

Mortality rate estimation for eelgrass *Zostera marina* (Potamogetonaceae) using projections from Leslie matrices

Olga Flores Uzeta, Elena Solana Arellano & Héctor Echavarría Heras

Departamento de Ecología Marina, Centro de Investigación Científica y Educación Superior de Ensenada, Ensenada, Baja California México, P.O. Box 430222, San Diego, CA. 92143-0222, USA. Fax: (646) 175 05 00; oflores@cicese.mx; esolana@cicese.mx; hechavar@cicese.mx

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Abstract: The main goal of this study is to provide estimations of mean mortality rate of vegetative shoots of the seagrass *Zostera marina* in a meadow near Ensenada Baja California, using a technique that minimizes destructive sampling. Using cohorts and Leslie matrices, three life tables were constructed, each representing a season within the period of monthly sampling (April 1999 to April 2000). Ages for the cohorts were established in terms of Plastochrone Interval (PI). The matrices were projected through time to estimate the mean total number of individuals at time t , $n(t)$ as well as mortality. We found no statistical differences between observed and predicted mean values for these variables ($t=-0.11$, $p=0.92$ for $n(t)$ and $t=0.69$, $p=0.5$ for mean rate of mortality). We found high correlation coefficient values between observed and projected values for monthly number of individuals ($r=0.70$, $p=0.007$) and monthly mortality rates ($r=0.81$, $p=0.001$). If at a certain time t a sudden environmental change occurs, and as long as the perturbation does not provoke the killing of all the individuals of a given age i for $0 \leq i \leq x - 1$, there will be a prevailing number of individuals of age or stage x at a time $t+1$. This non-destructive technique reduces the number of field visits and samples needed for the demographic analysis of *Z. marina*, and therefore decreases the disturbance caused by researches to the ecosystem. Rev. Biol. Trop. 56 (3): 1015-1022. Epub 2008 September 30.

Key words: matrix models, demography, *Zostera marina*, mortality rates.

Seagrass meadows are very important in the ecology of coastal waters (Mann 1982), but are very susceptible to both anthropogenic and climatic perturbations. This means that mortality and recruitment are going to be responsible for changes in density and shoot coverage (Olesen and Sand-Jensen 1994) and therefore the study of these processes is crucial to understand the demography of seagrasses. Since these plants are modular, with continuous branching and expansion of rhizomes (Gallegos 1995), it is reasonable to assume that recruitment is constant over time. Moreover Solana-Arellano (2004) showed through an analysis of variance, that recruitment rates for *Zostera marina* is constant ($F=1.4$, $p=0.48$) at our study site.

Even though seagrasses are widely distributed in temperate waters throughout the northern hemisphere (den Hartog 1970), shoots demography has not been studied is for terrestrial plants for which the general patterns of shoot density regulation via shoot natality or shoot mortality are known (Lovett 1981, Schmid and Harper 1985). Moreover demographic techniques to explore population have been applied to seagrasses only to evaluate the potential of a population in terms of density increments or decrements at the level of the whole meadow dynamics (Patriquin 1973, Durako and Moffler 1987, Cox and Tomlinson 1988, Duarte and Sand-Jensen 1990, Duarte *et al.* 1994). In addition, the use of these techniques has been restricted to only three sea-

grass species; *Posidonia oceanica* (Pergent and Pergent-Martini 1990), *Thalassia testudinum* (Patriquin 1973, Gallegos *et al.* 1993, Durako 1994) *Cymodocea nodosa* (Duarte and Sand-Jensen 1990, Pérez *et al.* 1994, Vidondo *et al.* 1997).

Specifically for *C. nodosa*, Pérez *et al.* (1994) gave an estimate of mean mortality rates (in logarithmic units) from the exponential reduction of shoots density grouped in consecutive annual cohorts structured by age. This relatively easy method, besides being destructive, gives no details of the mathematical assumptions necessary to obtain mortality rates for *C. nodosa*. On the other hand, the powerful reconstructive methods proposed by Pergent and Pergent-Martini (1990) and Duarte *et al.* (1994), which are non-destructive, are only suitable for long-lived seagrasses. For *Z. marina*, the study of Olesen and Sand-Jensen (1994) is the first work based on cohorts structured by age. They describe the relationship between growth and age of a shoot in terms of plastochrones, seasonal production, above ground biomass, and shoot density regulation through recruitment and mortality. Solana-Arellano (2004) used cohorts and the Pérez *et al.* (1994) method to describe the demography of a Mexican *Z. marina* meadow.

To study shoot demography, it is necessary to construct life history tables, defined by Ebert (1999), as the schedule of probabilities of things that may happen to individuals with particular attributes. The most general choice for attribute is age. We then projected these life tables to visualize trends in population demography, given parameters of birth and death. The projection of a life table matrix for a population was defined by Keyfitz (1972) as an assay that describes what is going to happen with the population density given certain parameters. According with Chapman (1986), life tables can be applied to plants and even marine algae if individuals can be differentiated.

The purpose of this work is to compare a series of observed mortality rates with projected rates based on only one or a few field samples. These rate estimations were obtained

through the projection of the Leslie matrix life table with entries given in terms of survival and fecundity rates. A reasonable correspondence of observed and predicted rates would allow us calculate some characteristic of the population without the use of destructive samplings. Besides contributing to the knowledge of demographic strategies, our findings would justify drastically reduce the number of field samplings needed per year.

MATERIALS AND METHODS

Field, laboratory and statistical methods:

data were generated biweekly from February 1999 to April 2000 in a *Z. marina* meadow. At Punta Banda Estuary on the Pacific coast of Baja California, México (31°40' - 31°48' N, and 116°37' -116°40' W). For a complete description of the sample site see Solana-Arellano (2004). On each sampling date, all shoots on 20 randomly selected rhizomes were marked using the technique of Kentula and McIntire (1986). After fifteen days, we harvested the previously marked shoots, and marked a new set. Samples were taken to the laboratory where the following measurements were made: number of leaves per shoot, number shoots per meter of rhizome, number of new shoots and leaves. Age of each shoot and biweekly Plastochrone Interval (PI) were calculated. The projections of the Leslie matrices were developed using the program Maple V (1998) for symbolic manipulation. The STATISTICA (1999) package was used for the data analysis.

Data management: the mean annual Plastochrone Interval (PI) was calculated using the criteria Jacobs (1979). The PI is defined as the time interval between the initiation of two successive leaves on one shoot. For *Z. marina*, Jacobs (1979) introduced a formula for the calculation of the PI which consists of multiplying the total number of marked shoots by times days of exposure and dividing by the number of new leaves on marked shoots. The age of each shoot, in Plastochrone Intervals, was calculated adding the number of standing

leaves plus leaf scars from shed leaves (Duarte and Sand-Jensen 1990). Monthly cohorts and the life tables were constructed using 1999 data as a preliminary analysis. Based on the result of this analysis, (see result section), we constructed three life tables: Life table I, with data gathered on April 1999 and projecting the interval of May-August 1999, Life table II, with data based on August 1999 and projecting the interval September-December 1999 and Life table III, with data of December 1999 that projected the time interval of January-April 2000. With each projection, mean monthly mortality rates were calculated and compared statistically with observed values.

Theoretical methods: we define a cohort as a collection of individuals that share some trait or a group of traits for purposes of study. Then, our collection of shoots from a particular sampling date classified by age can be considered as a cohort (Harper 1994, Ebert 1999). The groups of statistics that conform the life tables were defined as in Ebert (1999) so that n_x denotes is the number of individuals of age x . By means of m_x we represent the number of offspring produced by an individual that enters the age x and survives the entire period x to $x+1$. A correction has to be made for m_x to adjust fir changes of fecundity during the time interval. This fecundity can be concentrated at one point during a time interval as a pulse, or incorporated continuously over the interval t , to $t + 1$. Since our intervals are of different sizes then a second correction (m_x^*) has to be made to the value of m_x that will be concentrated at x (Ebert 1999). The symbol l_x stands for the probability that an individual (age 0) is alive at the beginning of age x . Finally p_x represents the probability that an individual of age x survives one time period and so is conditional on an individual being alive at the beginning of the time interval. Then, in terms of number of individuals we can define:

$$l_x = \frac{n_x}{n_0} \quad (1)$$

$$p_x = \frac{l_x + l_{x+1}}{l_{x-1} + l_x} \quad (2)$$

$$p_0 = \frac{l_0 + l_1}{2} \quad (3)$$

and

$$m_x^* = \frac{m_{x-1} + p_x m_x}{2} \quad (4)$$

The recruitment rate was calculated as:

$$b = \frac{\text{number of new born at time } t}{\text{number of survivors from } t - 1 \text{ to } t} \quad (5)$$

Similarly, survival rate is:

$$s = \frac{\text{number of survivors from } t - 1 \text{ to } t}{\text{number individuals at time } t - 1} \quad (6)$$

Then we can obtain mortality rate defined as

$$q = 1 - s \quad (7)$$

Now, let A be the life table matrix for our population, so A will contain information about the above statistics and will be projected in time to produce the number of individuals of age x at time $t+1$. That is,

$$n_x(t+1) = A n_x(t) \quad (8)$$

Or, in matrix form, we have.

$$\begin{bmatrix} n_0(t+1) \\ n_1(t+1) \\ \cdot \\ \cdot \\ n_n(t+1) \end{bmatrix} = \begin{bmatrix} m_1 & m_2 & \dots & m_n \\ p_1 & 0 & \dots & 0 \\ \cdot & & & \\ \cdot & & & \\ 0 & p_n & \dots & 0 \end{bmatrix} \begin{bmatrix} n_0(t) \\ n_1(t) \\ \cdot \\ \cdot \\ n_n(t) \end{bmatrix}$$

Matrix A on the right side of equation (8) is known as a Leslie matrix (Caswell 2001). With information on the above equation we can obtain the projected number of individuals and with equation (6) and (7) the mean monthly mortality rates.

RESULTS

We sampled 820 rhizomes with a total of 4 014 shoots. The mean annual PI was 8.2 with an standard error of ± 0.26 . Using the 1999 data, we made monthly projections and compared all projected densities, finding that projections of April and August were statistically different from the rest of the projected months (ANOVA, $p=0.02$). Based on these results and taking into account an assumed seasonal growth pattern for *Z. marina*, (McRoy 1970, Jacobs 1979, Solana-Arellano *et al.* 1997) we defined three seasons, winter-spring (January-April); Summer-Autumn (May-August) and Autumn-Winter (September-December). These three seasons were projected using life tables I, II and III as described in the data management section and we found a good correspondence between projected and observed values for the total number of individuals at time t ($n(t)$) (Fig. 1). Moreover we found no statistical differences between projected and observed values for $n(t)$ ($t=-0.11$, $p=0.92$). Using this information and equation (6), we estimated the mean

monthly rates of mortality and (Fig. 2) shows the projected and observed rates for the sampled period. We performed a Student's t test, and found no statistical differences between mean projected and mean observed values for mortality rate ($t=0.69$, $p=0.5$). The correlation between projected and observed rates was 0.81 ($p=0.001$). This along with Fig. 2 showed that the projections of life tables gave a reliable prediction of the aforementioned rates for *Z. marina*.

DISCUSSION

Competitive species regulate their population sizes through shoot recruitment (Schmid and Harper 1985). Furthermore, some authors (Miyanishi *et al.* 1979, Duarte and Kalf 1987, Hutchings and Mogie 1990) argue that seagrasses and in general clonal plants have a vigorous vegetative growth that results in vegetative expansion of clones. This is considered density-independent mortality which is a consequence of their growth strategy and not competition for substratum. On the other

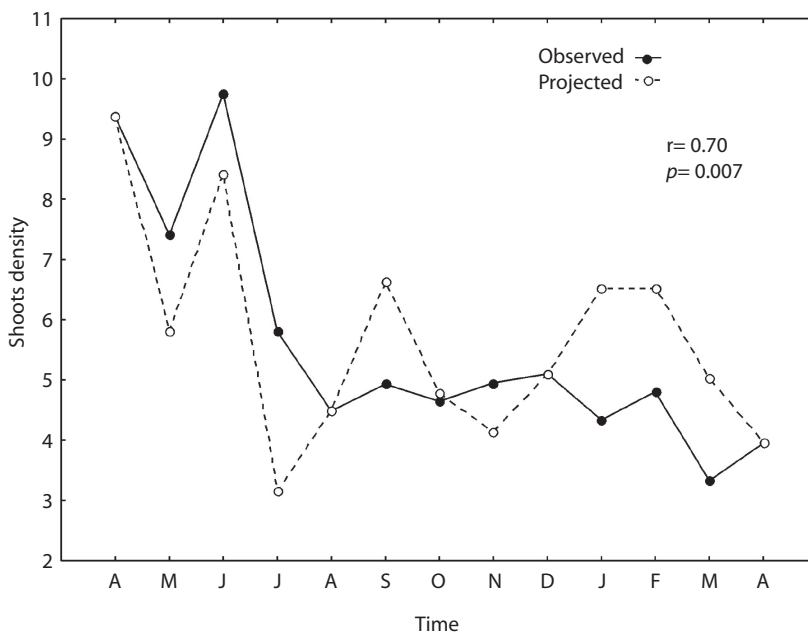


Fig. 1. Projected and observed values for shoots density at time t ($n(t)$).

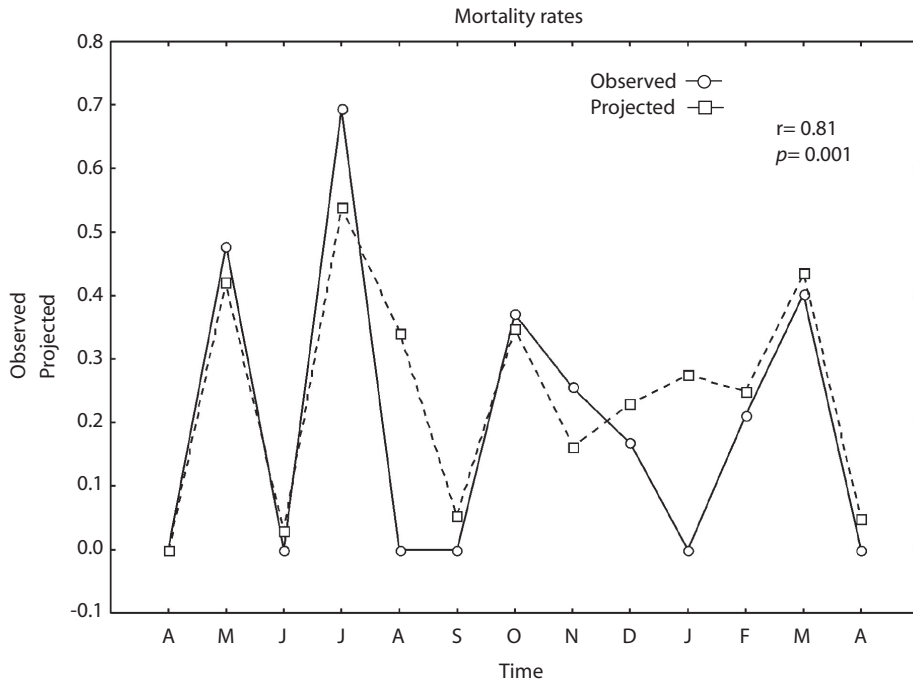


Fig. 2. Projected and observed values for mortality rate for *Z. marina*.

hand, plants with an opportunistic growth strategy tend to control $n_x(t)$ through mortality rate (Lovett 1981, Schmid and Harper 1985). Therefore, the study of mortality rates in these ecosystems is very important.

Seagrass meadows are very vulnerable to sudden climatic changes and anthropogenic perturbation (Short and Neckless 1999, Echavarría-Heras *et al.* 2006). In fact, Olesen and Sand-Jensen (1994) stated that shoot density is influenced by environmental factors and/or internal growth rhythms in seagrasses. This and the expensive and tedious marking techniques and lengthy laboratory procedures make necessary the development of rapid and non-destructive sampling techniques. The projection of a few seasonal life tables proved to be an effective tool for demographic analysis.

Within reasonable limits, whenever a population is ecologically stable, any sudden perturbation would induce a response allowing the population to prevail. Parameter identification and statistical contrastation of predictions

versus observed dynamics are basic steps of a model selection procedure. Once suitability has been found to be satisfactory, the assumptions that sustain the model are expected to be reasonably close to the phenomenological properties of the real system. Particularly a matrix population model as the one invoked in the present work is able to predict the persistence of the population after a sequence of adaptive steps characterized by delayed responses induced by environmental perturbations. Moreover the demographic parameters are defined in such a way that their values at a given time depend recursively on former values. For instance, l_x can be defined as the product of p_i values which stand for the probability that an individual of age i survives one period of time, for $0 \leq i \leq x - 1$. Hence l_x the probability that a given individual is alive at age x will take a value different from zero if and only if none of the p_i values vanishes. If at a certain time t a sudden environmental change occurs, then as long as the perturbation does

not provoke the killing of all the individuals of a given age i for $0 \leq i \leq x - 1$, then at a time $t+1$ there will be a prevailing number of individuals of age or stage x . That is, a sample taken in a given interval is demographically determined by the past history of the population up to that time. Hence the identification of the addressed model permitted to obtain reasonable values for the invoked demographic parameters. In other words, population stability and model suitability explain why our projection method produces consistent predictions.

Clonal plants respond to perturbations or environmental variability by modifying the morphological arrangement of their modules (Franco 1986). Furthermore, the utilization of plant resources depends on their ability to make physiological adjustments (Kays and Harper 1974). Other authors argued that both perturbations and environmental changes determine the expansion and growth patterns of modular populations but that the degree of adjustment is a characteristic of each species. Nevertheless, the adjustments are not instantaneous, in fact they depend on the different growth strategies with morphological extremes denoted by Lovett-Doust (1981) as guerilla and flange. The plasticity of the plant to adopt each of these strategies requires an amount of time. Solana-Arellano 2004 found that *Z. marina* adjusted its growth pattern from guerilla to flange and vice versa depending on the environment but this adjustment is not instantaneous. Therefore, even on sudden environmental changes the meadow will adjust after period of time. Hence the surviving individual will determine the demographic parameters; this implies that just two or three sampling visits per year on a stable population will provide a good basis for projection. On the other hand, suppose we have a projection based on data unaffected by sudden changes; assume that this has been proved to produce reasonable estimations. If a sudden non-catastrophic event occurs, then a sample taken after that event will give us estimations of the effect of such event on the meadow. This will be a crucial output for monitoring and management of the ecosystem.

The high correlation coefficient (0.81) between observed and projected mortality rates values through time showed that there is a good correspondence between observed and projected rates. Therefore we conclude that the contribution of a non-destructive technique for demographic analysis of a *Z. marina* population reduces number of field visits and quantity of samples and therefore the disturbances by researches to the ecosystem and is then an important practical contribution.

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RESUMEN

El propósito principal de este estudio es el de proveer estimaciones de tasas promedio de mortalidad de tallos vegetativos de *Zostera marina* en una pradera cercana a Ensenada Baja California, utilizando una técnica que minimiza los muestreos destructivos para estos pastos marinos. Mediante la utilización de cohortes y matrices de Leslie, se construyeron tres tablas de vida, cada una representando a una estación dentro de período anual de muestreos mensuales (Abril 1999 a Abril 2000). Las edades de los cohortes fueron estimadas en términos de Intervalos de Plastocrono (IP). Las matrices de Leslie fueron proyectadas a través del tiempo para estimar el número total de individuos a un tiempo t , $n(t)$ así como las tasas de mortalidad. No se encontraron diferencias significativas entre los valores medios observados y proyectados para estas variables ($t=0.11$, $p=0.92$ para $n(t)$ y $t=0.69$, $p=0.5$ para la tasa media de mortalidad). Se encontraron altas correlaciones los valores observados y proyectados tanto en el número de individuos ($r=0.70$, $p=0.007$) como las tasas mensuales de mortalidad ($r=0.81$, $p=0.001$). Si a cierto tiempo t ocurre un cambio ambiental repentino, de tal manera que la perturbación no provoque la muerte de todos los individuos de una edad i para $0 \leq i \leq x-1$, prevalecerá un número de individuos de edad o estadio x al tiempo $t+1$. Esta técnica no destructiva reduce el número de visitas al campo y de muestras necesarias para un análisis demográfico de *Z. marina* y por lo tanto decrece el disturbio causado al ecosistema.

Palabras clave: modelos de matrices, demografía, *Zostera marina*, tasa de mortalidad.

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