THE NORTHERN ANCHOVY SPAWNING BIOMASS FOR THE 1982-83 CALIFORNIA FISHING SEASON

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ABSTRACT
The 1982 equivalent larval census estimate of the spawning biomass of the northern anchovy (Engraulis mordax) central subpopulation is 1,866,000 MT (2,060,000 short tons). This estimate is based on data gathered by an egg production survey conducted from January 18 to March 9, 1982. The abundance of larvae is projected from the daily production of eggs and the subsequent mortality of eggs and larvae. This equivalent larval census estimate will be the basis of the anchovy fishery optimum yield (OY) for 1982-83.

The egg production method estimate of anchovy spawning biomass is 378,000 MT (417,000 short tons). This is based on a production of $12.30 \times 10^{12}$ eggs/day and a population (males and females) fecundity of $3.253 \times 10^7$ eggs/day/MT. The standard error of the egg production method estimate is 97,200 MT, for a coefficient of variation of 25.7%.

In 1982 the central subpopulation of northern anchovy was geographically distributed in the inshore portion of the Southern California Bight, from the Santa Barbara Channel Islands to northern Baja California. A smaller group was detected between Monterey and San Francisco Bays.

INTRODUCTION
The biomass of the central subpopulation of northern anchovy (Engraulis mordax) has been assessed periodically using three independent methods: larval census (Smith 1972; Stauffer and Parker 1980; Stauffer 1980; Stauffer and Picquelle 1981); sonar mapping (Smith 1970; Mais 1974; Hewitt et al. 1976); and egg production (Parker 1980; Stauffer and Picquelle). The anchovy management plan adopted by the Pacific Fishery Management Council (PFMC 1978) specifies that harvest quotas will be established by an optimum yield formula based on annual estimates of spawning biomass. Because of technical complexities in converting acoustic returns to fish biomass, the sonar mapping method is most useful for describing the disposition of adult schools. The egg production method is an improvement over the larval census method because it does not require multiple surveys and because it measures and incorporates variability in adult reproductive output. However, the optimum yield (OY) formula was developed based on larval census biomass estimates. The alternative estimation methods mentioned above produce estimates that consistently differ in magnitude from larval census estimates. Consequently, use of one of these estimates would systematically alter the optimum yields calculated from the larval-census-based formula. In order to avoid this
problem, the PFMC has interpreted the OY formula to require a "larval census equivalent" spawning biomass estimate.

The Scientific and Statistical Committee of the PFMC reviewed egg production and larval census surveys conducted in 1979 and 1980, and recommended that replicate egg production surveys, concurrent with a larval census survey, be conducted in 1981. These were successfully accomplished, and the harvest quotas continued to be established using larval census estimates of biomass. In 1982 the National Marine Fisheries Service, in cooperation with the California Department of Fish and Game, conducted only an egg production survey. This report describes the survey results, the egg production estimate of biomass, and the equivalent larval census estimate of biomass.

The egg production method defines the spawning biomass as the quotient of the daily production of eggs in the sea and the daily fecundity (per ton of spawners) of the population (Parker 1980). The larval census method defines the spawning biomass as proportional to the average standing stock of larvae summed over four quarters of the year, and assumes constant reproductive output (per ton of spawners) and constant survival of the young. The proportionality constant was defined from a regression of sardine biomass on sardine larvae and assumes a relative fecundity between anchovy and sardine (Smith 1972).

No larval census was conducted in 1982; however, an equivalent larval census can be estimated by measuring larval mortality and projecting the number of larvae resulting from the measured egg production.

DESCRIPTION OF SURVEY

The 1982 egg production survey for the central subpopulation of the northern anchovy was conducted on board NOAA ship David Starr Jordan and F/V Oregon Beaver during the period January 18 through March 9, 1982. (For detailed descriptions of operations, see the respective cruise reports on file at the Southwest Fisheries Center, La Jolla, California, and California Department of Fish and Game, Long Beach, California.) A total of 992 egg samples was obtained on the Jordan using a small-mesh plankton net retrieved vertically from 70 m (Figure 1); adults were sampled on the Beaver and the Jordan with 187 midwater trawls (Figure 2); and the disposition of adult schools was described by sonar operated aboard the Beaver (Figure 3). In addition, the larvae were sampled with the 505-micrometer mesh bongo plankton net used for the larval census method. These samples are used to determine the posthatch survival of the larvae. Sea temperature, salinity, and weather observations were also recorded.
Anchovy eggs were distributed along the northern Baja California coast and in the Southern California Bight as far north as the Santa Barbara Channel Islands (Figure 1). As in previous years, the geographic pattern of spawning was correlated with the pattern of surface temperature isotherms (Lasker et al. 1981). Spawning did not occur in water colder than 14°C south of Point Conception, although this is not considered a lethal temperature. Cold water was observed as a large plume extending southeastward from Point Conception into the Southern California Bight and immediately adjacent to the northern Baja California coastline (Figure 1). A much smaller area of spawning was sampled between Monterey and San Francisco Bays in 11-12°C water.

The geographic distribution of positive trawl samples agrees with the distribution of eggs (Figure 2). Adult fish were caught in the Southern California Bight, along the northern Baja California coast, and between Monterey and San Francisco. Adults were not caught in trawls taken in the Santa Barbara Channel and immediately adjacent to the northern Baja California coastline.

The geographic distribution of fish schools as detected by sonar agrees with the distribution of eggs and positive trawls (Figure 3). The area of heaviest anchovy concentration extends along the northern Baja California coastline (approximately 50 km offshore) and into the inshore portion of the Southern California Bight as far north as the Santa Barbara Channel Islands. Two smaller areas of heavy concentration were mapped in the offshore portion of the Southern California Bight.

In summary, northern anchovy, in the spring of 1982, were distributed continuously in the Southern California Bight from the Santa Barbara Channel Islands south to Cape Colnett. A second group was distributed between Monterey and San Francisco bays. In both groups fish were spawning, thus facilitating the use of reproductive surveys to estimate adult biomass.

**EGG PRODUCTION ESTIMATE**

**Estimation Equation**

The egg production estimate of anchovy spawning biomass, derived by Parker (1980) and modified by Stauffer and Picquelle (1980), is

\[
B = P_o A k W R F S
\]  

where \(B\) = spawning biomass (MT), \(P_o\) = daily egg production, number of eggs produced per 0.05 meter\(^2\), \(W\) = average weight of mature females (grams), \(R\) = sex ratio, fraction of population that are female, by weight (grams), \(F\) = batch fecundity, number of eggs spawned per mature female per batch, \(S\) = fraction of mature females spawning per day, \(A\) = total area of survey (0.05 meter\(^2\) units), \(k\) = conversion factor for grams to metric tons.

An approximate sample variance for the egg production spawning biomass estimator, derived from the delta method (Seber 1973), is a function of the sample variance and covariance of the parameters

\[
\text{Var}(B) = B^2 \times \left\{ \frac{\text{Var}(P_o)}{P_o^2} + \frac{\text{Var}(W)}{W^2} + \frac{\text{Var}(R)}{R^2} + \frac{\text{Var}(F)}{F^2} + \frac{\text{Var}(S)}{S^2} + \frac{\text{Cov}(P_o,W)}{P_o W} + \frac{\text{Cov}(P_o,R)}{P_o R} + \frac{\text{Cov}(P_o,F)}{P_o F} + \frac{\text{Cov}(P_o,S)}{P_o S} + \frac{\text{Cov}(W,R)}{W R} + \frac{\text{Cov}(W,F)}{W F} + \frac{\text{Cov}(W,S)}{W S} + \frac{\text{Cov}(R,F)}{R F} + \frac{\text{Cov}(R,S)}{R S} + \frac{\text{Cov}(F,S)}{F S} \right\}. \tag{2}
\]

**Daily Production of Eggs in the Sea**

The parameter \(P_o\), the daily production of eggs in the sea, is the number of eggs spawned per night, per unit area, averaged over the range and duration of the
survey. An ichthyoplankton survey is used to sample anchovy eggs to provide data on the density of the eggs by age. An exponential mortality model is then fit to the data, and the time-zero intercept of the fitted function is the estimate of egg production.

The sampling design of the survey is two-stage sampling with post-survey stratification. The first stage is a systematic sample of block areas. The total area of the survey is divided into 4 x 20- (or 4 x 10-) nm blocks. The second stage is the selection of a 0.05-m² sampling unit in the center of the block (Stauffer and Picquelle). This sampling design assumes that the distribution of eggs within one block is independent of the distribution within adjacent blocks. This assumption is based on experimental data showing that 4 miles is sufficient distance to ensure a negligible autocorrelation (P. Smith, pers. comm.). Advantages of this sampling plan are convenient and efficient use of ship time, even coverage of the total sample area, and improved precision of abundance estimates by maximizing the heterogeneity between adjacent sampling units (Jessen 1978).

Because of time and budget constraints, the sampling intensity was decreased in regions where fewer eggs were expected to be found. In the 1981 surveys and the 1980 survey, the majority of the anchovy population was found in the Southern California Bight; a narrow band of eggs was typically found off Baja California; and very few eggs were collected off the central California coast. Therefore, the sampling fraction in the bight is increased to one sample per 4 x 10-nm block; elsewhere there is one sample per 4 x 20-nm block. To compensate for the uneven sampling intensity, each station is assigned a weight, $w_{ij}$, that is proportional to the relative area that the $j$th station represents in the $i$th stratum.

The total survey area is divided into two strata in order to reduce the variability about the egg abundance estimates. The geographic area of the survey is specified without knowing the actual area that the anchovy stock is currently occupying. Therefore, a portion of the surveyed area is beyond the range of the stock, contributing a large number of zero stations, which potentially inflates variance estimates. After the data are examined, a boundary for the current anchovy habitat can be drawn. The boundary is determined by following each line seaward until the last positive tow was taken. Thus all stations seaward of the boundary are zero, and all positive stations are shoreward of the boundary, along with many imbedded zero stations. The area within the boundary is allocated to stratum 1, and the area outside the boundary is put into stratum 0.

Because the strata can be defined only after the data are collected, this technique is called post-survey stratification. The area of each stratum, $A_i$, is not predetermined, so $A_i$ is a random variable, and all variance estimates must be adjusted to include this source of variation (Jessen 1978).

The anchovy eggs from each sample are counted and staged according to the degree of embryonic development. The eggs are then classified into one-day age intervals based on the time of collection, surface water temperature, developmental stage, and laboratory-determined development rates. The ages are then further refined by adding the portion of a day that has elapsed between 2200 (the assumed time of spawning) and the time of collection. This aging procedure is facilitated by the fact that eggs in a particular sample are either spawned on the same evening, or are separated by one-day increments, thus making it easier to separate the modes of egg abundance over stages into one-day age groups. Sea-surface temperatures for the majority of the cruise ranged between 13° and 17°C with an egg-weighted average of 14.9°C. At this temperature, eggs begin to hatch after 2.91 days. Eggs less than 2.67 days old were used in the analysis.

The daily production of eggs, $P_{ik}$, and its variance are then estimated by regressing the counts of eggs on their age, using the exponential mortality model:

$$P_{ik} = P_{0k} e^{-Z_{ik}} + \epsilon_{ijk}$$

where $P_{ik}$ is the number of eggs in the $i$th day age category from the $j$th station in the $i$th stratum; $t_{ik}$ is the age in days measured as the elapsed time from the time of spawn for the $i$th day age category eggs to the time of sampling of the $j$th station in the $i$th stratum; $P_{0k}$ is the daily production of eggs per unit area (0.05 m²) in stratum $i$; $Z$ is the daily rate of instantaneous egg mortality; and $\epsilon_{ijk}$ is the additive error term. This model assumes that all eggs are spawned and fertilized each day at time 2200, and eggs have a constant positive rate of instantaneous mortality.

The mortality function (equation [3]) is fit to the data by regressing the aged egg counts $\{P_{ik}\}$ on their age $\{t_{jk}\}$ for stratum 1 with a weighted nonlinear
Figure 4  Mean egg production per 12-hour interval for stratum 1 plotted against age in days. The exponential mortality model was estimated by regressing the individual egg counts on age.

least squares routine using a pseudo-Gauss-Newton algorithm (Dixon and Brown 1979). The individual weighting factors are the station area weights, \( w_i \). The resulting estimate of \( P_o \) is 7.40 eggs per day per 0.05 m², and \( Z \) is 0.158 per day with variances 1.25 and 0.0110, respectively. A plot of the mean egg production per 12-hour interval and the estimated regression for stratum 1 are shown in Figure 4.

The final stratified estimate of \( P_o \) is the weighted average of the two strata where the strata weights \( u_i \) are proportional to \( A_i \), the area of the \( i \)th stratum. \( P_{oi} \) for stratum 0 is zero by definition, thus:

\[
\hat{u}_i = \frac{A_i}{A_0 + A_i},
\]

\[
\hat{P}_o = \hat{u}_i P_{oi} + u_0 P_{o0} = \hat{u}_i P_{oi}, \tag{4}
\]

and the variance, adjusted for postsurvey stratification (Jessen 1978), is

\[
\text{Var}(\hat{P}_o) = \left( 1 + \frac{1}{n} \right) \left[ \sum_{i=0}^{2} u_i \left( \text{Var}(\hat{P}_{oi}) \right) \right] \tag{5}
\]

where \( n \) is the total number of observations = 992; \( \text{Var}(\hat{P}_{oi}) = 1.25 \) is estimated from the regression (3) for stratum 1, and; \( \text{Var}(\hat{P}_{o0}) = 0 \) by definition.

The stratified estimate of \( P_o \) is 3.023 eggs per day per 0.05 m² for the entire survey area, with an estimated variance of 0.5119 and coefficient of variation of 23.67%. This estimate of \( P_o \) applies over the 59,299-nm² area of the survey (\( A = 4.068 \times 10^{12} \) area units of 0.05 m² each).

**Adult Parameters \( W, F, S, \) and \( R \)**

The parameters \( W, F, S, \) and \( R \) are estimated from samples of adult anchovies collected by the midwater trawl survey. The sampling design consists of three stages: (1) placement of trawl stations, (2) trawl catch, and (3) subsample of fish. The statistical technique of judgment sampling is used in selecting the location for the trawls in order to achieve a high proportion of positive trawls. Station locations were selected where concentrations of anchovies were detected by the occurrence of anchovy eggs or larvae in the plankton samples, and the presence of apparent schools on the sonar (Figures 1, 2, and 3). Because more stations are selected where heavy concentrations of anchovies are suspected, the sample design follows the precepts of probability sampling (Cochran 1963).

One trawl is taken at each selected location, and the trawl catch is assumed to be a random sample of fish at the station. Each station is given equal weight by subsampling an equal number of fish from each trawl, \( m^* \). This is appropriate if the probability of choosing a station is exactly proportional to the number of anchovies at the station; however, these probabilities are impossible to enumerate, so that the actual sampling design is only an approximation of the ideal design (judgment sampling rather than probability sampling). If the trawl catch size were a good measure of the abundance of fish at the station, this information could be used to improve the approximation of the ideal sample design. However, trawl catch size and abundance of fish are unrelated for anchovies; catch size depends more on the depth of the school, the avoidance of the net by the fish, and luck. This conclusion is supported by the historical lack of agreement between trawl catch size and the factors that indicate concentrations of anchovies (sonar detection, eggs and larvae in plankton samples).

Even though equal subsample sizes are attempted, they are not always attainable: the catch may have few mature females, or the catch may be extremely small. In these cases the stations should receive less weight to compensate for the error in judgment sampling; if there are few mature females in the catch then it is assumed that there are few mature females at the station, and if the catch is very small then it is assumed that the actual number of anchovies at the station is relatively small. Thus,
each station is given a weight of the relative subsample size.

The estimates and variances, $(\bar{y})$ and $\text{var}(\bar{y})$, of the four parameters—$W$, $R$, $F$, and $S$—are the weighted sample mean and sample variance (Cochran 1963)

$$
\bar{y} = \frac{1}{n} \sum_{i=1}^{n} \left( m_i \right) \bar{y}_i \tag{6}
$$

$$
\text{Var}(\bar{y}) = \frac{1}{n(n-1)} \sum_{i=1}^{n} \left( \frac{m_i}{m} \right)^2 (\bar{y}_i - \bar{y})^2 \tag{7}
$$

where $m_i$ is the number of fish subsampled from the $i$th trawl, $m$ is the average number of fish subsampled per trawl, $n$ is the number of positive trawls, $y_{ij}$ is the observed value for the $j$th fish in the $i$th trawl, and $\bar{y}_i = \frac{1}{m_i} \sum_{j=1}^{m_i} y_{ij}$ is the average for the $i$th trawl.

The method described by equations (6) and (7) is used in the succeeding sections to estimate parameters $W$, $F$, $S$, and $R$ and their respective variances.

**Average Female Weight**

The average female weight, $W$, is calculated as the weighted average of $W_{ij}$, the weight of the $j$th mature female from trawl $i$, using equation (6). The weight of females with hydrated eggs in their ovaries is temporarily inflated because of water retention, therefore their weight is adjusted using the regression of whole-body weight on ovary-free weight estimated from females that do not have hydrated eggs,

$$
\tilde{W}_{ij} = -0.0701 + 1.06 W_{ij}^* \tag{8}
$$

where $W_{ij}^*$ is the ovary-free weight; the regression has an $r^2 = 99.6\%$. The desired subsample size is $m^* = 15$ mature females. The estimated average female weight is 18.83 g with a variance of 0.1319 (equation [7]) and coefficient of variation of 1.937%. The mode of the frequency distribution of $W$, the average weight of mature female anchovies in the $i$th trawl, is at 19 g (Figure 5).

**Batch Fecundity**

For the estimate of batch fecundity, $F$, the individual observations, $F_{ij}$, are not observed directly but are estimated from a relationship between batch fecundity and ovary-free weight. This relationship is estimated from a sample of 109 hydrated females that were collected over the duration of the cruise. These hydrated females have a distribution of adjusted weights (equation [8]) similar to the weight distribution of females for the whole cruise (Figure 6). A linear regression of fecundity on weight was selected because the fitted exponent for the power function was not significantly different from one, indicating that any curvature present in the data is
very slight. The selected regression model is

\[ F_i = -179.7 + 617.2 W_i^* \] (9)

with an \( r^2 = 52.4\% \) (Figure 7).

Based on this regression, the \( F_i \) are estimated for each of the \( m^* = 15 \) mature females from each trawl. Estimated average batch fecundity is 10845 (equation [6]), with a variance of 171730 (equation [10]) and coefficient of variation of 3.82\%. Because the trawl averages \( (F_i) \) are not based on actual observations, but are estimates with their own variance, the variance estimate for \( F \) includes this additional source of variation (Draper and Smith 1966):

\[
\widehat{\text{Var}} (F) = \sum_{i=1}^{n} \left( \frac{m_i}{n} \right)^2 \left[ \left( \frac{F_i - \bar{F}}{n - 1} \right)^2 + \frac{S^2}{109} + (\bar{W}_i^* - \bar{W}_T^*)^2 \text{Var}(b) \right]
\] (10)

where \( S^2 = 6658675 \) is the variance about the regression (9); \( \bar{W}_i^* \) is the average ovary-free weight for the \( i \)th trawl; \( \bar{W}_T^* = 16.54 \) is the average ovary-free weight for the 109 hydrated females; \( \text{Var}(b) = 3172 \) is the variance of the slope of regression (9); and \( n = 135 \) positive trawls.

Estimated average batch fecundity is 10845 (equation [6]), with a variance of 171730 (equation [10]) and coefficient of variation of 3.82\%.

Specific fecundity, expressed as eggs per gram of whole-body weight, is similar to those observed in 1981.

<table>
<thead>
<tr>
<th></th>
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<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>Survey</td>
<td>8003/4</td>
<td>8102</td>
<td>8104</td>
<td>8202</td>
</tr>
<tr>
<td>Mean weight (g)</td>
<td>17.4</td>
<td>13.4</td>
<td>16.2</td>
<td>18.8</td>
</tr>
<tr>
<td>Eggs/gram</td>
<td>444.4</td>
<td>623.0</td>
<td>546.0</td>
<td>575.9</td>
</tr>
</tbody>
</table>

**Spawning Fraction**

Spawning fraction, \( S_i \), is estimated from \( m^* = 15 \) mature females, and \( S_i^* = 1 \) if the ovary of the \( j \)th mature female in the \( i \)th trawl is classified as having day-1 postovulatory follicles (day-1 spawners), and \( S_i^* = 0 \) otherwise (see Hunter and Macewicz 1980 for histological definition of day-1 and day-0 spawners). Thus \( S_i^* \) is the proportion of mature females in the \( i \)th trawl that are in the day-1 spawning category \( (S^1) \).

Another measure of spawning activity is the proportion of mature females that are classified as day-0 spawners, \( S^0 \). In past surveys, it has been suspected that the proportion of day-0 females is a biased estimate because of oversampling of females that spawn on the night of capture. This was indicated by a large discrepancy between \( S^0 \) and \( S^1 \), with the deviation between these two measures reaching a maximum during the peak hours of spawning. There was also a co-occurrence of high values of \( S^0 \) and low values of sex ratios for trawls taken during these hours, which suggested that males and day-0 spawning females segregate out from other females at the peak hours of spawning at a depth where they are more vulnerable to the midwater trawl\(^5\). In previous surveys, \( m_i \) was adjusted by assuming that the actual number of day-0 females is the same as the observed number of day-1 females.

The apparent oversampling of day-0 females was reduced in this year's data; this may have been due to a slight change in the trawl gear. The difference between \( S_i^0 \) and \( S_i^1 \) is still greatest during the peak spawning time 2200-2259 (Figure 8), and there is still a dip in the sex ratio over the time 2000-2259 (Figure 9), but the discrepancy is very small relative to past years. In fact, the values of \( S^0 \) and \( S^1 \), 0.120 and 0.127, respectively, are not significantly different from each other. Thus, we concluded that day-0 spawning females were not oversampled this year. Hence, \( m_i \) is not adjusted and is simply the number of mature females in the \( i \)th trawl. The estimate of spawning fraction is 0.120 (equation [6]) with variance = \( 9.47 \times 10^{-5} \) (equation [7]) and coefficient of variation = 8.13\%.

**Sex Ratio**

The parameter \( R \), sex ratio, is the fraction of females in the anchovy stock based on fish weight.
The sex ratio for each trawl is determined from the first 50 fish. Average weights by sex per trawl are estimated from the first 5 males and the first 20 females, where the weight of hydrated females is adjusted using equation (8). The trawl average, $R$, is the estimated total weight of the females in the first 50 fish subsampled, divided by the estimated total weight of the first 50 fish. The $m$, in equations (6) and (7) for this parameter is the estimated total weight of the first 50 fish. For both sexes, both mature and immature fish are included because of the difficulty of distinguishing between mature and immature males. The estimate for sex ratio is 0.472 (equation [6]), and its estimated variance (equation [7]) and coefficient of variation are $4.90 \times 10^{-4}$ and 4.69%, respectively.

### Biomass Estimate and Variance

The parameter estimates and their variances given above are summarized in Table 1. The resultant estimate of anchovy spawning biomass using equation (1) is 378,000 MT (417,000 short tons). The approximate variance is calculated according to equation (2), where covariance terms are included only for those pairs of variables with sample correlations that are significantly different from zero. The sample correlations between $P_o$ and the adult parameters are assumed to be zero because they are derived from two different surveys (the plankton survey and the trawl survey) and thus cannot be estimated. Of the adult parameters, only cov($W,F$), cov($R,W$), and cov($F,R$) are significant, with values of 110, 0.00274, and 1.57, respectively. The approximate variance of the biomass estimate is $9.45 \times 10^4$, and standard error is 97,200 MT, for a coefficient of variation of 25.7%. The approximate 95% confidence interval is ± 194,000 MT.

**Table 1**

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Symbol</th>
<th>Mean</th>
<th>Variance</th>
<th>Coefficient of Variation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Egg production (eggs/0.05 m²-day)</td>
<td>$P_o$</td>
<td>3.023</td>
<td>0.519</td>
<td>0.237</td>
</tr>
<tr>
<td>Area of survey (0.05 m² units)</td>
<td>$A$</td>
<td>$4.068 \times 10^{12}$</td>
<td>—</td>
<td>—</td>
</tr>
<tr>
<td>Average female weight (grams)</td>
<td>$W$</td>
<td>18.8</td>
<td>0.132</td>
<td>0.0193</td>
</tr>
<tr>
<td>Sex ratio by weight (females/total)</td>
<td>$R$</td>
<td>0.472</td>
<td>$4.90 \times 10^{-4}$</td>
<td>0.0469</td>
</tr>
<tr>
<td>Batch fecundity (eggs per batch per mature female)</td>
<td>$F$</td>
<td>10845</td>
<td>171730</td>
<td>0.0382</td>
</tr>
<tr>
<td>Fraction of mature females spawning per day</td>
<td>$S$</td>
<td>0.120</td>
<td>$9.47 \times 10^{-5}$</td>
<td>0.0813</td>
</tr>
<tr>
<td>Spawning biomass (MT)</td>
<td>$B_s$</td>
<td>378,000</td>
<td>945 \times 10^4</td>
<td>0.257</td>
</tr>
</tbody>
</table>
EQUIVALENT LARVAL CENSUS ESTIMATE OF BIOMASS

The anchovy management plan presently requires that the annual harvest quota for the U.S. fishery be based on an annual estimate of spawning biomass (Smith 1972; Huppert et al. 1980). The PFMC has interpreted the plan to require an equivalent larval census biomass estimate. The 1982 survey was designed to generate an estimate of the biomass based on the egg production and fecundity of the population, but was not designed to directly generate a larval census estimate of biomass.

It is possible to generate an equivalent larval census estimate of biomass by estimating the number of larvae resulting from the measured production of eggs (Stauffer 1983). To do so, the mortality rates of both eggs and larvae are required. The abundance of larvae can then be extrapolated to an annual census using historical proportions of quarterly abundances. The spawning biomass is assumed to be proportional to the annual census; the constant of proportionality was determined from a regression of adult sardine biomass on the annual census of sardine larvae, and an assumption of sardine to anchovy annual fecundity of 1:2 (Smith 1972).

During the 1982 egg production cruise, 96 plankton samples were obtained using the bongo net at standard CalCOFI stations (Figure 10). Anchovy larvae were found at 69 of the stations. In the following sections we discuss: (1) the production curve of the larvae derived from the size-specific catch curve; (2) an equivalent annual larval census derived from the population production of eggs and the subsequent mortality of both eggs and larvae; and (3) the discrepancy between the egg production estimate of biomass and the equivalent larval census estimate.

Production Curve

The larvae were grouped into 12 size categories (yolk-sac to 15 mm), and catches were adjusted for variations in the volume of water filtered per m of depth. Bias corrections were also applied for extrusion of small larvae through the meshes of the net and avoidance of the net by large larvae. The adjusted catches were divided by the duration of growth, through each size class, to estimate the age-specific production of larvae ($P_t$). The adjustments were accomplished by fitting a weighted negative binomial model to the sample frequency distributions of each size class. Each observation is weighted by a factor that is the product of the various adjustments, and the means of the final distributions are unbiased estimates of production ($P_t$). The procedure was developed in a series of papers: Bissel 1972; Zweifel and Smith 1981; Hewitt 1981, 1982; Hewitt and Methot 1982; Hewitt and Brewer 1983. The calculations are summarized in Table 2, and the results are described in Figure 11.

It is clear, from a log-transform plot, that a constant mortality model, $dp/dt = -cP$ will not adequately describe the data. A variable mortality model may be devised by defining $c$ as a function of age. Several forms would adequately describe the data, but for reasons of mathematical tractability we may use:

$$\frac{dp}{dt} = -\frac{\beta}{t^\gamma} P_t.$$  \hspace{1cm} (11)

As $t$ increases, the instantaneous rate of change of production decreases (improving survival with age). The above expression may be rearranged, integrated, and the integration constant determined at the age of hatching ($t_h$):

$$P_t = P_h \left(\frac{t}{t_h}\right)^{-\beta} \text{ for } t \geq t_h.$$  \hspace{1cm} (12)

*Variable mortality rate may be defined as a function of age, size, or population abundance. The important point is that the number of deaths is modeled as a portion of those living.*

N. Lo. SWFC supplied the form used here.
TABLE 2

Size-Specific Catches of Anchovy Larvae and Age-Specific Estimates of Daily Production

<table>
<thead>
<tr>
<th>Preserved size (mm)</th>
<th>Fraction retained by net</th>
<th>Average adjustment for sampling and sorting variability</th>
<th>Average duration of growth (days)</th>
<th>Average catch (larvae/tow)</th>
<th>( P_l ) (larvae/m²-day)</th>
<th>Age since fertilization (days)</th>
</tr>
</thead>
<tbody>
<tr>
<td>2.50</td>
<td>0.21</td>
<td>1.91</td>
<td>3.59</td>
<td>29.99</td>
<td>20.82</td>
<td>4.42</td>
</tr>
<tr>
<td>3.75</td>
<td>0.51</td>
<td>1.99</td>
<td>3.51</td>
<td>26.91</td>
<td>7.56</td>
<td>8.00</td>
</tr>
<tr>
<td>4.75</td>
<td>0.69</td>
<td>1.98</td>
<td>2.58</td>
<td>16.91</td>
<td>4.80</td>
<td>11.07</td>
</tr>
<tr>
<td>5.75</td>
<td>0.73</td>
<td>2.18</td>
<td>2.38</td>
<td>12.12</td>
<td>3.20</td>
<td>13.62</td>
</tr>
<tr>
<td>6.75</td>
<td>0.73</td>
<td>2.18</td>
<td>2.24</td>
<td>8.93</td>
<td>2.51</td>
<td>15.74</td>
</tr>
<tr>
<td>7.75</td>
<td>0.68</td>
<td>2.20</td>
<td>2.16</td>
<td>6.75</td>
<td>2.09</td>
<td>18.13</td>
</tr>
<tr>
<td>8.75</td>
<td>0.61</td>
<td>2.23</td>
<td>2.12</td>
<td>5.10</td>
<td>1.78</td>
<td>20.24</td>
</tr>
<tr>
<td>9.75</td>
<td>0.60</td>
<td>2.09</td>
<td>2.11</td>
<td>3.38</td>
<td>1.28</td>
<td>22.67</td>
</tr>
<tr>
<td>10.75</td>
<td>0.54</td>
<td>2.86</td>
<td>2.12</td>
<td>1.70</td>
<td>0.52</td>
<td>24.92</td>
</tr>
<tr>
<td>11.75</td>
<td>0.56</td>
<td>2.27</td>
<td>2.14</td>
<td>1.06</td>
<td>0.39</td>
<td>27.01</td>
</tr>
<tr>
<td>12.75</td>
<td>0.46</td>
<td>1.75</td>
<td>2.19</td>
<td>0.51</td>
<td>0.29</td>
<td>28.88</td>
</tr>
<tr>
<td>13.75</td>
<td>0.57</td>
<td>2.11</td>
<td>2.29</td>
<td>0.33</td>
<td>0.12</td>
<td>31.93</td>
</tr>
</tbody>
</table>

1Extrusion corrections are based on relative retention rates between 75-micrometer and 505-micrometer mesh nets (Lo, SWFC pers. comm.) Avoidance corrections are based on night and day catch comparisons using bongo data from 1978 and 1979 (Hewitt and Methot 1982). Fraction of larvae retained in the net is estimated as \( \frac{5}{3} \) col. 5/col. 2; \( r = 0.4 \).

2Accounts for partial sorting of samples and standardizes sampling volume to 1 m³ per m of depth.

3Temperature-dependent embryonic growth is determined from laboratory experiment (Lo, SWFC pers. comm.), and post-yolk-sac growth follows Methot's (1981) description.

4Production rates (\( P_l \)) may be estimated by dividing the average catch by the product of columns 2, 3, and 4. In practice, however, \( P_l \) is the mean of a weighted negative binomial model fit to the distribution of individual observations (i.e., plankton tows).

This expression defines the larval production curve. The average standing stock of larvae is the area under the curve, and it may be determined by integrating the above expression between \( t_h \) and 30 days (the maximum age effectively caught with a standard plankton tow). Thus:

\[
N = A \int_{t_h}^{30} P_h \left( \frac{t}{t_h} \right)^{-\beta} dt
\]

\[
= A \frac{P_h t_h}{\beta - 1} \left( 1 - \left( \frac{t_h}{30} \right)^{\beta - 1} \right) \tag{13}
\]

where \( N \) is the average standing stock of larvae and \( A \) is the area of the survey.

**Equivalent Annual Larval Census**

An equivalent larval census estimate of biomass may be derived by reducing the production of eggs to the production of hatching larvae via egg mortality and then integrating the larval production curve over age to estimate average standing stock of larvae. The average standing stock is converted to the annual larval census by a factor \( 2.12 \). The standard sampler would retain a fraction \( r \) of these larvae; the remainder are extruded through the meshes, and a small portion avoid capture. Finally, the annual larval census is converted to spawning biomass using the historical conversion factor of \( 8.9 \times 10^{-8} \) MT/larva. Thus:

\[
\hat{B}_t = N(2.12) r(8.9 \times 10^{-8}) \tag{14}
\]

where: \( r = \frac{\text{fraction of larvae retained in standard sampler}}{\text{fraction of larvae retained in standard sampler}} \),

\[
P_h A = P_o A
\]

\[
N = P_o A \left( \frac{t_h}{30} \right)^{-\beta} \left( 1 - \left( \frac{t_h}{30} \right)^{\beta - 1} \right)
\]
TABLE 3
Projected and Observed Larval Census Estimates of Spawning Biomass and Related Parameters

<table>
<thead>
<tr>
<th></th>
<th>1980</th>
<th>1981</th>
<th>1982</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Feb</td>
<td>Apr</td>
<td></td>
</tr>
<tr>
<td>Population production $P_A$ of eggs (eggs/day)</td>
<td>$2.97 \times 10^5$</td>
<td>$9.07 \times 10^5$</td>
<td>$1.46 \times 10^5$</td>
</tr>
<tr>
<td>Egg mortality rate $z$</td>
<td>0.453</td>
<td>0.138</td>
<td>0.158</td>
</tr>
<tr>
<td>Time to hatch (days) $t_h$</td>
<td>2.71</td>
<td>2.75</td>
<td>2.75</td>
</tr>
<tr>
<td>Survival to hatch $s = e^{-t_h}$</td>
<td>0.293</td>
<td>0.684</td>
<td>1.000</td>
</tr>
<tr>
<td>Coefficient in larval mortality $\beta$ model</td>
<td>1.7999</td>
<td>1.7058</td>
<td>1.7058</td>
</tr>
<tr>
<td>Retention $r$</td>
<td>0.4</td>
<td>0.4</td>
<td>0.4</td>
</tr>
<tr>
<td>Projected larval census $B_l$ estimate of spawning biomass (MT)</td>
<td>$1.533 \times 10^6$</td>
<td>$3.126 \times 10^6$</td>
<td>$2.746 \times 10^6$</td>
</tr>
<tr>
<td>Larval census estimate of $B_l$, spawning biomass (MT)</td>
<td>$1.611 \times 10^6$</td>
<td>$2.544 \times 10^6$</td>
<td>$2.544 \times 10^6$</td>
</tr>
<tr>
<td>Daily fecundity ($\text{eggs/day}$) $q$</td>
<td>$30.28 \times 10^6$</td>
<td>$33.03 \times 10^6$</td>
<td>$33.84 \times 10^6$</td>
</tr>
</tbody>
</table>

$P_A$ is the population production of hatching larvae, $P_A$ is the population production of eggs, $z$ is the fraction of egg production surviving to hatch, and $B_l$ is the equivalent larval census estimate of spawning biomass (MT).

Equation (14) was applied to the data presented in this report; the parameter values are listed in Table 3. The 1982 equivalent larval census estimate of spawning biomass is $1.866 \times 10^8$ MT (2.060 $\times 10^8$ short tons); the equivalent annual larval census is 20,966 $\times 10^6$ larvae.

Discrepancy between the Biomass Estimates
The equivalent larval census estimate of spawning biomass ($1.866 \times 10^8$) is nearly five times the egg production estimate of spawning biomass ($0.378 \times 10^8$ MT). One or more of three factors may be the cause of this discrepancy: (1) sampling errors between the ichthyoplankton gears; (2) variability in the conversion factor relating the larval census to the spawning biomass (resulting from variability in the population fecundity and survival of young, which are assumed to be constant in the larval census method); and (3) systematic inaccuracy in the conversion factor relating anchovy larvae to anchovy spawners.

The projected abundance of larvae and the measured abundance of larvae agree reasonably well for three surveys (Table 3). This indicates that the two sampling gears (CalVET net, used to sample eggs, and the CalCOFI bongo net, used to sample larvae) yield consistent and compatible estimates of ichthyoplankton production. It is possible to predict the catch of one gear from the catch of the other. The discrepancy between the biomass estimates is thus not due to inaccuracy or imprecision in the ichthyoplankton sampling.

The population fecundity and survival of larvae did not vary considerably among the three surveys (Table 3). Egg survival, however, did vary, ranging from an estimated 29% survival in 1980 to 100%
Adjustment of Larval Census Estimates of Spawning Biomass

CalCOFI Rep.: Vol. XXIV, 1983

TABLE 4

<table>
<thead>
<tr>
<th>Year</th>
<th>1980</th>
<th>1981</th>
<th>1982</th>
</tr>
</thead>
<tbody>
<tr>
<td>Adjusted larval census estimates of spawning biomass for variability in egg survival</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Feb</td>
<td>Apr</td>
<td></td>
</tr>
<tr>
<td>S</td>
<td>0.293</td>
<td>0.684</td>
<td>1.000</td>
</tr>
<tr>
<td>Bc</td>
<td>792 x 10^3</td>
<td>577 x 10^3</td>
<td>339 x 10^3</td>
</tr>
<tr>
<td>Bc</td>
<td>1611 x 10^3</td>
<td>2544 x 10^3</td>
<td>2544 x 10^3</td>
</tr>
<tr>
<td>Bc</td>
<td>0.650</td>
<td>2.218</td>
<td>0.950</td>
</tr>
<tr>
<td>Adj. S</td>
<td>3.573 x 10^4</td>
<td>2.417 x 10^4</td>
<td>1.654 x 10^4</td>
</tr>
<tr>
<td>Adj. Bc</td>
<td>4.5</td>
<td>4.2</td>
<td>4.9</td>
</tr>
<tr>
<td>Adj. Bc</td>
<td>0.300</td>
<td>0.977</td>
<td>0.439</td>
</tr>
<tr>
<td>Adj. S</td>
<td>1.000</td>
<td>3.412</td>
<td>1.462</td>
</tr>
<tr>
<td>Adj. Bc</td>
<td>5.498 x 10^4</td>
<td>3.719 x 10^4</td>
<td>2.544 x 10^4</td>
</tr>
<tr>
<td>Adj. Bc</td>
<td>6.9</td>
<td>6.4</td>
<td>7.5</td>
</tr>
</tbody>
</table>

Thus the discrepancy between the larval census estimates and the egg production estimates of spawning biomass can be explained by variation in egg survival and inaccuracy in estimating the average proportionality between the larval census and the spawning biomass. We further consider the egg production method to be more precise (because it considers the rate at which eggs die) and more accurate (because of the underestimate of anchovy fecundity) than the larval census method.

ACKNOWLEDGMENTS

The work reported here is the result of a cooperative survey conducted by the National Marine Fisheries Service (NMFS) and the California Department of Fish and Game (CDF&G). Field collections, laboratory processing, and data analysis were conducted under the direction of Dr. Reuben Lasker, James Thrailkill, Dr. John Hunter, and Dr. Geoff Moser of NMFS, and Kenneth Mais and Alec MacCall, CDF&G. We especially thank the crews of NOAA R/V David Starr Jordan and of F/V Oregon Beaver, and also Eric Lynn, William Flexx, Jack Metoyer, Dr. John Graves, Richard Charter, Elaine Sandknop, Betsy Stevens, Morgan Busby, Barbara Sumida, Scott Miller, Beverly Maciewicz, Pete Paloma, and Julie Shoemaker of NMFS, J. Jensen of Southwestern College, and Lucy Dunn. Mary Farrell, Jeannie Haddox, Mary Lumpkins and Frances Pocinhich of Scripps Institution of Oceanography. We also thank Dr. Richard Methot and Dr. Gary Stauffer, NMFS, and Alec MacCall, NMFS (formerly CDF&G), for their suggestions and review.

LITERATURE CITED


