0.52</td>
<td>0.0</td>
</tr>
<tr>
<td>Pygospio elegans</td>
<td>12</td>
<td>199709.64</td>
<td>4.87095x10⁷</td>
<td>0.04</td>
<td>0.84</td>
<td>0.0</td>
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<tr>
<td>Oligochaeta spp.</td>
<td>12</td>
<td>6296138.69</td>
<td>4.26131x10⁴</td>
<td>1.48</td>
<td>0.25</td>
<td>0.04</td>
</tr>
<tr>
<td>Capitella spp.</td>
<td>12</td>
<td>1339.827</td>
<td>4.89079x10⁷</td>
<td>0.00</td>
<td>0.99</td>
<td>0.00</td>
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<tr>
<td>Hydrobia sp.</td>
<td>12</td>
<td>1.10609x10⁷</td>
<td>3.78483x10⁷</td>
<td>2.92</td>
<td>0.12</td>
<td>0.15</td>
</tr>
<tr>
<td>Gemma gemma</td>
<td>12</td>
<td>1.87427x10⁷</td>
<td>3.01665x10⁴</td>
<td>6.21</td>
<td>0.03</td>
<td>0.32</td>
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<tr>
<td>Mya arenaria</td>
<td>12</td>
<td>915003.51</td>
<td>3.97313x10⁷</td>
<td>2.30</td>
<td>0.16</td>
<td>0.11</td>
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</table>

Regression analyses indicated that, of the dominant macrofauna present in the Sheltered Infauna cores, only Gemma gemma was a significant predictor of Mya arenaria settlement (Table 3). G. gemma densities in Sheltered Infauna cores explained a significant amount of variation in M. arenaria settlement ($r^2 = 0.32$), and higher G. gemma densities were associated with higher M. arenaria settlement. Densities of Pygospio elegans, Polydora comuta, Oligochaeta spp., adult M. arenaria, and Hydrobia sp. were not significant predictors of M. arenaria larval settlement.

**DISCUSSION**

The establishment of pattern in soft-sediment communities and the role that larval settlement plays is a largely unresolved debate in benthic ecology (e.g. Olafsson et al. 1994, Snelgrove & Butman 1994). Although there are important variables that can only be tested in situ, field experiments in which conditions are relatively 'natural' suffer from problems of appropriate time scales and stochastic processes that may or may not be known (see Gallagher et al. 1983). Laboratory experiments mimic nature poorly, and there is strong evidence that processes as varied as feeding (Miller et al. 1992), nutrient flux (Huettel & Gust 1992) and larval settlement behaviour (Grasse et al. 1992a, Snelgrove et al. 1998) may change considerably in different flow regimes. As such, laboratory flumes that reproduce representative flow regimes have been instrumental in demonstrating selective capacity of settling larvae.
Previous flume experiments on larval settlement have deliberately used sediments that have been altered in order to test explicit hypotheses on the importance of grain size, organic content and other key variables. The use of azoic sediments in settlement studies has been criticized (Smith & Brumsickle 1989) because they are 'unnatural'. Some studies utilized sediments that were azoic and homogenized (Butman et al. 1988, Grassle & Butman 1989, Butman & Grassle 1992, Grassle et al. 1992a,b, Snelgrove et al., 1993, 1998), others used sediments that were azoic but with specific chemical attributes (Woodin et al. 1993) and some used sediments with a very simplified fauna (André & Rosenberg 1991). These flume experiments suggest that near-bed flow (Snelgrove et al. 1993), active habitat selection for sedimentary characteristics such as high organic content (e.g. Butman et al. 1988, Butman & Grassle 1992, Grassle et al. 1992a,b), coarse grain size/low organic content (Snelgrove et al. 1998) or geochemical cues associated with disturbance (Woodin et al. 1995) and predation by suspension feeders (André et al. 1993) may all contribute to spatial pattern. Although flume experiments with altered sediments have greatly improved our understanding of larval settlement, the next logical application of flume simulations is to use natural communities in manipulations to bridge the gap with field experiments.
Resident infauna can impact settling larvae via many different mechanisms. Larvae could actively respond to or avoid certain species or associated compounds (e.g. Woodin et al. 1993), fall prey to resident fauna (e.g. Woodin 1976) or fail to compete effectively with other species (e.g. Whitlatch 1980). Structures that extend above the bottom also disturb boundary-layer flow (Eckman 1985, Erman & Jumars 1988), as can currents generated by suspension feeders (Monismith et al. 1990); these disturbances could influence settlement patterns of larvae that are delivered to the bed as passive particles (Armonies & Hellwig-Armonies 1992, Butman & Grasse 1992, Snelgrove et al. 1993).

Despite similarities in sediment composition (Emerson & Grant 1991) the fine-scale surface structure and faunas at the Sheltered and Exposed Sites are remarkably different and present interesting questions regarding larval settlement. As is the case with the other dominant taxa at Eastern Passage, *Mya arenaria* occur at both sites but are more abundant at the Sheltered Site during the summer months when we sampled (Fig. 2). Differences in larval settlement in response to sediment structure or species present could contribute to density differences. The high densities of interface feeders such as *Polydora cornuta* and *Pygospio elegans* probably contribute to the concentration of fine material at the sediment-water interface and the distinctly different appearance of the sediment. An additional factor that may contribute to differences in this ‘fluff layer’ is a sticky microbial film associated with sediment grains at the Sheltered Site that is lacking at the Exposed Site, likely relating to hydrodynamic activity and abrasion (Grant & Emerson 1994). Alternatively, the species that contribute to these habitat differences could impact *M. arenaria* settlement.

Given that *Mya arenaria* is more abundant at the Sheltered Site than the Exposed Site during summer, higher settlement should be expected in Sheltered Site cores in flume simulations if habitat selection contributes to field distributions. Indeed, larvae settled in significantly higher numbers in Sheltered Site cores than in corresponding Exposed Site cores, suggesting that settling *M. arenaria* are attracted to some sediment attribute that is only partly destroyed by defaunation/freezing. Although Sheltered No Fauna cores had significantly higher settlement than Exposed No Fauna cores, numbers were far less than in Sheltered Infauna cores. Explanations for this result include active choice by settling larvae for some aspect of the sediment that was greatly reduced by freezing or by removal of adults, differences in sediment biofilming ('stickiness') and larval retention, differences in sediment roughness elements, and thus in passive larval flux to the bed, and increased larval flux to the bed via infaunal feeding activity.

Differences in sediment biofilm may have been a factor, given that biofilming is greater in Sheltered Site sediments than Exposed Site sediments (Grant & Emerson 1994) and sieving and freezing would likely have altered any biofilm present. Interestingly, Ólafs-
son (1988) observed higher settlement of *Mya arenaria* in substratum without red algal mat cover compared to natural or defaunated substratum with mats; combined with our results, this finding suggests that any linkage between settling *M. arenaria* and algae/biofilming differs substantially depending on the specific nature of the organic matter associated with the sediment. Roughness is less likely as an explanation for our treatment differences because the No Infauna cores were not obviously different from Infauna cores with respect to this variable. In addition, none of the core surfaces was perfectly flat, and larger-scale topography that was randomly distributed among treatments would likely have masked any roughness effect induced by tubes etc. If differences in settlement are related to increased larval flux via infaunal feeding currents, or active selection (or avoidance) by larvae for the presence of some species, then the densities of that species should be a significant predictor of *M. arenaria* settlement.

Regression analysis to determine which individual taxa were significant predictors of *Mya arenaria* settlement was significant only for *Gemma gemma*. *G. gemma* is a suspension-feeding bivalve that has been observed to enhance *Mercenaria mercenaria* settlement in still-water experiments (Ahn et al. 1993). The authors concluded that sediment alteration by *G. gemma* resulted in active choice by settling *M. mercenaria* in response to multiple cues including sediment geochemistry, roughness, and biodeposits. They eliminated entrapment related to stickiness because individuals were crawling freely. In the present study, if *G. gemma* did modify sediment in a way that made it more attractive to *M. arenaria* larvae, then freezing destroyed much of whatever made the sediment attractive. *G. gemma* may help maintain an attractive biofilm layer on sediments (that would be reduced by freezing), either by reducing bioturbation by other species or perhaps by enhancing nutrient flux within the sediment (Weinberg & Whitlatch 1983 provide an example of a polychaete enhancing nutrient flux and thereby enhancing *G. gemma* growth). Species-specific compounds (Woodin et al. 1993) that may largely break down in freezing also cannot be dismissed. Given the small size of *G. gemma* siphons, it is unlikely that their feeding currents increased larval flux to the bed. In any case, the weakness of the relationship indicates that whatever the effect of *G. gemma*, it is probably indirect or possibly even spurious. Interestingly, the finding of enhanced *Mya arenaria* settlement in the presence of *G. gemma* is in direct contradiction to previous predictions (Sanders et al. 1962, Emerson & Grant 1991), based in the former case on an observed inverse relationship between *G. gemma* and *M. arenaria* abundance in the field. Clearly a more complex explanation is needed to clarify these field distributions.

Competency tests indicated highest settlement approximately 24 h after (presumably competent) pediveligers were observed in the culture. It is interesting, though perhaps coincidental, that settlement in flume experiments tended to peak around this time. Although competency tests offer a crude measure of whether larvae are competent, the degree of variability between replicates and batches punctuates the need for a more definitive and repeatable means of evaluating competency. The absence of high levels of spontaneous settlement in cultures in the later runs of flume experiments suggests that competency in individuals probably lasts for several days but additional data to explicitly test this question are sorely needed. The significant Run effect resulted from changes in total settlement depending on relative age of larvae used in experiments. Given that the degree of selectivity appeared to be highest in Run 2 of all 3 experiments (Fig. 3), and that it tended to be lowest by the final run, it is unlikely that the first 2 sets of experiments were run prematurely. A more likely scenario is that larvae were reaching a point where swimming capacity was reduced by increasing larval size (see Jonsson et al. 1991) with age, and passive deposition became increasingly important in the later flume runs.

An obvious conclusion that might be drawn from this study is that differences in larval settlement between sites could contribute to higher *Mya arenaria* adult densities at the Sheltered Site. Nonetheless, a more complicated explanation must be invoked. From high frequency sampling throughout the summer months, Emerson & Grant (1991) documented a complete absence of *M. arenaria* spat at the Sheltered Site and extremely high abundances of spat at the Exposed Site. It is quite possible that differences in larval supply to the 2 sites during their study period contributed to the absence of spat at the Sheltered Site; Günther (1992) noted differences in *M. arenaria* settlement over scales of 100s of metres in the Wadden Sea. Another possibility is higher densities of predators at the Sheltered Site may quickly remove settled *M. arenaria*, and successful recruitment to the Sheltered Site may be accomplished by post-settlement transport of larger and less vulnerable individuals. In several instances in our flume experiments, guts of spionid polychaetes were full of *M. arenaria* larvae, a finding that is consistent with previous studies that have documented spionid predation on bivalve larvae (Breese & Phibbs 1972, Daro & Polk 1973). Predation has been invoked to explain *M. arenaria* patterns elsewhere as well (Möller & Rosenberg 1983).

Given that the absence of spat at the Sheltered Site was observed during a different year than the present study, and the strong selective behaviour exhibited by *Mya arenaria* for Sheltered Site cores in the flume
experiments, the most parsimonious explanation for these findings is that habitat selection can play a role in determining *M. arenaria* pattern but it is not the sole determinant. In some years, supply differences may pre-empt the opportunity to select, and in these years recruitment is either limited or accomplished by post-settlement immigration (Emerson & Grant 1991, Roegner et al. 1995). Thus, *M. arenaria* pattern at Eastern Passage is likely determined by a combination of pre- and post-settlement processes.

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