Use of null models to assess patterns of size class migrations in the copepod Zausodes arenicolus

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ABSTRACT: Size-specific patterns in the migration (emergence) of a sediment-associated harpacticoid copepod were examined in a subtidal seagrass habitat using a null model approach. Size frequency distributions of all Zausodes arenicolus Wilson, 1932 and of females, males and copepodites collected on 2 dates in the water column and sediment were analyzed. Migrating and non-migrating size frequency distributions for all Z. arenicolus were different only in November 1983. The migration of significantly larger females in November 1983 influenced the size distribution for all migrating Z. arenicolus and was not the result of random emergence from the sediment. Size distributions of migrating and non-migrating males and copepodites were not significantly different and could not be distinguished from null distributions on either date. Evidence for size influencing migratory behavior of Z. arenicolus is weak, but the non-random migration of larger females on one date requires further investigation.

INTRODUCTION

The size of an individual often influences migratory behavior in marine organisms, especially Crustacea (Sulkin 1984, Epifanio 1988). Typically, the effects of body size on migration are inferred from differences in water column vertical distributions of different sized individuals (Huntley & Brooks 1982, Rothlisberg 1982, Williams & Fragopoulou 1985). Size influences mating success in several arthropod taxa (Ward 1983, Lawrence 1987, Grad & Maly 1988), and increased mating contacts is proposed as one advantage of vertical migration for benthic species (Preece 1971, Williams & Bynum 1972), including harpacticoid copepods (Bell et al. 1988). Size-selective predation also can influence migrations of freshwater insects (Allan 1978) and crustaceans (Newman & Waters 1984, Malmqvist & Sjostrom 1987), and marine zooplankters (Robertson & Howard 1978, Fancett & Kimmerer 1985). Based on observed relationships between body size and migration, a number of deterministic hypotheses, for example differential predation, have been proposed to explain size-related differences in migration (Huntley & Brooks 1982, Ward 1983, Lampert 1987).

An alternative to concluding that size differences in migration result from deterministic factors would be to suggest that observed distributions simply result from non-deterministic or random factors. The role of non-deterministic factors in shaping observed patterns can be examined using a null model. Although the term null model was coined by Colwell & Winkler (1983), equivalent models have been in use for some time (see Williams 1947). While most hypotheses assume a deterministic cause, null model hypotheses investigate the contribution of stochastic factors in shaping observed patterns. Null models can serve as controls for non-experimental data by evaluating the role of chance in structuring patterns (Strong 1980).

Procedures for constructing a null model will vary considerably from study to study. Fundamentally, null models are used to generate randomly a new data set that is compared statistically with data actually collected. The newly constructed data set is generated by Monte Carlo simulations, a series of random draws with or without replacement, from a previously identified source pool that usually consists of a combination of all data collected. Numbers drawn from a source pool in simulations can be classified by variables such as species names, flowering times, or body sizes. Harvey et al. (1983) and Nitecki & Hoffman (1987) contain reviews and examples of the use of null models.

Our study applies a null model approach to analyze behavioral differences in migration among size classes.
of a sediment-associated harpacticoid copepod, *Zausodes arenicolus* Wilson, 1932. This is a very mobile species in subtropical seagrass beds and is commonly found in phytal (Bell et al. 1987), benthic and pelagic (Walters & Bell 1986) habitats. Pelagic occurrences of *Z. arenicolus* in subtidal sand and seagrass habitats typically result from active migration into the water column (Walters & Bell 1986, Walters 1987). *Z. arenicolus* commonly is involved in precocial clasping (Kern et al. 1984), and migration has been linked to reproductive activity (Bell et al. 1988). Preliminary observations of *Z. arenicolus* migrations indicated that behavioral differences among size classes may exist (Walters unpubl.). In this paper, we examine the size frequency distributions of migrating and non-migrating *Z. arenicolus* using a null model approach and ask whether the migratory behavior of this species is influenced by size.

**METHODS**

**Field methods.** *Zausodes arenicolus* that migrated into the water column were collected from a *Thalassia testudinum* (König) seagrass bed near the mouth of Tampa Bay, Florida, USA, on 23 November 1983 and 17 November 1984. Dates were selected to coincide with seasonal increases in *Z. arenicolus* sediment densities and numbers migrating (Walters unpubl.). An emergence trap (Walters 1988) was used to collect migrating harpacticoids. The trap consisted of a 10.4 cm diameter PVC base pushed 15 cm into the sediment, and a clear acrylic catch-tube positioned securely on the PVC base. An inverted, transparent funnel in the base of the catch-tube and a 0.063 mm mesh cap directed movement into and restricted movement out of the catch-tube, respectively. Organisms had to swim 15 cm from the sediment-water interface to be collected in the catch-tube. A cork stoppered the funnel while the catch-tube was filled with 0.063 mm filtered seawater and placed on the PVC base. Sampling periods began when the cork was removed.

On each date, 10 to 12 emergence traps were placed in an ca 2 m² area of the seagrass bed immediately before sunset, since sunset represents the time of maximum harpacticoid migration within Tampa Bay seagrass habitats (Walters 1987, 1988). Water depth during 2 h sampling periods was ca 1 m. Migrating (within catch-tube) and non-migrating (within sediment) harpacticoids were collected at the end of a 2 h sampling period. Non-migrating harpacticoids were estimated from 3 subsamples which consisted of a 2.7 cm diameter core pushed 3 cm into the sediment within the PVC base of each trap. All catch-tube and sediment samples were preserved in 10 % buffered formalin and seawater, with Rose Bengal added to facilitate sorting.

**Laboratory methods.** Non-migrating harpacticoids were separated from sediments by the shake and decant method (Wieser 1960), and all harpacticoids sorted at 25× under a dissecting microscope. Individual *Zausodes arenicolus* were removed from samples, mounted in glycerin and measured with an ocular micrometer under a compound microscope. All individuals were measured from the anterior margin of the cephalothorax (excluding rostrum) to the posterior end of the anal somite (excluding terminal setae). Later stage copepodites and adult copepods were sexed, and observations regarding the number of gravid females and clasping pairs per sample were recorded.

**Statistical analyses.** To compare the overall size frequency distributions of migrating and non-migrating *Zausodes arenicolus* within the seagrass bed, we combined individuals collected in different traps. In 1983, size frequency distributions representing the sum of *Z. arenicolus* from 10 traps were analyzed. Size frequency analyses of 1984 data only used 4 of 12 emergence traps collected because 4 traps adequately represented the size frequency distribution of migrating and non-migrating harpacticoids in 1983; there was no statistical difference between distributions from 4 and > 4 traps [Kolmogorov-Smirnov 2-sample test, (K-S) Z = 1.276, p > 0.05]. We realize population size distributions may change yearly, but believe our method was a reasonable approach to reduce the 1984 sample size without loss of information.

The frequency of non-migrating individuals and the total frequency, migrating and non-migrating, of a given size class were adjusted for area sampled because non-migrating *Zausodes arenicolus* were estimated from 3 subsamples in each trap. Total frequency of a given size class was calculated by:

\[ f_i = m_i + s_i (5.2^2/3(1.35^2)) \]

where \( f_i \) = frequency of size class i (water column + sediment); \( m_i \) = frequency of size class i in the water column; \( s_i \) = frequency of size class i in the sediment; 5.2 = radius of water column trap; 1.35 = radius of sediment core subsample.

*Zausodes arenicolus* were grouped into 20 μm size classes and the size frequency distributions of migrating and non-migrating individuals compared on each date using the K-S 2-sample test at the 0.05 level of significance (Sokal & Rohlf 1981). Analyses also were run separately for males and females because *Z. arenicolus* is sexually dimorphic with respect to size as well as structure. Comparison of null-model-generated size frequency distributions (see below) for migrating *Z. arenicolus* were also compared with observed water column distributions using the K-S 2-sample test.

**Null model techniques.** Null model size distributions of migrating *Zausodes arenicolus* were generated by
Monte Carlo simulation from a source pool produced by summing actual water column and sediment numbers within each size class (Table 1). The null distribution generated from the model consisted of individuals chosen from a size pool weighted by the actual proportion of individuals in each size class. Distributions generated by the null model were compared to actual distributions of migrants. The null model hypothesis tests whether observed distributions of migrating *Z. arenicolus* could have been generated by random factors; i.e., the numbers of a given size class in the water column simply reflected the proportion of that size class in the sediment. In summing together water column and sediment copepods to produce the source pool, the model assumes all migrators came from the sediment population, and that the source pool is representative of all available migrating size classes and relative abundances of *Z. arenicolus* in each size class. Given the haphazard positioning of traps in the seagrass bed and the large sample size of measured *Z. arenicolus* (3511 in 1983 and 1032 in 1984), assumptions about sampling all size classes and accurately determining the relative abundance of each size class appear to have been satisfied.

Monte Carlo simulations were written using SPSS® and run on the IBM 3033 at the University of South Florida Regional Computer Center and the IBM 3081 at the University of South Carolina. In each simulation, a different random number sequence was used to ensure independence of the results. Individuals were sampled from each source pool without replacement; once an individual copepod migrated into the water, sediment re-entry through the catch-tube funnel was unlikely. Numbers drawn from the source pool for each model were determined by the actual numbers of *Z. arenicolus* found to have migrated into the water column.

### RESULTS

*Zausodes arenicolus* were 19.3 and 21.0 % of the total harpacticoid assemblage on sampling dates and represented a greater percentage of the Tampa Bay seagrass harpacticoid assemblage compared to other November dates (Table 2). Although typically not a predominant migrating species in subtidal habitats (Walters 1987), November percentages of migrating *Z. arenicolus* were similar to migration percentages of the entire harpacticoid assemblage (Table 2). In the present study, 8.9 % of the total harpacticoid sediment population migrated into the water column during the 1983 postsunset sampling period, while in 1984 only 5.5 % migrated. For *Z. arenicolus*, 20.1 and 5.2 % of the sediment-associated population migrated into the water column in 1983 and 1984, respectively (Table 2). In both years, females predominated over males in the sediment, while males predominated in the water column in 1983 and females in 1984 (Table 3). Equal proportions of gravid and non-gravid females were found in the water column and sediment on both dates; however, a greater percentage of females were gravid in 1983 than 1984 (Table 3). A greater percentage of the total *Z. arenicolus* population was involved in precopulatory clasping in the water column in 1984 compared to 1983 (Table 3).

The size distributions of migrating and non-migrating *Zausodes arenicolus* (Fig. 1) were significantly different in 1983 (K-S, $Z = 2.183$, $p < 0.05$) but not in 1984 (K-S, $Z = 1.216$, $p > 0.05$). Examining the data categorized by sex and stage, the 1983 size distribution of females in the water column was significantly different from that in the sediment (K-S, $Z = 4.538$, $p < 0.05$). The size distribution of migrating females was skewed towards larger size classes in 1983 (Fig. 2). Females between 377 and 439 μm comprised 28 % of all migra-

<table>
<thead>
<tr>
<th>Size class (μm)</th>
<th>Number in water</th>
<th>Number in sediment</th>
<th>Model source pool</th>
<th>Sample random water column distribution</th>
</tr>
</thead>
<tbody>
<tr>
<td>251–271</td>
<td>403</td>
<td>950</td>
<td>1353</td>
<td>413</td>
</tr>
<tr>
<td>272–292</td>
<td>224</td>
<td>590</td>
<td>814</td>
<td>247</td>
</tr>
<tr>
<td>293–313</td>
<td>148</td>
<td>315</td>
<td>463</td>
<td>131</td>
</tr>
<tr>
<td>314–334</td>
<td>54</td>
<td>80</td>
<td>134</td>
<td>43</td>
</tr>
<tr>
<td>335–355</td>
<td>19</td>
<td>30</td>
<td>49</td>
<td>17</td>
</tr>
<tr>
<td>356–376</td>
<td>3</td>
<td>0</td>
<td>3</td>
<td>0</td>
</tr>
<tr>
<td>377–397</td>
<td>0</td>
<td>5</td>
<td>5</td>
<td>1</td>
</tr>
<tr>
<td>398–418</td>
<td>1</td>
<td>0</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>Total</td>
<td>852</td>
<td>1970</td>
<td>2822</td>
<td>852</td>
</tr>
</tbody>
</table>

Table 1. *Zausodes arenicolus*. Example of source pool generation for size-class null models using data from the male size frequency distribution in 1983. Comparison of the actual number collected in the water column (Column 2) with the randomly generated water column distribution (Column 5) tests whether observed size-class migration patterns are significantly different from migration pattern expected under the null model.
Table 2. Numbers (100 cm\(^{-2}\)) of migrating and non-migrating *Zausodes arenicolus* and total harpacticoids collected during postsunset periods on various dates in a Tampa Bay, Florida, subtidal seagrass habitat. Data are from Walters & Bell (1986), Walters (1987) and the present study.

<table>
<thead>
<tr>
<th>Date</th>
<th>Z. arenicolus</th>
<th>Total harpacticoids</th>
<th>% Z. arenicolus of total</th>
<th>% Z. arenicolus migrating</th>
<th>% Total migrating</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Migrating</td>
<td>Non-migrating</td>
<td>Migrating</td>
<td>Non-migrating</td>
<td>Migrating</td>
</tr>
<tr>
<td>Walters &amp; Bell (1986)</td>
<td>Nov 1981</td>
<td>38</td>
<td>132</td>
<td>711</td>
<td>2106</td>
</tr>
<tr>
<td>Present study</td>
<td>Nov 1983</td>
<td>242</td>
<td>957</td>
<td>555</td>
<td>5653</td>
</tr>
<tr>
<td></td>
<td>Nov 1984</td>
<td>69</td>
<td>1261</td>
<td>355</td>
<td>6118</td>
</tr>
</tbody>
</table>


<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>Female</td>
<td>35</td>
<td>47</td>
<td>42</td>
<td>51</td>
</tr>
<tr>
<td>Male</td>
<td>46</td>
<td>29</td>
<td>38</td>
<td>25</td>
</tr>
<tr>
<td>Copepodite</td>
<td>19</td>
<td>24</td>
<td>20</td>
<td>24</td>
</tr>
<tr>
<td>Gravid female (as % of all females)</td>
<td>17</td>
<td>15</td>
<td>7</td>
<td>5</td>
</tr>
<tr>
<td>Clasping (as % of total <em>Zausodes</em>)</td>
<td>11</td>
<td>13</td>
<td>17</td>
<td>6</td>
</tr>
</tbody>
</table>

Fig. 1. *Zausodes arenicolus*. Size frequency distributions of migrating and non-migrating copepods collected in November 1983 and 1984. Frequency is expressed as the percentage in each size class. n: actual numbers collected.
between migrating and non-migrating females (K-S, Z = 0.559, p > 0.05), males (K-S, Z = 0.955, p > 0.05) or copepodites (K-S, Z = 1.114, p > 0.05).

Results of comparisons between actual and null-model-generated size distributions indicated that the size distribution of migrating Zausodes arenicolus collected in 1983 was significantly different from randomly generated water column distributions in 10 out of 10 runs of the model (Table 4). The actual 1984 size distribution of migrating Z. arenicolus, however, was indistinguishable from randomly generated size distributions in 9 out of 10 trials. We also compared observed and null-model-generated size distributions for the 1983 data categorized by sex and stage. The distribution of migrating females was significantly different from randomly generated distributions; however migrating male and copepodite distributions were indistinguishable from distributions generated by the null model (Table 4).

### Table 4. Summary of results for comparisons of actual and null-model-generated water column size frequency distributions in 1983 and 1984. Listed are the number of times out of 10 that the null hypothesis was rejected. (H0: observed water column size frequency distribution not significantly different from randomly generated distribution)

<table>
<thead>
<tr>
<th>Year</th>
<th>Group</th>
<th>Model results</th>
</tr>
</thead>
<tbody>
<tr>
<td>1983</td>
<td>Total distribution</td>
<td>10</td>
</tr>
<tr>
<td></td>
<td>Females</td>
<td>10</td>
</tr>
<tr>
<td></td>
<td>Males</td>
<td>0</td>
</tr>
<tr>
<td></td>
<td>Copepodites</td>
<td>0</td>
</tr>
<tr>
<td>1984</td>
<td>Total distribution</td>
<td>1</td>
</tr>
</tbody>
</table>

**DISCUSSION**

Size significantly influenced the migration of Zausodes arenicolus only in 1983 and the overall effect resulted from significantly larger females entering the water column. Migrations of males and copepodites in 1983 and any sex or stage in 1984 were not influenced by size. The 1983 migration of larger females into the water column was also significantly different from that predicted by the null model, implying that deterministic forces are affecting female migrations. Different sized males and copepodites appeared to migrate randomly into the water column on both dates.

The non-random migration of larger females in 1983 is inconsistent with the hypothesis that harpacticoid migration is linked to mating activity (see Bell et al. 1988). Precocious clasping, grasping of subadult stage IV or V females by adult males, is typical in Zausodes arenicolus (Kern et al. 1984) suggesting that clasping pairs, adult males and smaller female copepodites should predominate in the water column if migration is linked to mating. Clasping pairs were not found more commonly in the water in 1983, and while adult males are predominant migrants, it is the larger size classes of females that were more common in the water column in 1983. The migration of larger females and the observation that larger, gravid females are more common in the water column compared to the sediment (Service pers. obs.) suggests that female migration may be linked to other aspects of reproductive behavior, such as naupliar release. A more detailed study of reproduction in Z. arenicolus is required to test the naupliar release hypothesis.

The singular 1983 occurrence of significantly larger female migrators may be related to variability in the
year-to-year reproductive cycle of Zausodes arenicolus. Compared to 1984, a greater percentage of the Z. arenicolus population was gravid in both the water column and sediment in 1983. Reduced numbers of gravid females and an increase in the percentage of claspers of pairs in the water column suggests that the 1984 population was in an earlier phase of population growth. It is possible that larger females in the appropriate stage of the reproductive process were not available in 1984, resulting in the lack of a significant size difference in female migrations.

Null or neutral models have been used to examine numerous patterns in biology (Gilpin & Diamond 1982, Matthews 1982, Silvertown 1983, Simberloff 1984, Gotelli et al. 1987, Tokeshi & Townsend 1987, Clarke 1988, Lambshedd & Platt 1988), but few investigations of single species population processes, as in the present study, have employed null models. The utility of null models has remained controversial (Harvey et al. 1983, Nitecki & Hoffman 1987), and a number of potential problems with the approach are illustrated by studies attempting to detect the effects of competition in species co-occurrence data (Connor & Simberloff 1979, 1983, 1984, Diamond & Gilpin 1982, Gilpin & Diamond 1982, 1984). Formulation of a realistic source pool, selection of unlikely species combinations, and construction of an unbiased model that does not reflect past competitive interactions or account for the special properties of individual species all confound the multi-specific use of null models. If competition, or other multispecies processes, is not the major focus of a study, deciding on an appropriate null model and accompanying source pool is less of a problem.

Examining patterns within populations avoids many of the problems associated with multispecies null models. Single species models appear to require fewer assumptions compared to models analyzing community processes. Unreasonable combinations generated during Monte Carlo simulations from source pools containing species that naturally do not co-occur are unlikely in population level analyses. In the present study, the null model combined all sizes of Zausodes arenicolus since all sizes typically co-occur in the sediments. Our approach also avoids problems concerning construction of the source pool from which random draws are taken. The size source pool is readily identifiable and, unlike models attempting to detect patterns in multi-species distributions, there is no question as to the origins of migrating individuals. Sample sizes were adequate to represent relative size frequency distributions and emergence traps were sealed to the sediment surface.

Comparison of actual with null-model-generated size distributions of migrating Zausodes arenicolus suggested that statistical differences in the sizes of migrating individuals in 1983 resulted from a real difference in behavior. We can conclude that the migration of larger females was not a chance event and probably resulted from some deterministic cause (i.e. naupliar release).

Null model results increase our confidence in stating that size, at times, influences Z. arenicolus migrations. Although null model results corroborated initial K-S findings that significantly larger females migrated, many instances exist where use of null models leads to different interpretations (Rathke 1984, Simberloff 1984, Gotelli et al. 1987). Such studies emphasize that it is not always necessary to ascribe a deterministic cause to a pattern when a stochastic one will suffice.

Null models can have an important role in exploratory data analysis; however they generally have been underutilized in marine studies (but see Platt & Lambshedd 1985, McCoy et al. 1986, Gotelli et al. 1987, Clarke 1988, Lambshedd & Platt 1988, Meyers et al. 1988). Properly constructed null models identify patterns that can be statistically distinguished from random and pinpoint areas for future experimental research. Null model results in the present study suggest that the size distribution of migrating females was not influenced solely by stochastic factors. Experimental investigations of deterministic factors affecting the size distribution of migrating Zausodes arenicolus are warranted.

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LITERATURE CITED

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