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Variation in egg spawning among subpopulations of sea urchins *Strongylocentrotus droebachiensis*: a theoretical approach

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ABSTRACT: Variation in zygote production, by determining the initial size of the larval pool, can affect the population dynamics of marine invertebrates with a planktonic larval stage. In this paper, we model temporal and spatial patterns in egg spawning of the sea urchin Strongylocentrotus droebachiensis in the shallow subtidal zone along the Atlantic coast of Nova Scotia. In this region, population outbreaks of sea urchins have caused major transitions from kelp beds to barrens (areas devoid of fleshy macroalgae). We define 3 community states and associated subpopulations of sea urchins: (1) the established kelp bed state, where sea urchins are in low density and small; (2) the transition state, with sea urchins in kelp beds, newly created (transitional) barrens, and grazing fronts (dense aggregations of large sea urchins along the edges of kelp beds); and (3) the barrens state after fronts have dispersed and intermediate densities of moderately sized sea urchins remain in post-transitional barrens. Using data from the literature and unpublished sources, we parameterize mathematical models to predict egg spawning for each subpopulation, both on an areal basis and for the entire coast. The predicted number of eggs spawned per m² is 1 order of magnitude higher in grazing fronts (7.1×10^7) than in transitional and post-transitional barrens (5.8×10^6) and 4.4×10^6 , respectively), and 4 to 6 times higher in barrens than in established kelp beds (1.0×10^6) . Differences among subpopulations in the number of eggs spawned are directly related to differences in adult density, female gonad output, and body size. The total number of eggs spawned (all subpopulations combined) on a coastal scale increases \sim 6-fold from the established kelp bed state (5.6 \times 10¹⁴) to the late transition state (3.1×10^{15}) and then drops slightly in the barrens state (2.4×10^{15}) . During most of the transition state, sea urchins in barrens spawn the greatest number of eggs. Based on published values, we estimate that fertilization rates are highest in grazing fronts (62%), intermediate in transitional and post-transitional barrens (36 and 43 %, respectively) and lowest in kelp beds (15 %), resulting in up to 2 orders of magnitude differences in the number of zygotes produced in these subpopulations $(4.4 \times 10^7, 2.1 \times 10^6, 1.9 \times 10^6, \text{ and})$ 1.5×10^{5}). Total zygote production (all subpopulations combined) on a coastal scale increases ~16-fold from the established kelp bed state (8.1×10^{13}) to the late transition state (1.3×10^{15}) and then drops slightly in the barrens state (1.1×10^{15}) . During most of the transition state, sea urchins in barrens contribute the greatest number of zygotes to the total zygote pool, with those in grazing fronts contributing up to 44% of all zygotes. Sensitivity analysis and evaluation of model assumptions indicate that our predictions should be correct in a relative sense (i.e. comparing among subpopulations), although absolute numbers may be overestimated. A disparity between temporal patterns of estimated zygote production and observed settlement rates of S. droebachiensis in the shallow subtidal zone suggests that zygote production interacts with other factors to determine settlement rates of this species.

KEY WORDS: Sea urchins \cdot Strongylocentrotus droebachiensis \cdot Egg spawning \cdot Fertilization rate \cdot Zygote production \cdot Kelp beds \cdot Barrens \cdot Grazing fronts

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INTRODUCTION

In marine benthic invertebrates with a planktonic larval stage, variation in the supply of settling larvae to

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adult populations is an important determinant of population structure and dynamics (e.g. Pearse & Hines 1987, Minchinton & Scheibling 1991, Grosberg & Levitan 1992). The overall supply of settlers is fundamentally related to the total number of zygotes produced at a multitude of sites. Zygote production in free-spawn-

ing marine invertebrates is determined by the number of eggs spawned and the proportion of those eggs that is fertilized. Both egg spawning and fertilization rates are influenced by a variety of environmental (e.g. food supply, hydrodynamic regime) and demographic factors (e.g. population density, size structure, gamete output), which show marked spatial and temporal variation among different adult populations. As a result, adult populations do not contribute equally to the overall pools of eggs and zygotes produced in a region. To date, there have been few attempts to evaluate how populations vary in their relative contribution to egg or zygote pools (Keats et al. 1984), or how the size of the pools may vary over time. Empirical analysis of this problem is hampered by logistical difficulties. However, an extensive body of work on factors that influence egg spawning in combination with an increased understanding of fertilization ecology (reviewed by Levitan 1995) and published fertilization estimates enable a theoretical approach.

Fecundity and fertilization rate, which together determine zygote production, are influenced by the simple and interactive effects of a variety of factors. Fecundity usually increases in both absolute and relative terms with body size (e.g. Gonor 1972, Kettle & Lucas 1987, Meidel & Scheibling 1998b) and with higher food quality and/or quantity (e.g. Vadas 1977, Scheibling 1981). A high density of spawning adults generally increases the number of eggs spawned per unit area although it may negatively influence body size or gamete output of individuals (Branham et al. 1971, Levitan 1988a, 1991, Wahle & Peckham 1999). Fertilization rates have been shown to increase with adult density (Pennington 1985, Grosberg 1991, Levitan 1991, Oliver & Babcock 1992, Yund 1995, Coma & Lasker 1997) and in some cases with sperm production (Babcock et al. 1994, Yund 1998). Empirical (Pennington 1985, Levitan et al. 1992, Wahle & Peckham 1999) and theoretical (Denny & Shibata 1989, Young et al. 1992, Levitan & Young 1995) studies have also indicated that a low flow regime during spawning increases fertilization rate by reducing gamete dilution.

Well-studied species which exhibit large fluctuations in population density, demography, and reproductive output are ideally suited for modelling variation in egg spawning and zygote production. One such species is the sea urchin *Strongylocentrotus droebachiensis* (Müller), for which changes in population size and structure have been well documented along the Atlantic coast of Nova Scotia over the last 3 decades (Lang & Mann 1976, Wharton & Mann 1981, Scheibling et al. 1999). When sea urchins are in low abundance, kelp beds (consisting mainly of the kelps *Laminaria longicruris* and *L. digitata* with an understory of coralline and fleshy macroalgae) thrive in the shallow

rocky subtidal zone (Edelstein et al. 1969, Mann 1972). Sea urchins exposed to an abundance of food in the form of drift algae and small understory plants grow rapidly and produce large gonads (Lang & Mann 1976, Meidel & Scheibling 1998a,b). As the density of sea urchins increases, large individuals start to aggregate at the edge of kelp beds, forming fronts (Lang & Mann 1976, Wharton 1980b, Bernstein et al 1981, Scheibling et al. 1994, 1999) which destructively graze the kelp. Sea urchins in fronts climb up onto plants, weigh them down and consume them entirely (Breen & Mann 1976a, Mann 1977). The facilitative effect of group feeding on an abundant food source enables sea urchins in this zone to grow and reproduce at a high rate despite high densities (Meidel & Scheibling 1998a,b). The formation of grazing fronts initiates a large-scale transition in community state, as fronts advance into the kelp bed and create, in their wake, barren grounds dominated by coralline algae (Bernstein et al. 1981, Johnson & Mann 1986, Chapman & Johnson 1990). This transition from the kelp bed state to the barrens state occurs within about a decade in Nova Scotia (Breen & Mann 1976a, Scheibling et al. 1999). After the destruction of kelp beds, sea urchins in barrens remain relatively abundant but their nutritional state deteriorates, leading to declines in growth and gonad production (Lang & Mann 1976, Meidel & Scheibling 1998a,b). The subtidal zone persists in the barrens state until sea urchins are eliminated by disease (Miller 1985, Scheibling 1986). The ensuing reduction in grazing pressure results in rapid growth of macroalgae, and kelp beds are reestablished within 2 to 3 yr (Scheibling 1986, Novaczek & McLachlan 1986, Johnson & Mann 1988). Following mass mortalities, sea urchins begin to recolonize the developing kelp beds via larval settlement (Miller 1985, Raymond & Scheibling 1987), but the recovery of adult populations may be limited by intensive predation on recruits (Scheibling 1996). The mechanisms causing population outbreaks of S. droebachiensis have not yet been conclusively demonstrated, but increased survival of the planktonic larvae (Hart & Scheibling 1988) and migration of sea urchins from deeper waters into the shallows (Scheibling et al. 1999) have been hypothesized. The role of variation in zygote production in initiating outbreaks has not been explored.

In this paper, we use mathematical models to predict the number of eggs spawned by *Strongylocentrotus droebachiensis* per unit area, and on a coastal scale, in kelp beds, barrens, and grazing fronts along the Atlantic coast of Nova Scotia. We parameterize these models using sea urchin population data obtained from the literature and unpublished observations. We also estimate the relative contributions of sea urchins in these different habitats to the overall number of eggs

spawned in this region during the transition from kelp beds to barrens. Our goal is to use our estimates of egg spawning and published estimates of fertilization rates to determine spatial and temporal variation in zygote production and to evaluate the importance of this variation to the population dynamics of *S. droebachiensis* along this coast.

MODEL CONSTRUCTION AND PARAMETER ESTIMATION

Density and demography of subpopulations

Fluctuations in populations of Strongylocentrotus droebachiensis, and hence in community state, occur along much of the Atlantic coast of Nova Scotia, where disease outbreaks periodically eliminate sea urchins (Miller 1985, Scheibling 1986, Scheibling & Hennigar 1997; Fig. 1). In our model we include 3 different community states and the associated sea urchin subpopulations in this region of coast: (1) the established kelp bed state, with sea urchins in kelp beds; (2) the transition state, with sea urchins in kelp beds, transitional barrens, and grazing fronts; and (3) the barrens state, with sea urchins in post-transitional barrens. These community states generally succeed each other in any one area but also can occur simultaneously in different areas. We also consider the other transition state when kelp beds develop following a mass mortality of sea urchins. Although sea urchins recruit to the developing kelp beds, this state is distinguished from the established kelp bed state by the absence of adult sea urchins and therefore is not included in our model. We use the term 'subpopulation' for sea urchins in a particular habitat (kelp beds, transitional or post-transitional barrens) or zone (grazing fronts) in the ecological sense, not implying genetic differentiation.

To parameterize our model, we compile data on the population density, the proportion of adults, and the mean and maximum body size of adults in each of the 3 subpopulations we consider (Table 1). These data come from investigations on populations of Strongylocentrotus droebachiensis along the Atlantic coast of Nova Scotia over a 25 yr period. We estimate the proportion of adults and the mean size and maximum adult size from recorded size-frequency distributions which we reconstruct (where necessary) in 5 mm size classes. The proportion of adults depends on the size (test diameter) at first reproduction, which ranges between 18 and 25 mm in S. droebachiensis (Vadas 1977, Thompson 1979, Scheibling 1986, Raymond & Scheibling 1987, Munk 1992). Because sea urchins tend to reproduce earlier in life, and therefore at a smaller size, in a nutritionally rich environment

(Buchanan 1966, Sivertsen & Hopkins 1995, Meidel & Scheibling 1999), we use 20 mm as the size at first reproduction in kelp beds and grazing fronts, and 25 mm in transitional and post-transitional barrens. Mean adult size is estimated by multiplying the midpoint of each 5 mm size class by the proportion of adult sea urchins in that class and summing the resulting values over all adult size classes.

Egg spawning

Eggs spawned per unit area

We calculate the total number of eggs spawned (O) per unit area (U, m^2) in subpopulation k (sea urchins in kelp beds, grazing fronts, transitional or post-transitional barrens) as the sum of the eggs spawned by females in all size classes of adults:

$$O_k^{(U)} = \sum_j D_k a_k f \ p(j)_k \frac{E_j}{5.56 \times 10^{-7}}$$
 (1)

where D is sea urchin density (no. m⁻²), a is the proportion of adults, f is the proportion of females, p is the proportion of sea urchins in a 5 mm size class of adults denoted by its mid-point j, E is fecundity (dry weight of eggs released, g) of a female sea urchin and 5.56×10^{-7} is the dry weight per egg (g, Thompson 1979). All symbols used in this and subsequent models are summarized in Table 2.

Mean density and mean proportion of adults in each subpopulation (Table 3) are estimated from published data (Table 1). Because *Strongylocentrotus droe-*

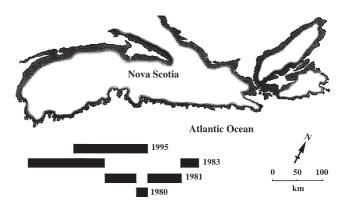


Fig. 1. Map of Nova Scotia showing the extent of mass mortalities of *Strongylocentrotus droebachiensis* during 1980 to 1983 and in 1995. Near-complete mass mortality is indicated by black bars. The maximum range of mass mortality (1983) denotes the extent of the shallow subtidal zone over which the number of eggs spawned and total zygote production is predicted. Gaps between bars in 1981 and 1983 indicate areas where sea urchins had been eliminated by disease in previous years. (Adapted from Scheibling & Hennigar 1997)

bachiensis usually has a sex ratio of 1:1 (Munk 1992, Meidel & Scheibling 1998b), we set the proportion of females to 0.5 in all subpopulations (Table 3). The

range of size classes of adults differs among subpopulations depending on the size at first reproduction (see 'Density and demography of subpopulations') and the

Table 1. Summary of published data on population density, proportion of adults, mean and maximum adult size of *Strongylocentrotus droebachiensis* in kelp beds, grazing fronts, transitional and post-transitional barrens along the Atlantic coast of Nova Scotia

Location	Sampling period	Population density (no. m ⁻²)	Proportion of adults	Mean adult size (mm)	Maximum adult size (mm)	Source
Kelp beds						
St. Margaret's Bay	1973	230	0.02	32.5	35	Breen & Mann (1976b) ^a
Boutilier Point	1977	29.5				Chapman (1981)
Boutilier Point	10/1977-1/1980	0				Bernstein et al. (1981)
Coyle Cove	6/1981	13	0.20	30.4	50	Miller (1985a)
Little Duck Island	1992-1993	20	0.45	30.7	60	Scheibling et al. (1999) ^b
Mill Cove	1992-1993	5	0.23	24.8	35	Scheibling et al. (1999)
Grazing fronts						
St. Margaret's Bay	1973		0.73	37.2	55	Breen & Mann (1976a)
St. Margaret's Bay	1973	240	0.88	39.8	60	Breen & Mann (1976b)
Clark's Harbour	1978/79	48	0.79	40.3	75	Wharton & Mann (1981) ^c
Boutilier Point	1978	50				Bernstein et al. (1981) ^d
Little Duck Island	1992-1993	256	0.95	43.1	65	Scheibling et al. (1999) ^b
Mill Cove	1992-1993	84	0.73	34.8	55	Scheibling et al. (1999)
Transitional barrens						3 (, ,
St. Margaret's Bay	1973	75	0.54	36.1	55	Breen & Mann (1976b) ^e
Boutilier Point	10/1977-1/1980					Bernstein et al. (1981) ^f
Eagle Head	3/1983	24	0.48	34.9	50	Scheibling & Stephenson (1984
Eagle Head	3/1983	26	0.71	32.9	50	Scheibling & Stephenson (1984
Coyle Cove	6/1981	18	0.60	38.6	70	Miller (1985)
Little Duck Island	1992-1993	80	0.20	33.4	60	Scheibling et al. (1999) ^b
Mill Cove	1992-1993	48	0.18	31.4	50	Scheibling et al. (1999)
Post-transitional barrens ^g						3 (, ,
Horse Island (2)	1973	80	0.22	31.4	45	Breen & Mann (1976b)
Luke Island (5)	1973	110	0.04	30.3	35	Breen & Mann (1976b)
Northwest Cove (1)	5-9/1975	50				Lang & Mann (1976)
Northwest Cove (3)	5-9/1975	70				Lang & Mann (1976)
Northwest Cove (4)	5-9/1975	95				Lang & Mann (1976)
Ingomar	1978/79	105.4	0.10	38.0	60	Wharton (1980a)
Port Mouton	1978/79	107.9	0.13	35.4	60	Wharton (1980a)
Blue Rocks	1978/79	61.9	0.04	30.0	40	Wharton (1980a)
Northwest Cove	1978/79	67.5	0.13	30.8	50	Wharton (1980a)
Mushaboom	1978/79	55.0	0.58	32.6	55	Wharton (1980a)
Drum Head	1978/79	29.4	0.92	38.6	55	Wharton (1980a)
Dover	1978/79	45.6	0.71	30.7	50	Wharton (1980a)
Northwest Cove (1)	5-9/1975	53	0.64	33.4	50	Wharton (1980a)
Northwest Cove (4)	5-9/1979	96	0.23	29.3	40	Wharton (1980a)
Luke Island (5–6)	1973	112	0.05	30.5	35	Wharton (1980a)
St. Margaret's Bay (8)	1975 (?)	43	0.09	31.9	45	Wharton (1980a)
St. Margaret's Bay (11–12)	1970s	25	0.42	30.4	45	Wharton (1980a)
Louse Head	8/1984	89.3	0.45	41.0	60	Scheibling (1986)
Cape Mocodome	8/1984	49.8	0.59	46.5	70	Scheibling (1986)

^aPopulation density and proportion of adults from this study are not included in calculating the means of these 2 parameters as they appear to be distorted by a heavy recruitment event

^bData from 1994 and 1995 are not used because of a mass mortality in fall 1993 at Little Duck Island

^cSite 2b in this study

^dOnly data from July and December 1978 are used because densities in other months do not indicate the existence of a grazing front

 $^{{}^{\}rm e}{\rm Data}$ collected 5 m from the edge of the kelp bed

^fData averaged from exposed and hidden sea urchins collected during the day

 $^{{}^{}g}$ Numbers in parenthesis indicate years since kelp bed destruction

Table 2. List of symbols used

- a Proportion of adults
- A Total area occupied by a subpopulation
- B Total wet body weight
- c Constant in logistic Eq. (6)
- D Population density
- E Dry weight of eggs released
- f Proportion of females
- F Fertilization rate
- F_0 Fertilization rate in the absence of spawners
- F_{m} Asymptotic fertilization rate
- I Proportion of total body wet weight that is gonads
- *j* Size (mid-point of a size class)
- k Subpopulation
- ${\cal O}^{(U)}$ Number of eggs spawned per unit area by a subpopulation
- $\mathcal{O}^{(C)}$ Number of eggs spawned on a coastal scale by a subpopulation
- $O^{(T)}$ Total number of eggs spawned on a coastal scale
- p(j) Proportion of sea urchins in a size class of adults
- r Proportion of spawning adults
- R Proportion of I that is spawned
- s Density of spawners
- S Proportional sensitivity (elasticity)
- v Original model parameter in sensitivity analysis
- v* Altered model parameter in sensitivity analysis
- X Original model output in sensitivity analysis
- X^* Altered model output in sensitivity analysis
- $Z^{(U)}$ Number of zygotes produced per unit area by a subpopulation
- $Z^{(C)}$ Number of zygotes produced on a coastal scale by a subpopulation
- $Z^{(T)}$ Total number of zygotes produced on a coastal scale

maximum size. For each subpopulation, the mean proportion of sea urchins in each size class of adults (Table 3) is estimated from the sources in Table 1 by summing the proportions in each class over all studies and dividing that sum by the total proportion in all adult classes. Because the proportion of sea urchins >60 mm is low (≤ 0.02) in all subpopulations, we use 55-60 mm as the largest size class.

Fecundity (E) for size classes j=22.5-57.5 mm in kelp beds and grazing fronts, and j=27.5-57.5 mm in transitional and post-transitional barrens, is estimated from the spawned wet weight of gonads and a wet to dry weight conversion coefficient for gonads of *Strongylocentrotus droebachiensis* estimated by regression $(r^2=0.970, n=87)$:

$$E_{j,k} = 0.249 I_{j,k} R_k B_j (2)$$

where I is the maximum proportion of the total body wet weight that is gonads, R is the proportion of gonad weight that is spawned, and B (g) is the total wet body weight. In kelp beds, grazing fronts and transitional

barrens, we estimate I (Table 3) from published data (Meidel & Scheibling 1998b). Because I increases with size in young adult sea urchins, we estimate I for j =22.5–32.5 mm using a logistic equation given by Meidel & Scheibling (1998b). For larger adults, I is at an asymptotic level and is estimated as the average I at the peak of the reproductive cycle, based on measurements over 2 yr and at 2 sites for each subpopulation (Meidel & Scheibling 1998b). We assume that in posttransitional barrens I is 30% of that in kelp beds (Lang & Mann 1976). In kelp beds, grazing fronts and transitional barrens, we estimate R, the proportion of gonads spawned (Table 3), based on published data as the decrease in wet gonad mass directly after spawning (Meidel & Scheibling 1998b). We assume that sea urchins in post-transitional barrens release the same proportion of gonadal material as those in transitional barrens. We estimate B at each size j using the following regression ($r^2 = 0.995$, n = 424; Scheibling et al. 1999):

$$\ln B_i = 2.81 \ln j - 7.0 \tag{3}$$

Eggs spawned on a coastal scale

The number of eggs spawned (O) on a coastal scale (C) in subpopulation k is calculated as

$$O_k^{(C)} = O_k^{(U)} A_k (4)$$

where A is the total area occupied by a subpopulation (km²). The total number of eggs spawned by all subpopulations combined in the shallow rocky subtidal zone (T) is calculated as

$$O^{(T)} = \sum_{k} O_{k}^{(C)} \tag{5}$$

We consider only subpopulations along that section of the coast of Nova Scotia where near-complete mass mortalities of Strongylocentrotus droebachiensis have been documented (Fig. 1). We also limit the offshore distribution of these subpopulations to a depth of 15 m, where >90 % of the total algal biomass in this region is found (Mann 1972, Moore & Miller 1983). The total area occupied by a subpopulation is estimated using surveys of the shallow (to 15 m depth) subtidal zone of the Atlantic coast of Nova Scotia by Moore & Miller (1983) and Moore et al. (1986). These surveys determined the total area (km 2) that consisted of >50 % hard substratum and was deemed suitable habitat for sea urchins and macroalgae. The section of the coast we consider includes the entire area surveyed by Moore & Miller (1983), which they estimate to have hard substratum covering 512 km². Of the area surveyed by Moore et al. (1986), only the portion of eastern Halifax 40-45 mm, j = 42.5 mm

45-50 mm, j = 47.5 mm

50-55 mm, j = 52.5 mm

55-60 mm, j = 57.5 mm

20-25 mm, j = 22.5 mm

25-30 mm, j = 27.5 mm

30-35 mm, j = 32.5 mm

35-40 mm, j = 37.5 mm

40-45 mm, j = 42.5 mm

45-50 mm, j = 47.5 mm

50-55 mm, j = 52.5 mm

55-60 mm, j = 57.5 mm

Density of spawners

Proportion of spawning adults

Proportion of total body weight that is gonads

Proportion of gonad weight that is spawned

Parameter Symbol Units Transitional Post-transitional Kelp Grazing beds fronts barrens barrens Dno. m^{-2} Population density (SE) 14 (5.3) 136 (46.4) 41 (10.1) 71 (6.4) Proportion of adults (SE) а 0.29(0.09)0.82(0.04)0.45(0.07)0.33(0.07)f 0.50 0.50 Proportion of females 0.50 0.50 Proportion of adults in size class p 20-25 mm, j = 22.5 mm0.409 0.123 25-30 mm, j = 27.5 mm0.243 0.133 0.288 0.338 30-35 mm, j = 32.5 mm0.1610.1720.323 0.232 35-40 mm, j = 37.5 mm0.059 0.157 0.206 0.165

0.089

0.026

0.007

0.006

0.008

0.030

0.085

0.205

0.205

0.205

0.205

0.205

0.70

0.33

1.3

 $no.\ m^{-2}$

R

r

 \boldsymbol{S}

0.146

0.131

0.084

0.033

0.010

0.038

0.098

0.159

0.159

0.159

0.159

0.159

0.72

0.33

37

0.097

0.057

0.014

0.010

0.025

0.062

0.112

0.112

0.112

0.112

0.112

0.77

0.33

6

0.128

0.077

0.035

0.016

0.009

0.026

0.062

0.062

0.062

0.062

0.062

0.77

0.33

8

Table 3. Basic parameters used to calculate the number of eggs spawned and zygotes produced per unit area in kelp beds, grazing fronts, transitional and post-transitional barrens

county (\sim 27 km²) is relevant here, which brings the total estimated area of suitable habitat for sea urchins and macroalgae to 539 km².

When the shallow subtidal zone is in the established kelp bed or the barrens state, the entire area (A) is occupied by sea urchins either in kelp beds or in post-transitional barrens. During the transition state, when large kelp beds are slowly grazed down and replaced by transitional barrens, we gradually decrease the area of kelp beds from 100 to 0% in 1% steps, and correspondingly increase the area of barrens. Following the removal of all kelp, transitional barrens change into post-transitional barrens relatively quickly. To account for this change, we decrease (increase) the area of transitional (post-transitional) barrens incrementally by 10% from the onset of the barrens state until the entire rocky subtidal zone is converted to post-transitional barrens.

The total area represented by grazing fronts on a coastal scale is estimated by assuming a front width of 2 m (Bernstein et al. 1981, Scheibling et al. 1999) along the entire length of the rocky coast. As grazing fronts may be convoluted or discontinuous along the edge of a kelp bed (pers. obs.), our estimate of the area represented by grazing fronts is a rough approximation. The

surveys (Moore & Miller 1983, Moore et al. 1986) give a total length of coastline suitable for sea urchins of 1609 km (1400 km for the southwestern shore and 209 km for eastern Halifax County). Based on these estimates of front width and length, the total area occupied by grazing fronts is $\sim 3.2~{\rm km^2}$. This area is assumed to be constant during most of the transition state, and subtracted from the estimated area of transitional barrens. Because grazing fronts form and disperse over time, we incrementally increase the area occupied by this subpopulation in the early stages of the transition state by 10% and similarly decrease it in the early stages of the barrens state.

Fertilization rate

We estimate the fertilization rate in each subpopulation based on Wahle & Peckham's (1999) field experiments on the effect of $Strongylocentrotus\ droebachiensis\ density\ (2.25,\ 36,\ and\ 144\ sea\ urchins\ m^{-2})\ on$ fertilization rate in weak currents (speed < 5 cm s⁻¹) and in the absence of macroalgae. To estimate fertilization rates at the densities of spawners in our subpopulations, we fit Wahle & Peckham's data (as

presented in their Fig. 6) with the following logistic function:

$$F = \frac{F_0 F_{\rm m}}{F_0 + (F_{\rm m} - F_0) e^{-cF_{\rm m}s}}$$
 (6)

where F is fertilization rate (%), F_0 is fertilization rate in the absence of spawners, $F_{\rm m}$ is asymptotic fertilization rate, c is a constant and s is spawner density (no. m⁻²). Wahle & Peckham also determined the interactive effects of population density (2.25 and 36 sea urchins m⁻²) and macroalgal (*Laminaria saccharina*) cover on fertilizations in strong currents (mean speed \pm SE: 9.3 ± 3.7 cm s⁻¹ outside kelp cover). Because current speed and absence/presence of macroalgae had little effect on fertilizations at the low population density, our approach of estimating fertilization rates in kelp beds based on a weak current in the absence of macroalgae is appropriate.

The density of spawners (s, Table 3) in each subpopulation is estimated based on population density (D), the proportion of adults (a), and the proportion of adults that spawn during 1 spawning event (r, Table 3). Previous studies have shown that natural spawning events in Strongylocentrotus droebachiensis may involve only a small proportion of the population (Keats et al. 1987, Pearse et al. 1988). To account for spawning asynchrony, we multiply adult densities in the subpopulations by the proportion of adults that spawn during a spawning event (Table 3). This parameter is estimated as the mean proportion of adult sea urchins that were partially spawned or spent on a single sampling date during the peak of the main spawning season (March to May), pooled over kelp beds, grazing fronts and transitional barrens (Meidel & Scheibling 1998b).

Zygote production

The number of zygotes produced (Z) per unit area (U) in subpopulation k is estimated as

$$Z_k^{(U)} = \frac{O_k^{(U)} F_k}{100} \tag{7}$$

where F is the fertilization rate (%) estimated from Wahle & Peckham (1999) (see 'Fertilization rate' above).

The number of zygotes produced (Z) on a coastal scale (C) in subpopulation k is estimated as

$$Z_k^{(C)} = Z_k^{(U)} A_k (8)$$

The total number of zygotes produced by all subpopulations combined in the shallow rocky subtidal zone (T) is calculated as

$$Z^{(T)} = \sum_{k} Z_k^{(C)} \tag{9}$$

MODEL PREDICTIONS

Eggs spawned

We estimate that the total number of eggs spawned per m² (Eq. 1; Fig. 2) differs among subpopulations by more than 1 order of magnitude. It is highest in grazing fronts (7.1×10^7) , intermediate in transitional and post-transitional barrens $(5.8 \times 10^6 \text{ and } 4.4 \times 10^6)$, respectively), and lowest in kelp beds (1.0×10^6) .

The temporal variation in the number of eggs spawned by the different subpopulations on a coastal scale (Eq. 4; Fig. 3) is related to changes in the extent of the subtidal zone that these subpopulations occupy. Egg spawning in kelp beds is highest when the system is in the established kelp bed state (5.6×10^{14}) and declines at an accelerating rate as this habitat is destroyed during the transition state. In the expanding transitional barrens, egg spawning concomitantly increases to reach a maximum (3.1×10^{15}) late in the transition state once kelp beds have been destroyed. While grazing fronts are fully established, sea urchins in this zone occupy, according to our assumption, a constant area and spawn a constant number of eggs (2.3×10^{14}) . In the post-transitional barrens after the fronts have dispersed, egg spawning (2.4×10^{15}) is somewhat lower than in transitional barrens but over 4-fold higher than in established kelp beds.

We predict total egg spawning by *Strongylocentrotus droebachiensis* in the shallow subtidal zone (Eq. 5; Fig. 3) to vary markedly in time depending on the community state. The number of eggs spawned is lowest in the established kelp bed state and progressively

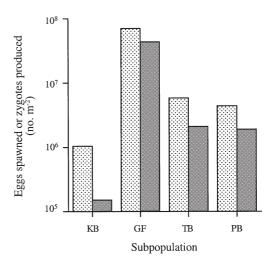


Fig. 2. Estimated number (m⁻²) of eggs spawned (stippled bars) and zygotes produced (grey bars) in kelp beds (KB), grazing fronts (GF) and transitional (TB) and post-transitional (PB) barrens along the Atlantic coast of Nova Scotia

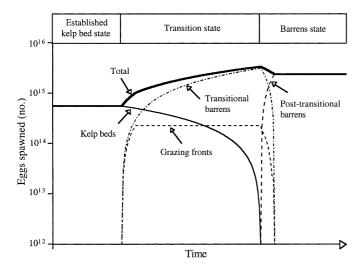


Fig. 3. Estimated number of eggs spawned in kelp beds, grazing fronts, transitional and post-transitional barrens on a coastal scale, and total number of eggs spawned in the entire subtidal zone (bold line, which partly overlaps with lines indicating the number of eggs spawned in kelp beds and post-transitional barrens), as the community moves from the established kelp bed via the transition to the barrens state over time

increases during the transition state as kelp beds are replaced by transitional barrens, reaching a maximum (3.4 \times 10¹⁵) just before grazing fronts disperse. Although sea urchins in fronts occupy <1% of the total subtidal area, they contribute 7 to 22% of all eggs spawned during the transition state. As the entire sub-

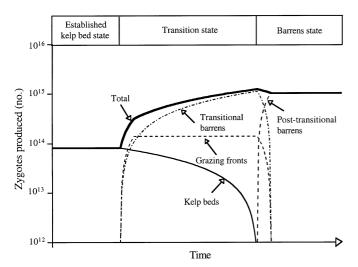


Fig. 4. Estimated number of zygotes produced in kelp beds, grazing fronts, transitional and post-transitional barrens on a coastal scale, and total number of zygotes produced in the entire subtidal zone (bold line, which partly overlaps with lines indicating the number of zygotes produced in kelp beds and post-transitional barrens), as the community moves from the established kelp bed via the transition to the barrens state over time

tidal zone is converted to post-transitional barrens, total egg spawning drops by ~30 % (to 2.4×10^{15}) in the barrens state.

Fertilization rate

Based on Wahle & Peckham's (1999) data, the parameters of the logistic function (Eq. 6) we use to estimate fertilization rates in the subpopulations are F_{0} , fertilization rate in the absence of spawners, 10.6%; $F_{\rm m}$, asymptotic fertilization rate, 62%; and $c_{\rm r}$ a constant, -0.005. The resulting curve provides a better fit to the data $(r^2 = 0.569)$ than Wahle & Peckham's (1999) power function ($r^2 = 0.485$; their Fig. 6). The estimate of 10.6% for the fertilization rate in the absence of spawners is clearly an overestimate which resulted from the particular distribution of data points (12 to 20 fertilization rates at each of 3 densities). The adequacy of our parameter estimates is confirmed, however, by the good match between the fertilization rates estimated from the logistic function (18% at 2.25 sea urchins m⁻², and 62% at both 36 and 144 sea urchins m⁻²) and the mean fertilization rates measured by Wahle & Peckham (18, 62 and 61%, respectively; their Fig. 5). Using the logistic function, we estimate that fertilization rates vary more than 4-fold among subpopulations. The rate is highest in grazing fronts (62.0%), intermediate in transitional and post-transitional barrens (36.1 and 43.3%, respectively), and lowest in kelp beds (14.6%).

Zygote production

The number of zygotes produced per m² (Eq. 7; Fig. 2) differs among subpopulations by more than 2 orders of magnitude. It is highest in grazing fronts (4.4×10^7) , intermediate in transitional and post-transitional barrens (2.1×10^6) and (2.1×10^6) , respectively), and lowest in kelp beds (1.5×10^5) .

Like egg spawning, zygote production by the different subpopulations on a coastal scale (Eq. 8; Fig. 4) shows temporal variation that is again related to changes in the extent of the subtidal zone that these subpopulations occupy. Zygote production in kelp beds is highest when the system is in the established kelp bed state (8.1×10^{13}) and declines at an accelerating rate as this habitat is destroyed during the transition state. In grazing fronts, zygote production is constant (1.4×10^{14}) , while in transitional barrens it increases to a maximum (1.1×10^{15}) late in the transition state once kelp beds have been destroyed. Zygote production is slightly lower (1.0×10^{15}) in post-transitional than in transitional barrens but more than 1

order of magnitude higher than in established kelp beds.

Like total egg spawning, total zygote production by *Strongylocentrotus droebachiensis* in the shallow subtidal zone (Eq. 9; Fig. 4) is estimated to vary markedly in time depending on the community state. Total zygote production is at a minimum in the established kelp bed state and progressively increases as kelp beds are replaced by transitional barrens, reaching a maximum (1.3×10^{15}) just before grazing fronts disperse. Sea urchins in grazing fronts contribute 11 to 44% of all zygotes produced during the transition state. As the entire subtidal zone is converted to post-transitional barrens, total zygote production drops by 16% (to 1.1×10^{15}) in the barrens state.

SENSITIVITY ANALYSIS

We analysed mean proportional sensitivity (*S*, also termed elasticity) based on all 4 subpopulations to measure how sensitive a model output is to small changes in basic parameters:

$$S = \frac{1}{4} \sum_{k} \frac{\frac{X_{k}^{*} - X_{k}}{X_{k}}}{\frac{Y_{k}^{*} - V_{k}}{V_{k}}}$$
(10)

where X^{\star} is the output of the altered model, X is the output of the original model, v^{\star} is the parameter value increased by a small amount, and v is the original parameter value. A proportional sensitivity of 1 means that the model output increases by the same proportion by which the parameter value is increased. Sensitivity analysis indicates which parameters need to be most carefully estimated to obtain accurate model predictions. For our purposes, the analysis also indicates which parameters may be important for explaining differences in model output among subpopulations.

We calculated the effect of an increase in a basic parameter by 1% on the number of eggs spawned per unit area, $(O_k^{(U)}, \text{ Eq. 1})$ and on fertilization rate (F,Eq. 6). Basic parameters for $O_k^{(U)}$ were population density (D) and the proportion of: adults (a), females (f), body weight that is gonads (I), gonad weight that is spawned (R), and females in a size class (p(j)), respectively. To determine the effect of an increase in the proportion of females in a size class, we increased this proportion in 1 size class by 1% and decreased the proportion in each of the other size classes in proportion to the abundance in that class. Basic parameters for F were population density (D), the proportion of adults (a), and the proportion of adults that spawn during a spawning event (r). Only 1 parameter was varied in each analysis.

We estimated the proportion of spawning adults based on the proportion of adult sea urchins that were partially spawned or spent on a single sampling date. Because there was a 1 mo interval between that sample and the previous one, this estimate reflects the proportion of adult sea urchins that had spawned over a month. As a result, we may overestimate r during the short time interval over which fertilizations are likely to occur. To determine how a potential overestimate of r may affect fertilization rate, we calculated the reduction in F when the proportion of adult spawners was reduced to 0.17 and to 0.08 (i.e. representing 2 and 4 spawning events within the 1 mo period; a reduction in r by 50 and 75%, respectively).

Our analysis indicated that the sensitivity of the number of eggs spawned per unit area was identical ($S \pm SE: 1.0 \pm 0.0$) for increases in population density and in the proportion of adults, females, body weight that is gonads, and gonad weight that is spawned, respectively. Sensitivity was variable, and generally low, to changes in the proportions of females in a size class of adults (Fig. 5). The sensitivity to an increase in the proportions of females in the 2 smallest size classes was intermediate (-0.41 ± 0.26 and -0.31 ± 0.07 for j = 22.5 and 27.5 mm, respectively), while sensitivities to increases in other size classes were low (-0.14 ± 0.04 to 0.06 ± 0.01).

The sensitivity of fertilization rate was identical ($S \pm SE: 0.46 \pm 0.19$) for increases in population density, the proportion of adults, and the proportion of adults that spawn during a spawning event. A reduction in the proportion of spawning adults by 50% to account for the

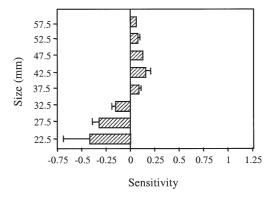


Fig. 5. Mean (±SE) sensitivity of the number of eggs spawned per unit area to a small change in the proportion of females in a size class (represented by its mid-point) of adults. Means at 22.5 mm are based on sensitivities calculated in kelp beds and grazing fronts, while means at all other sizes are based on sensitivities calculated in all 4 subpopulations (kelp beds, grazing fronts, transitional and post-transitional barrens). Proportional sensitivities shown here are equal to the proportional change in the number of eggs spawned when the proportion of females is increased by 1%

possibility of 2 spawning events within a 1 mo period reduced fertilization rates by 14% (to 12.5%) in kelp beds, by 2% (to 61.0%) in grazing fronts, by 40% (to 21.6%) in transitional barrens, and by 50% (to 25.3%) in post-transitional barrens. A reduction in the proportion of spawning adults by 75% to account for the possibility of 4 spawning events within a 1 mo period reduced fertilization rates by 21% (to 11.5%) in kelp beds, by 22% (to 48.2%) in grazing fronts, by 57% (to 15.5%) in transitional barrens, and by 61% (to 17.0%) in post-transitional barrens. These reductions in fertilization rates correspond to intermediate sensitivities of 0.49 ± 0.20 for a 50% reduction in the proportion of spawning adults and of 0.54 ± 0.14 for a 75% reduction.

DISCUSSION

Model assumptions and parameterization

The accuracy of our model predictions depends on both the accuracy of parameter estimates and the validity of assumptions. In this section, we consider the accuracy of certain estimates of model parameters, and we evaluate the validity of 2 important assumptions and whether they are equally valid for all subpopulations. This assessment will indicate the extent to which our predictions are accurate in an absolute or relative (i.e. among subpopulations) sense. Finally, we also consider the effect of omitting deeper-water populations from our models.

Sensitivity analysis indicates that the accuracy of the predictions of our egg spawning model depends largely on the accuracy of estimates of population parameters. We estimate these parameters based on several studies of populations of *Strongylocentrotus droebachiensis* along the Atlantic coast of Nova Scotia. Although absolute parameter values may vary over time and space within subpopulations, the rank order of those values among subpopulations is likely correct. Therefore, our models should accurately predict the relative numbers of eggs spawned by different subpopulations.

In an extension of our models of egg spawning, we estimate fertilization rates and the number of zygotes produced by subpopulations of *Strongylocentrotus droebachiensis* off the coast of Nova Scotia. The approach we use to approximate the proportion of spawning adults may overestimate that parameter. In nature, spawning of *S. droebachiensis* has only been observed outside the major spawning season when it involved a small percentage of the population (up to 15%, Keats et al. 1987; 5%, Pearse et al. 1988). Spawning by a low number of individuals has previously been reported for other sea urchins (Pennington 1985, Levitan 1988b), although 'mass spawning' has been observed in *S. pur*-

puratus, an unidentified sea urchin (Pennington 1985), and Evechinus chloroticus (Lamare & Stewart 1998). When we reduce the proportion of spawning adults to 0.17 and to 0.08 (i.e. to proportions similar to what has been observed in the field), estimated fertilization rates in the different subpopulations decrease by 2 to 50 % or by 21 to 61 %, respectively, suggesting that an overestimate of this parameter would have a marked influence on the estimated number of zygotes produced.

Our estimates of fertilization rates may be affected also by variability in the hydrodynamic regime. In the shallow subtidal zone, current velocity varies in space and time as different types of currents (tidal or wind driven) interact with one another. We estimate fertilization rates in the subpopulations based on field experiments conducted in a weak current (<5 cm s⁻¹; Wahle & Peckham 1999). If sea urchins spawn at stronger current speeds, fertilization rates are likely to decline because of increased gamete dilution rates (e.g. Pennington 1985, Denny & Shibata 1989, Levitan & Young 1995). Wahle & Peckham (1999) demonstrated a decline in fertilizations from 62 to 17% outside kelp cover as current speed increased ~2-fold, although this effect was noted only at a density of 36 spawners m⁻² (not at 2.25 m⁻²). Therefore, if sea urchins spawn in currents much greater than 5 cm s⁻¹, we may be overestimating fertilization rates, and consequently the number of zygotes produced. Because this overestimate is more likely to occur in grazing fronts (where the density is 37 spawners m⁻²) than in other subpopulations (1.3 to 8 spawners m⁻²), our estimated level of zygote production by individual subpopulations may be inaccurate in a relative (i.e. among subpopulations) sense. If this is the case, our estimated level of total zygote production also may be inaccurate in an absolute sense.

Our estimates of egg spawning and the contribution of different subpopulations to the overall zygote pool are limited to the shallow subtidal zone above 15 m depth. Although the abundance and size of Strongylocentrotus droebachiensis declines considerably at greater depths (Propp 1977, Logan et al. 1984, Martin et al. 1988), no information is available on the reproductive output or demography of these populations. Also, no measures exist on the extent of hard substratum below 15 m depth along the Atlantic coast of Nova Scotia. Therefore, we cannot evaluate the potential of deeperwater populations for egg spawning (or zygote production). While populations below 30 m are likely to have a low reproductive output due to a lack of macroalgal food, those at intermediate depths (15 to 30 m) may spawn a considerable number of eggs as their food supply is augmented by drift algae from the shallows. It is thus likely that we are underestimating total egg spawning in the subtidal zone, especially during the established kelp bed state when the reproductive potential of deeper-living populations may be enhanced by inputs of drift algae (Scheibling et al. 1999).

Our evaluation of assumptions and sensitivity analysis suggest that our estimates of egg spawning are likely to be accurate in a relative sense although absolute numbers may be inaccurate. Our estimates of fertilization rates, and thus levels of zygote production, in the different subpopulations are most influenced by potential variability in spawning synchrony and in hydrodynamic conditions. Total egg spawning and zygote production on a coastal scale may be underestimated due to the omission of sea urchins at depths >15 m. To improve model predictions, we need to obtain a better estimate of spawning synchrony, determine under which flow conditions Strongylocentrotus droebachiensis are most likely to occur and spawn, and collect reproductive and demographic data on sea urchin populations at depths >15 m.

Variation in the number of eggs spawned

The predicted number of eggs spawned per m² was influenced equally by changes in population density, and in the proportions of adults, females, body weight that is gonads, and gonad weight that is spawned (as indicated by sensitivity analysis). Sea urchins in grazing fronts release the largest number of eggs per unit area because extremely high densities of large females spawn a high proportion of their relatively large gonads. In transitional and post-transitional barrens, where females are less dense and smaller, and have a smaller gonad output than in grazing fronts, the predicted number of eggs spawned per unit area is about 1 order of magnitude lower than in fronts. In kelp beds, where relatively small females occur at very low densities, the predicted number of eggs spawned per unit area drops to about one-fifth of that spawned in barrens. In a study of variation in the gonad output of Strongylocentrotus droebachiensis at different depths (0 to 18 m), Keats et al. (1984) estimated that sea urchins in kelp beds produced a gamete biomass per unit area (12 g m^{-2}) that was similar to or less than that produced by sea urchins at higher densities in barrens at greater depths (9 to 49 g m⁻²).

The contribution of each subpopulation to the total number of eggs spawned on a coastal scale is determined largely by the extent of the subtidal zone that a given subpopulation occupies. As a result, sea urchins in grazing fronts, which spawn the largest number of eggs per unit area but occupy <1% of the rocky subtidal zone, generally spawn fewer eggs in total than sea urchins in other subpopulations. For sea urchins in kelp beds and transitional barrens, egg spawning

decreases or increases respectively, as kelp beds are destroyed by grazing fronts and replaced with barrens during the transition state. The total number of eggs spawned on a coastal scale increases approximately 6-fold from the established kelp bed state to a peak in the late transition state before sea urchins in grazing fronts disperse. In post-transitional barrens, 4 times more eggs are spawned than in the former kelp beds.

Although numerous studies have determined the number of eggs produced by free-spawners either per individual (e.g. Thompson 1979, Scheibling & Lawrence 1982, Kettle & Lucas 1987, McShane et al. 1996) or per unit area (Coma et al. 1995, Coma & Lasker 1997), few attempts have been made to estimate the total number of eggs produced by an entire population (Coma & Lasker 1997). Little information exists on variation in egg spawning among populations of different size or demography. Our model predictions suggest that both the size and demography, i.e. structure of a population, must be taken into account when estimating the number of eggs that it spawns. Because egg spawning in combination with fertilization success determines zygote production, it is crucial to establish how egg spawning varies temporally and spatially when assessing how populations may vary in their contribution to the total zygote pool.

Variation in fertilization rate

Sensitivity analysis indicates that the estimated fertilization rate is similarly sensitive to changes in population density, in the proportion of adults, and in the proportion of adults that spawn during a spawning event. We assume that the proportion of spawning adults is identical in all subpopulations. The other 2 parameters, however, vary markedly among subpopulations of Strongylocentrotus droebachiensis (Table 3), resulting in a 4-fold difference in estimated fertilization rates. Fertilization rate is highest (62%) for sea urchins in grazing fronts where adults occur at extremely high densities (i.e. at small distances between spawners). In barrens, where adult densities are lower, fertilization rates are estimated to be intermediate (36 and 43% in transitional and post-transitional barrens, respectively). In kelp beds, the low density of adults causes the fertilization rate to drop to the lowest level (15%).

Variation in zygote production and sea urchin population dynamics

Because estimated fertilization rates differ between subpopulations by up to a factor of 4 (between sea urchins in grazing fronts and in kelp beds), estimated variation in zygote production per unit area among subpopulations of Strongylocentrotus droebachiensis is more pronounced than variation in the number of eggs spawned. The contribution of each subpopulation to total zygote production on a coastal scale is, however, still determined largely by the extent of the subtidal zone that a given subpopulation occupies. As a result, the general patterns of egg spawning and zygote production on a coastal scale are similar. Although sea urchins in grazing fronts still produce generally fewer zygotes than sea urchins in other subpopulations, a high fertilization rate enables them to make a significantly larger contribution to the overall zygote pool than to the overall egg pool (11 to 44 vs 7 to 22%). Because of the large variation in fertilization rate, there is an approximately 16-fold increase in total zygote production from the established kelp bed state to the late transition state, compared to an approximately 6-fold increase in egg spawning during the same interval.

To assess the importance of variation in zygote production on the population dynamics of Strongylocentrotus droebachiensis along the Atlantic coast of Nova Scotia, we compare our estimate of zygote production with natural variation in settlement. On a coastal scale, we estimate that total zygote production in the shallow subtidal zone differs 16-fold among community states that include adult sea urchins. Because no zygotes are produced in the shallow subtidal zone for at least 2 yr after a sea urchin mass mortality, variation in total zygote production is even greater if we include the developing kelp bed state (from 0 to $\sim 10^{15}$ m⁻²). However, settlement can occur shortly after a sea urchin die-off (Miller 1985, Scheibling 1986, Raymond & Scheibling 1987, Scheibling & Raymond 1990) and can vary interannually by more than an order of magnitude during the transition from kelp beds to barrens (Balch & Scheibling 2000), when we predict a more gradual increase. This discrepancy between the temporal patterns of estimated zygote production and observed settlement rates suggests that settlement of S. droebachiensis in the shallow subtidal zone is not primarily determined by zygote production within that zone. Other important factors may include advective transport of larvae to and from the region (Scheltema 1986, Shanks 1995), and sea temperature which regulates the rate of larval development and hence survival in the plankton (Hart & Scheibling 1988). Nevertheless, the estimated large increase in zygote production during the transition state may enhance settlement and thus sustain, or even accelerate, a sea urchin outbreak.

A clearer resolution of the importance of variation in egg spawning and fertilization rate, and consequently variation in zygote production, to population dynamics of *Strongylocentrotus droebachiensis*, relative to other

biological and physical factors, requires further study. For example, longitudinal studies spanning different community states may elucidate the relationship between patterns of zygote production and the frequency and magnitude of settlement events. Genetic analysis of larvae and settlers also may be helpful in determining the origins of larvae settling along this coast (Medeiros-Bergen et al. 1995, Palumbi 1995). In particular, investigation of deep-water populations is needed to assess their contribution to the larval pool.

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