# A length-based model for cockles on Snake Bank, Whangarei Harbour, 2002 

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Final Research Report for Ministry of Fisheries Research Project COC2001/01<br>Objective 2

## Final Research Report

Title:

Authors:

1. Date:
2. Contractor:
3. Project Title: Stock assessment of cockles in Whangarei Harbour including Snake Bank
4. Project Code:
5. Project Leader: Martin Cryer

## 6. Duration of Project:

Start date: 1 October, 2001
Expected completion date: 30 September, 2002

## 7. Executive Summary:

A stochastic, dynamic, length-based, observation-error, time-series model for cockles on Snake Bank, Whangarei Harbour, is described. The model is driven by reported landings, and fitted using Bayesian techniques to estimates of absolute biomass and length frequency distributions from roughly annual surveys on the bank since 1983, to periodic length frequency distributions from the commercial catch, and to length frequency distributions from roughly quarterly surveys on part of the bank between 1992 and 1995. Two alternative growth models are explored; one has normal error structure and allows negative growth whereas the other has log-normal error structure and does not allow negative growth. Both models seem over-prescribed and tests on MCMC chains suggested they did not converge on estimates for many important parameters. To explain rapid changes in biomass, both models favour implausibly high rates of natural mortality and growth. Neither model could resolve the apparent incompatibility of the biomass and length frequency data sets, and the results were very sensitive to the relative weights accorded each data set. Because of these problems, we conclude that the models do not provide a reasonable description of the observed data and we are not confident that the estimated parameter values are useful.

## 8. Objectives:

Only Objective 2 is pertinent to this report; the other objectives have been reported separately.

1. To estimate the size structure and absolute biomass of cockles on Snake Bank during the 2001/2002 fishing year. The target coefficient of variation (c.v.) of the estimate of absolute recruited biomass is $20 \%$.
2. To complete the cockle stock assessment and estimate yields for cockles on Snake Bank for the 2002/2003 fishing year.
3. To estimate the size structure and absolute biomass of cockles in other areas within Whangarei Harbour during the 2001/2002 fishing year. The target coefficient of variation (c.v.) of the estimate of absolute recruited biomass is $20 \%$.

## 9. Methods:

### 9.1 Background and previous stock assessments

Snake Bank is a sandbank of about $1.5 \mathrm{~km}^{2}$ close to the mouth of Whangarei Harbour in northern New Zealand (Figure 1). A commercial fishery for cockles has operated on the bank since at least the early 1980s, and this is the only bank within the harbour where commercial fishing is permitted. There are other cockle beds in the harbour but none has a density of large cockles similar to that on Snake Bank (Cryer et al., 2003).


Figure 1: Beaches and banks within Whangarei Harbour, including Snake Bank. Unhatched grey indicates sandy, intertidal areas, and black polygons on these substrates indicate sampling strata from surveys on Snake Bank and elsewhere (after Cryer et al. 2002, 2003).

Surveys of the cockle population on Snake Bank have been conducted since 1982, some time after the first fishing, but well before a substantial fishery developed. Grid surveys were used until 1996 (Cryer, 1997), and various stratified random designs since (e.g., Morrison \& Cryer 1999, Morrison 2000, Morrison \& Parkinson 2001, Cryer \& Parkinson 2001, Cryer et al. 2002). Whatever the design, the overall mean biomass of cockles on the bank (for a given size range) was estimated using the weighted average of the stratum estimates of mean biomass, weights being proportional to the relative area of each stratum:

$$
\begin{equation*}
\bar{x}_{y}=\sum_{i=1}^{n} W_{i} \bar{x}_{i} \tag{1}
\end{equation*}
$$

where $\bar{x}_{y}$ is the overall mean biomass, $W_{i}$ is the relative area and $\bar{x}_{i}$ the mean biomass in stratum $i$. These are considered estimates of absolute biomass (or abundance if estimated using numbers). Variance was estimated using:

$$
\begin{equation*}
s_{y}^{2}=\sum_{i=1}^{n} W_{i}^{2} s_{i}^{2} / n_{i} \tag{2}
\end{equation*}
$$

where $s_{y}^{2}$ is the variance of the estimated mean biomass, $s_{i}^{2}$ is the sampling variance in stratum $i$, and $n_{i}$ is the number of samples taken in stratum $i$ (Snedecor \& Cochran 1989).

Station length frequency distributions were estimated by scaling the recorded length frequency distributions by the inverse of the sampled fraction at each station and to a square metre of sediment. Stratum length frequency distributions were estimated as the average station length frequency distribution for that stratum scaled by the stratum area (in square metres). The population length frequency was estimated by adding the stratum length frequency distributions.

Based on these surveys, yield for Snake Bank cockles was estimated by Cryer et al. (2002) as Maximum Constant Yield (MCY, method 2. equation 3) and Current Annual Yield (CAY, using the full version of the Baranov Catch Equation, equation 4, see also Annala et al. 2002).

$$
\begin{gather*}
M C Y=0.5 \cdot F_{0.1} \cdot B_{a v}  \tag{3}\\
C A Y=\frac{F_{r e f}}{F_{r e f}+M}\left(1-e^{-\left(F_{r q}+M\right)}\right) B_{b e g} \tag{4}
\end{gather*}
$$

where $F_{0.1}$ and $F_{r e f}$ are reference rates of fishing mortality, $M$ is natural mortality, $B_{a v}$ is the average recruited biomass between 1991 and 2001, and $B_{b e g}$ is the start of season recruited biomass. Estimates of $\mathrm{M}=0.30, \mathrm{~F}_{0.1}=0.41$ (from Cryer 1997) were used. An estimate of $\mathrm{F}_{\max }=0.62$ is available (Cryer 1997) but was not used.

The above describes a fairly crude stock assessment, and the Ministry's Shellfish Fishery Assessment Working Group decided that length-based modelling would be a better use of the data and should provide better estimates of yield (and projections of future stock status under different management regimes).

### 9.2 Input data for the length-based model

The following data were available for the length-based model: catch records since 1982 when the population was assumed to be close to virgin (Table 1); 13 estimates of absolute abundance (with variances) for cockles on Snake Bank between 1982 and 2002 (Table 2, Figure 2); 11 estimates of population length frequency distribution from surveys other than that in 1982 (Figure 3); 9 estimates of length frequency distribution from one part of the bank between August 1992 and March 1995 (Figure 4); and 4 estimates of the length frequency distribution of cockles taken by commercial fishers in 1992, 1996, and 2001 (early and late in the year, Figure 5).

Table 1: Reported commercial landings of cockles from Snake Bank since 1982 (after Annala et al., 2002) (values for 1982-86 and 2002 were assumed). No recreational or customary catch is included as these are thought to be very small compared with commercial landings (e.g., Annala et al., 2002, p. 112).

| Year | Landings (t) | Year | Landings (t) |
| :--- | ---: | ---: | ---: |
|  |  |  |  |
| 1982 | 162 | 1993 | 316 |
| 1983 | 162 | 1994 | 566 |
| 1984 | 162 | 1995 | 501 |
| 1985 | 162 | 1996 | 495 |
| 1986 | 162 | 1997 | 457 |
| 1987 | 114 | 1998 | 439 |
| 1988 | 128 | 1999 | 473 |
| 1989 | 255 | 2000 | 473 |
| 1990 | 426 | 2001 | 423 |
| 1991 | 396 | 2002 | 423 |
| 1992 | 537 |  |  |

Table 2: Estimates of biomass (t) of cockles on Snake Bank from Cryer et al. (2002). Estimates marked with an asterisk (*) were made by analysis of length frequency distribution, others by directly weighing samples sorted into three size classes.

| Year | n | Total |  | $<30 \mathrm{~mm}$ |  | $>30 \mathrm{~mm}$ |  | $>35 \mathrm{~mm}$ |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | Biomass | c.v. | Biomass | c.v. | Biomass | c.v. | Biomass | c.v. |
| 1982 | 199 | 2556 | - | * 216 | - | * 2340 | - | 1825 | $\sim 0.10$ |
| 1983 | 187 | 2509 | - | * 321 | - | * 2188 | - | 1700 | $\sim 0.10$ |
| 1985 | 136 | 2009 | 0.08 | * 347 | $\sim 0.10$ | 1662 | 0.08 | 1174 | $\sim 0.10$ |
| 1988 a | 53 | - | - | - | - | 1140 | $>0.15$ | - | - |
| 1988 b | 53 | - | - | - | - | 744 | $>0.15$ | - | - |
| 1991 | 158 | 1447 | 0.09 | 686 | 0.10 | 761 | 0.10 | 197 | 0.12 |
| 1992 | 191 | 1642 | 0.08 | 862 | 0.10 | 780 | 0.08 | 172 | 0.11 |
| 1995 | 181 | 2480 | 0.07 | 1002 | 0.09 | 1478 | 0.07 | 317 | 0.12 |
| 1996 | 193 | 1755 | 0.07 | 959 | 0.09 | 796 | 0.08 | 157 | 0.11 |
| 1998 | 53 | 2401 | 0.18 | 1520 | 0.20 | 880 | 0.17 | 114 | 0.20 |
| 1999 | 47 | 3486 | 0.12 | 2165 | 0.12 | 1321 | 0.14 | 194 | 0.32 |
| 2000 | 50 | 1906 | 0.23 | 1336 | 0.24 | 570 | 0.25 | 89 | 0.32 |
| 2001 | 51 | 1405 | 0.17 | 970 | 0.18 | 435 | 0.17 | 40 | 0.29 |
| 2002 | 53 | 1618 | 0.14 | 1152 | 0.15 | 466 | 0.19 | 44 | 0.29 |



Figure 2: Estimated total biomass of cockles ( $\pm$ one standard error) on Snake Bank, 1982-2002, and the estimated proportion of biomass (dashed line) accounted for by cockles of 30 mm or more shell length.


Figure 3: Estimated population length frequency distribution of cockles on Snake Bank since 1983 (data for 1982 lost). The shaded area in each histogram shows those animals 30 mm shell length and larger (the nominal size at recruitment to the fishery).


Figure 4: Estimated length frequency distribution of cockles near the lagoon on Snake Bank between 1993 and 1995 (data collected to estimate growth rate. The shaded area in each histogram shows those animals 30 mm shell length and larger (the nominal size at recruitment to the fishery).

### 9.3 Structure of the model

A length based model was used to assess the snake bank cockle population. This model was adapted from a model developed by Breen et al. (2000) to assess paua (Haliotus iris) in PAU 5B and 5D. The model is a stochastic, dynamic, length-based observation-error time series model. It is stochastic because annual variations in recruitment can be estimated as a vector of free parameters. It is dynamic because no equilibrium, other than in the initial length structure, is assumed. Cockles are represented in the model as numbers-at-length rather than numbers-at-age. The error is assumed to be observation error rather than process error.

The model is length-based, with 47 length 'bins', each of 1 mm shell length. The left-hand edge of the first bin is 4 mm ; the largest bin is a plus-group representing cockles larger than 50 mm . Sexes are not distinguished. The time step is one season ( 3 months). The model is implemented in AD Model Builder ${ }^{\top \mathrm{M}}$ (Otter Research Ltd., http://otterrsch.com/admodel.htm). AD Model Builder ${ }^{\top M}$ incorporates a Markov chain Monte Carlo procedure for the calculation of Bayesian posterior distributions.


Figure 5: Estimated length frequency distribution of cockles in the commercial harvest from Snake Bank between in 1992, 1996, and 2001 ( $a=$ summer, $b=$ winter). The shaded part of each histogram contains animals 30 mm shell length and larger (the nominal size at recruitment to the fishery).

The model population is initialised and then driven by reported catches. The model is fitted, using maximum likelihood methods, to vectors of absolute abundance estimates from surveys, survey length frequency samples, and commercial length frequency samples. Outputs are the past, present, and (potentially) projected states of the stock, estimated using Bayesian methods. Parameters estimated in the model are:

| $\ln (R 0)$ | log of base recruitment $R 0$ |
| :---: | :---: |
| $l^{R}$ | mean size of new recruits to population |
| $\left(\sigma^{R}\right)^{2}$ | variance of the size distribution of newly recruited animals |
| M | instantaneous annual natural mortality rate |
| $L_{\infty}$ | asymptotic length |
| or g10 | mean growth increment at 10 mm (alternative linear model) |
| K | Brody coefficient, year ${ }^{-1}$ |
| or g30 | mean growth increment at 30 mm (alternative linear model) |
| $\alpha$ | c.v. of the expected growth increment |
| $\sigma_{M I N}$ | minimum standard deviation of expected growth increment |
| $v^{1 r}$ | shape parameter for the left-hand limb of research survey selectivity |
| $\nu^{1 s}$ | shape parameter for the left-hand limb for commercial catch |
| $\phi^{\text {diff }}$ | The average shift in length of $\phi_{1}^{s}$ from period to period. |
| $\tilde{\boldsymbol{\sigma}}$ | common component of observation error |
| $\varepsilon$ | vector of recruitment deviations |

Related parameters that are fixed in the model (see also Appendix 1) are:

```
\phi
vr shape parameter for the right-hand limb of research survey selectivity fixed at 10
\phi
v 2s shape parameter for the right-hand limb for commercial catch fixed at 200
```


### 9.4 Fitting the model and diagnostics

## Initial Conditions

The initial population is assumed to be in equilibrium with zero fishing mortality and the base recruitment. The model is run for 60 periods with no fishing to obtain near equilibrium in numbers-at-length. To start, recruitment is evenly divided among the bins (it is added to any animals remaining there after growth from the previous quarter) in proportions determined from:

$$
\begin{equation*}
\beta_{k}^{R}=\exp \left(-\left(\frac{l_{k}-l^{R}}{\sigma^{R}}\right)^{2}\right) \tag{5}
\end{equation*}
$$

where $l_{k}$ is the midpoint length of the $k$ th length class, $l^{R}$ and $\sigma^{R}$ are the mean and the standard deviation of the distribution of recruited cockle's length. The last bin acts as a "plus group". The recruitment happens annually in a chosen season for the burn-in and project periods, and in any period with $\tau_{t}^{R}=1$ for periods with data, hence the number of recruits in period $t$ is:

$$
\begin{equation*}
R_{k, t}=\tau_{t}^{R} \beta_{k}^{R} R 0 \tag{6}
\end{equation*}
$$

where the $\tau_{t}^{R}$ is a switch based on whether the recruitment happens in period $\mathrm{t}\left(\tau_{t}^{R}=1\right)$ or not ( $\tau_{k}^{R}=0$ ).

## Growth Transition Matrix

During initialisation, the vector $N_{t}$ of numbers-at-length is determined from numbers in the previous year, survival, the growth transition matrix $G$ and the vector of recruitment:

$$
\begin{equation*}
N_{t}^{\prime}=N_{t-1}^{\prime} \bullet G \exp (-\dot{M} / 4)+R_{t} \tag{7}
\end{equation*}
$$

where the prime $\left({ }^{\prime}\right)$ denotes vector transposition and the $\operatorname{dot}(\bullet)$ denotes matrix multiplication.
Incremental growth in the model was calculated using either a linear increment (von Bertalanffy) or exponential increment growth function. Variation in growth about the mean increment was determined using either normal or log normal error structure. Three options for describing incremental growth and variation were investigated.

## 1. von Bertalanffy function normal error structure

A growth transition matrix is calculated inside the model from the estimated growth parameters. The expected annual growth increment for the $k$ th length class is given by the von Bertalanffy growth function for $l_{k}<L_{\infty}$ and is zero where $l_{k} \geq L_{\infty}$ :

$$
\begin{array}{cl}
\Delta l_{k}=\left(L_{\infty}-l_{k}\right)(1-\exp (-K / 4)) & \text { for } l_{k}<L_{\infty} \\
\Delta l_{k}=0 & \text { for } l_{k} \geq L_{\infty} \tag{8b}
\end{array}
$$

The standard deviation of this increment is assumed to be proportional to the increment with a minimum $\sigma_{M I N}$, and is given by a smooth differentiable function:

$$
\begin{equation*}
\sigma^{\Delta l_{k}}=\left(\Delta l_{k} \alpha-\sigma_{\text {MIN }}\left(\frac{1}{\pi} \times \tan ^{-1}\left(\left(\Delta l_{k} \alpha-\sigma_{\text {MIN }}\right) \times 10^{6}\right)+0.5\right)+\sigma_{\text {MIN }}\right. \tag{9}
\end{equation*}
$$

From the expected increment and standard deviation for each length class, the probability distribution of growth increments for a cockle of length $l_{k}$ is calculated from the normal distribution, and translated into the vector of probabilities of transition from length bin $k$ to other length bins to form the growth transition matrix G. Zero and negative growth increments are permitted.

## 2. Linear increment model (von Bertalanffy) log normal error structure

For many fish species the von Bertalanffy function describes mean length at age well. The rate of change (differential) in mean increment defined by the von Bertalanffy function is linear. A linear increment model was formulated on the basis of two parameters g10 and g30 which represent the expected annual growth increment at 10 and 30 mm shell length, respectively.

$$
\begin{equation*}
\Delta l_{k}=\frac{30 g_{10}-10 g_{30}}{g_{10}-g_{30}} \text { (after Francis 1988) } \tag{10}
\end{equation*}
$$

It is preferable to estimate growth relative to these linear parameters and, thereby, avoid the high correlation that typically occurs when $K$ and $L_{\infty}$ parameters are estimated (Francis 1988).

The log-normal probability density for growth increment $\Delta l$, for a cockle of length $l$ is given by,

$$
\begin{equation*}
\mathrm{f}\left(\Delta l_{k}\right)=\frac{1}{\sqrt{2 \pi} \sigma_{k} \Delta l} \exp \left[-\frac{1}{2}\left(\frac{\log \left(\Delta l_{k}\right)-\log \left(\Delta l_{k}\right)+\frac{1}{2} \sigma_{k}^{2}}{\sigma_{k}}\right)^{2}\right] \tag{11}
\end{equation*}
$$

where $\sigma_{l}$ is the standard deviation of the expected increment at length $l$. Negative growth increments are not permitted.

## Dynamics

For each period $t$, the model calculates the biomass available to the fishery based on the selectivity $V_{k, t}^{s}$ and the average weight $w_{k}$ :

$$
\begin{equation*}
B_{t}=\sum_{k} N_{k, t} V_{k, t}^{s} w_{k} . \tag{12}
\end{equation*}
$$

The selectivity of the commercial fishery is assumed to have changed over time, taking smaller cockles in later years. Hence, the selectivity, $V_{k, t}^{s}$ is calculated from:

$$
\begin{equation*}
V_{k, t}^{s}=\lambda_{k, t}^{s} \exp \left(\frac{(\ln 0.5)\left(l_{k}-\phi_{t}^{s}\right)^{2}}{\left(v_{t}^{s s}\right)^{2}}\right)+\left(1-\lambda_{k, t}^{s}\right) \exp \left(\frac{(\ln 0.5)\left(l_{k}-\phi_{t}^{s}\right)^{2}}{\left(v_{t}^{2 s}\right)^{2}}\right) \tag{13}
\end{equation*}
$$

where $\phi_{t}^{s}=\phi_{1}^{s}-(t-1) \phi^{\text {diff }}$ and $\lambda_{k, t}^{s}=\frac{1}{1+\exp \left(-\left(l_{k}-\phi_{t}^{s}\right) \delta\right)}$
where $\delta$ is a shape parameter for mixing selectivity curves (assumed to be 5 ).
The observed catch is then used to calculate exploitation rate, which was limited to a $U^{\max }$ with the posfun function of AD Model Builder ${ }^{\text {TM }}$. If the ratio of catch to biomass exceeds this, then exploitation rate is restricted to just over $U^{\max }$ and a penalty is added to the total negative $\log$-likelihood function. Let $A_{\min }$ be the survival rate at $U^{\max }$, i.e. 1- $U^{\max }$, and $A_{t}$ be 1- $U_{t}$ :

$$
\begin{array}{r}
A_{t}=1-\frac{C_{t}}{B_{t}} \quad \text { for } \frac{C_{t}}{B_{t}} \leq U^{\max } \\
A_{t}=0.5 A_{\min }\left[1+\left(1+3-\frac{2\left(1-\frac{C_{t}}{B_{t}}\right)}{A_{\min }}\right)^{-1}\right] \text { for } \frac{C_{t}}{B_{t}}>U^{\max } \tag{16}
\end{array}
$$

The penalty invoked when exploitation rate is limited is:

$$
\begin{equation*}
1000000\left(A_{\min }-\left(1-\frac{C_{t}}{B_{t}}\right)\right)^{2} \tag{17}
\end{equation*}
$$

Survival from fishing is calculated as:

$$
\begin{equation*}
S F_{k, t}=1-\left(1-A_{7}\right) V_{k, t}^{s} \tag{18}
\end{equation*}
$$

The vector of numbers-at-length in the following year is calculated from:

$$
\begin{equation*}
N_{t}^{\prime}=\left(S F_{t-1} \otimes N_{t-1}\right)^{\prime} \cdot \mathrm{G} \exp (-M / 4)+R_{t} \tag{19}
\end{equation*}
$$

where $\otimes$ denotes a vector element product and recruitment is modified by the estimated deviations:

$$
\begin{equation*}
R_{k, t}=\tau_{t}^{R} \beta_{k}^{R} R 0 \exp \left(\varepsilon_{t}-0.5 \sigma_{\varepsilon}^{2}\right) \tag{20}
\end{equation*}
$$

## Predictions

The predicted survey index is calculated from model numbers in bins greater than 5 mm length, taking into account sampler selectivity-at-length:

$$
\begin{equation*}
\hat{J}_{t}=\sum_{k=2}^{47} N_{k, t} V_{k}^{r} \tag{21}
\end{equation*}
$$

where $V_{k}^{r}$ is calculated from:

$$
\begin{equation*}
V_{k}^{r}=\lambda_{k}^{r} \exp \left(\frac{(\ln 0.5)\left(l_{k}-\phi^{r}\right)^{2}}{\left(v^{1 r}\right)^{2}}\right)+\left(1-\lambda_{k}^{r}\right) \exp \left(\frac{(\ln 0.5)\left(l_{k}-\phi^{r}\right)^{2}}{\left(v^{2 r}\right)^{2}}\right) \tag{22}
\end{equation*}
$$

and

$$
\begin{equation*}
\lambda_{k}^{r}=\frac{1}{1+\exp \left(-\left(l_{k}-\phi^{r}\right) \delta\right)} \tag{23}
\end{equation*}
$$

where $\delta$ is assumed to be 5 .
The model predicts proportions-at-length for the research survey from numbers in each length class for lengths greater than 5 mm :

$$
\begin{equation*}
\hat{p}_{k, t}^{r}=\frac{N_{k, t} V_{k}^{r}}{\sum_{k=2}^{47} N_{k, 1} V_{k}^{r}} \tag{24}
\end{equation*}
$$

Predicted proportions-at-length for commercial catch sampling are similar starting at length 19 mm :

$$
\begin{equation*}
\hat{p}_{k, t}^{s}=\frac{N_{k, t} V_{k, t}^{s}}{\sum_{k=16}^{k=16} N_{k, t} V_{k, t}^{s}} \tag{25}
\end{equation*}
$$

## Fitting

The likelihood for the predicted survey index is:

$$
\begin{equation*}
L\left(\hat{J}_{t} \mid \theta\right)=\frac{\varpi^{J}}{\sigma_{t}^{J} \tilde{\sigma} \sqrt{2 \pi}} \exp \left[\frac{-\left(\ln \left(J_{t}\right)-\ln \left(\hat{J}_{t}\right)\right)^{2}}{2\left(\sigma_{t}^{J} \tilde{\sigma} / \sigma^{J}\right)^{2}}\right] \tag{26}
\end{equation*}
$$

where $\theta$ is the parameter vector. The following likelihood is used to fit model predictions to observed proportions-at-length from research surveys (Breen et al. 2002).

$$
\begin{equation*}
L\left(\hat{p}_{k, t}^{r} \mid \theta\right)=\frac{\kappa_{t}^{r} \nabla^{r} \sqrt{\left(p_{k, t}^{r}+0.1\right)}}{\tilde{\sigma} \sqrt{2 \pi}} \exp \left[\frac{-\left(p_{k, t}^{r}+0.1\right)\left(p_{k, t}^{r}-\hat{p}_{k, t}^{r}\right)^{2}}{2\left(\tilde{\sigma} / \kappa_{t}^{r} \sigma^{r}\right)^{2}}\right] \tag{27}
\end{equation*}
$$

where $\kappa_{t}^{r}$ is the square root of numbers measured in period $t$. The negative log-likelihood is summed for all years with observations and for all length classes.

The likelihood components for commercial catch sampling and the quarterly samples are analogous.

The optimum model fit to the data was deemed to be at the mode of the joint posterior distribution (MPD) i.e., the minimum negative log value of the combined likelihoods and priors. The MPD was estimated derived using the auto differentiation-based minimiser in procedure of ADMB.

The optimum model fit to the data was deemed to be at the mode of the joint posterior distribution (MPD) i.e., the minimum negative log value of the combined likelihoods and priors. The MPD was estimated using the auto differentiation-based minimiser in ADMB .

## Likelihood weighting

Changing the relative weightings on the likelihood terms was achieved by multiplying the common observational error term ( $\tilde{\sigma}$ ) by an inverse scalar ( $w_{\text {likex }}$ ).

$$
\begin{equation*}
\tilde{\sigma}_{w_{\text {itax }}}=\tilde{\sigma}\left(\frac{1}{w_{l i k e_{x}}}\right) \tag{28}
\end{equation*}
$$

## Priors, bounds and assumptions

Bayesian priors were established for all parameters. With the exception of natural mortality, $M$, and recruitment all were uninformative, incorporated simply as uniform distributions with upper and lower bounds set so wide as not to restrict the estimation unless highly implausible values were explored (Appendix 2)

The prior probability density for $M$ was calculated from the normal distribution (Appendix 3):

$$
\begin{equation*}
L(M)=\frac{1}{\sigma_{M}^{2} \sqrt{2 \pi}} \prod_{t} \exp \left(-\frac{M_{t}^{2}}{2 \sigma_{M}^{2}}\right) \tag{29}
\end{equation*}
$$

The prior probability density for the vector of estimated recruitment deviations vector, $\varepsilon$, was calculated from the normal distribution:

$$
\begin{equation*}
L(\varepsilon)=\frac{1}{\sigma_{\varepsilon}^{2} \sqrt{2 \pi}} \prod_{t} \exp \left(-\frac{\varepsilon_{t}^{2}}{2 \sigma_{\varepsilon}^{2}}\right) \tag{30}
\end{equation*}
$$

## Biological assumptions

The length-weight relation was taken from Cryer (1997) and was

$$
\begin{equation*}
w_{k}=1.6010^{-7} l_{k}^{3.29} \tag{31}
\end{equation*}
$$

where $l_{k}$ is the length in millimetres and $w_{k}$ is the weight in kilograms.
In calculating spawning biomass, maturity-at-length, $m_{k}$, was assumed to be knife-edged at 19 mm . Spawning biomass is:

$$
\begin{equation*}
S_{t}=\sum_{k} N_{k, t} m_{k} \tag{32}
\end{equation*}
$$

## Bayesian Posteriors

The posterior distributions of key model parameters were characterised using the HastingsMetropolis Markov Chain Monte Carlo (MCMC) sampling procedures (Hastings 1970). The MCMC procedure was run for one million iterations and was sampled at every $100^{\text {th }}$ iteration. The Geweke (1992) test of convergence was applied to each chain from the MCMC sampling process.

## 10. Results:

## Normal error model fits

Under uniform weighting across all likelihoods the model achieved an excellent fit to the biomass estimates (Figure 6). However, the biomass trajectory from this model was highly erratic, the estimate of natural mortality was implausibly high (on its upper bound of $M=0.8$, Table 3) and the fit to the survey length frequency distributions was poor (especially for the early years where the model did not reproduce the preponderance of very large cockles, Appendix 3). The fit to the commercial length frequency distributions was very poor and predicted a much wider range of cockles, especially small cockles, in the harvest than observed (Appendix 3). The fit to the seasonal samples was also very poor and the model did not reproduce the observed pattern of recruiting cohorts that are such a dominant feature of this data set. A poor fit to the seasonal samples is not necessarily fatal because these samples
were collected from one small part of the bank and there may be spatial variation in spat fall and growth. However, we consider poor fits to length frequency distributions integrated over the whole population and to measurements from the commercial catch to be very serious flaws of the model with "equal" likelihood weightings.


Figure 6: Model fits to survey biomass estimates under increasing weighting toward commercial and survey length frequency data (weighting ratios biomass : commercial LF : survey LF)

Improved fits to length frequency data and more plausible estimates of $M$ were achieved by increasing the relative weighting on the commercial and survey length frequency data (Table 3; Appendix 4). However, these improvements were achieved at the expense of the fit to the biomass series (Figure 6). A weighting factor of 5 and 20 on the commercial and survey length frequency likelihoods, respectively, was thought to give the best overall fit to the observed data and we have nominated this as the base case (Figure 6; Table 3; Appendix 4). As well as fitting the length frequency data better, this fit also has a much less erratic trajectory than that having "equal" weightings. The 1:5:20 model reproduced the commercial and seasonal length frequency distributions reasonably well, but still failed to predict the very large cockles that dominated the population in the early 1980s.

Assuming normal error around the mean growth increment allows negative growth of cockles in the model. In the base case, $38 \%$ of animals with starting widths 28 mm or greater shrank at least 1 mm over the annual growth period (Figure 7). We think this amount of negative growth to be biologically implausible, and this was our motivation for testing a lognormal error structure as an alternative.

## Normal error model base case diagnostics

Natural mortality ( $M$ ) and mean recruitment (Rcoff) were highly correlated in the fitting process ( 0.983 , Appendix 5a) and there was significant correlation between the two growth parameters $K$ and $\operatorname{Linf}$ (Appendix 6a). Most of the annual recruitment parameters were independent of one other.


Figure 7: Percentage of animals growing negatively under base case model fit.
Most of the posterior distributions of the main model parameters were unimodal, and the MLE values generally corresponded with the modes of posterior distributions (Appendix 6). Given that most of the model priors were uniform, this pattern in the posteriors would generally suggest that the model was able to find optimum solutions for most parameters. However, there was evidence of a high degree of autocorrelation in the MCMC traces (Appendix 6) and tests indicated that convergence was not attained for several parameters, including some important ones (Appendix 7a). We conclude that, although optimal solutions can be found for most parameters, the model provides a poor description of the observed data. Our confidence that the estimated parameter values have biological meaning is, therefore, low.

Table3: Parameter estimates from Maximum Probability Density (MPD) fits to observational data under various weighting scenarios (weighting ratios biomass : commercial LF : survey LF) [**Auto differentiation failed to converge (i.e., the Hessian matrix was not positive definite); shaded denotes parameter value on boundary; bold values denotes "base case"]

| Normal Error Model Linf \& K estimated |  |  |  |  |  |  | Lognormal Error Model g10 \& g 30 estimated |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Parameters | *1:1:1 | 1:1:5 | 1:5:5 | 1:5:10 | 1:5:15 | 1:5:20 | 1:1:1 | 1:1:5 | **1:5:5 | 1:5:10 | 1:5:15 | *1:5:20 |
| $\ln (R 0)$ | 19.38 | 19.23 | 19.29 | 18.91 | 18.09 | 17.73 | 19.01 | 19.04 | 19.08 | 18.97 | 18.40 | 17.92 |
| $l^{R}$ | 2.44 | 2.45 | 2.30 | 3.16 | 3.15 | 3.17 | 3.18 | 3.12 | 3.14 | 20.00 | 16.44 | 14.36 |
| M | 0,80 | 0.80 | 0.80 | 0.53 | 0.27 | 0.20 | 0.64 | 0.66 | 0.66 | 0.53 | 0.34 | 0.24 |
| Lsor or 930 | 31.63 | 32.02 | 31.96 | 31.78 | 31.70 | 31.74 | 0.55 | 0.54 | 0.50 | 0.68 | 0.68 | 0.65 |
| $K$ or gl0 | 1.45 | 1.40 | 1.38 | 1.10 | 0.85 | 0.77 | 6.01 | 5.91 | 5.89 | 1.90 | 2.24 | 2.18 |
| $v^{1 r}$ | 4.62 | 3.68 | 4.06 | 3.58 | 3.04 | 2.77 | 2.43 | 1.57 | 1.64 | $49.99$ | 49.98 | 49,99 |
| $v^{\text {ls }}$ | 0.00 | 2.19 | 1.30 | 1.57 | $0,09$ | 1.59 | 0.38 | 0.24 | 1.06 | 0.000 | 0.05 | 0.44 |
| $\phi^{\Delta / 5}$ | -0.58 | -0.02 | -0.03 | -0.03 | -0.05 | -0.02 | -0.04 | -0.05 | -0.04 | -0.05 | -0.05 | -0.04 |
| $\mathcal{E} \quad 1982$ | 0.09 | 0.45 | 0.37 | 0.14 | -0.70 | -1.07 | 0.55 | 0.51 | 0.47 | -0.79 | -0.84 | -0.78 |
| E 1983 | 0.10 | 0.13 | 0.11 | 0.00 | -0.32 | -0.64 | -0.02 | 0.01 | 0.02 | -0.02 | -0.22 | -0.58 |
| $\mathcal{E} 1984$ | 0.06 | 0.30 | 0.21 | -0.20 | -0.68 | -0.82 | 0.14 | 0.10 | 0.08 | -0.17 | -0.32 | -0.47 |
| $\mathcal{E} \quad 1985$ | -0.51 | -0.61 | -0.55 | -0.54 | -0.70 | -0.76 | -0.51 | -0.43 | -0.41 | -0.75 | -0.75 | -0.71 |
| $\mathcal{E} \quad 1986$ | -0.38 | -0.25 | -0.30 | -0.60 | -0.94 | -0.89 | -0.40 | -0.39 | -0.41 | -0.73 | -0.83 | -0.78 |
| E 1987 | -0.60 | -0.45 | -0.51 | -0.76 | -0.92 | -0.82 | -0.56 | -0.56 | -0.58 | -0.87 | -0.85 | -0.70 |
| $\varepsilon \quad 1988$ | -0.75 | -0.68 | -0.69 | -0.74 | -0.62 | -0.48 | -0.60 | -0.62 | -0.61 | -0.97 | -0.78 | -0.53 |
| E 1989 | -0.07 | -0.03 | 0.02 | 0.16 | 0.85 | 1.26 | 0.16 | 0.06 | 0.22 | 0.73 | 0.92 | 1.24 |
| $E \quad 1990$ | 0.20 | 0.35 | 0.28 | 0.30 | 0.33 | 0.30 | 0.26 | 0.33 | 0.21 | -0.61 | -0.41 | -0.19 |
| $\varepsilon \quad 1991$ | -0.45 | -0.60 | -0.66 | -0.70 | -0.33 | -0.09 | -0.63 | -0.72 | -0.72 | -1.19 | -0.93 | -0.59 |
| $\varepsilon \quad 1992$ | 0.30 | 0.37 | 0.33 | 0.59 | 1.14 | 1.35 | 0.53 | 0.50 | 0.49 | 0.26 | 0.82 | 1.27 |
| $\varepsilon 1993$ | 1.11 | 1.11 | 1.07 | 1.01 | 0.82 | 0.61 | 0.94 | 0.95 | 0.94 | 0.96 | 0.87 | 0.25 |
| $\varepsilon 1994$ | -0.04 | 0.01 | -0.04 | -0.40 | -0.58 | -0.34 | 0.15 | 0.03 | 0.01 | -0.89 | -0.61 | 0.24 |
| E 1995 | -0.36 | -0.37 | -0.36 | -0.27 | 0.22 | 0.52 | -0.02 | 0.01 | 0.00 | -0.35 | 0.05 | 0.25 |
| E 1996 | 0.16 | 0.10 | 0.05 | -0.04 | 0.31 | 0.52 | -0.36 | -0.26 | -0.27 | -0.11 | 0.18 | 0.46 |
| $\mathcal{E} \quad 1997$ | 0.22 | 0.41 | 0.39 | 0.87 | 1.03 | 0.95 | 0.74 | 0.72 | 0.70 | 0.36 | 0.72 | 0.73 |
| $\varepsilon 1998$ | 1.11 | 0.89 | 0.85 | 0.37 | 0.13 | 0.30 | 1.03 | 0.86 | 0.83 | -0.16 | -0.06 | 0.13 |
| $\varepsilon \quad 1999$ | -0.30 | -0.05 | -0.16 | -0.32 | -0.30 | -0.14 | -1.07 | -0.61 | -0.57 | 0.46 | 0.13 | 0.08 |
| $\varepsilon \quad 2000$ | -0.70 | -0.69 | -0.67 | -0.31 | 0.37 | 0.73 | -1.33 | -0.85 | -0.81 | -0.80 | 0.27 | 0.88 |
| $\varepsilon \quad 2001$ | -0.19 | -0.26 | -0.21 | 0.08 | -0.16 | -0.24 | 0.23 | 0.38 | 0.35 | -0.83 | -0.55 | -0.76 |
| $\varepsilon \quad 2002$ | 0.73 | 0.67 | 0.56 | -1.04 | -1.22 | -1.02 | 0.87 | -1.26 | -1.20 | -0.42 | -0.99 | -0.93 |
| Likelhoods |  |  |  |  |  |  |  |  |  |  |  |  |
| Like Bio | -63.29 | -37.98 | -34.20 | 13.80 | 80.80 | 90.23 | -62.12 | 5.39 | 2.97 | -10.47 | 63.27 | 99.04 |
| Like Survey LF | -2305.71 | -2355.60 | -2356.15 | -2417.18 | -2550.98 | -2627.02 | -2117.02 | -2293.78 | -2290.95 | -2384.71 | -2489.09 | $-2588.00$ |
| Like Com LF | -109.73 | -117.88 | -235.68 | -217.54 | -185.95 | -163.21 | -240.52 | -106.14 | -245.42 | -215.80 | -180.94 | -157.33 |
| Mprior | 3.77 | 3.82 | 3.82 | 2.43 | 1.56 | 3.02 | 3.22 | 3.27 | 3.30 | 2.40 | -1.81 | 2.21 |
| RecruitPrior | 17.23 | 16.79 | 15.49 | 19.49 | 30.94 | 36.55 | 26.80 | 22.73 | 21.40 | 29.88 | 28.34 | 30.54 |
| Total like | -2457.73 | -2490.85 | -2606.72 | -2599.00 | -2623.63 | -2660.43 | -2389.64 | -2368.52 | -2508.71 | -2578.70 | -2580.23 | -2613.53 |
| Indicators |  |  |  |  |  |  |  |  |  |  |  |  |
| B0 | 2628 | 2304 | 2384 | 2658 | 2842 | 2782 | 2306 | 2328 | 2356 | 3142 | 2994 | 2855 |
| BOrec | 1104 | 1471 | 1307 | 1543 | 1575 | 1956 | 1142 | 1104 | 1311 | 1639 | 1755 | 1912 |
| B2002rec | 1957 | 1136 | 1189 | 956 | 1104 | 1041 | 1229 | 930 | 951 | 820 | 979 | 1039 |
| B2002rec/B0rec | 1.77 | 0.77 | 0.91 | 0.62 | 0.70 | 0.53 | 1.08 | 0.84 | 0.73 | 0.50 | 0.56 | 0.54 |

## Lognormal error model fits

Under uniform weighting across all likelihoods, the model achieved a good fit to the biomass estimates (Figure 8). The estimated value of $M$ under uniform weighting was much higher than the mean of the prior ( 0.64 vs. 0.30 ), but was not as high as estimated by the normal error model and was not against its bound (Table 3). The lognormal error model's fit to the survey length frequency data was worse than that achieved by the normal error model under uniform weighting (Appendix 3 cf Appendix 8) and the fit was especially poor for the most recent surveys (2001 and 2002). Conversely, the lognormal error model reproduced some of the dominant cohort structure in the seasonal samples better than did the normal error model, and the fits to recent catch samples were good.

Unfortunately, increasing the relative weighting on the commercial and survey length frequency data resulted in a poorer fit to the biomass estimates (Figure 8) and did not markedly improve the fit to the length frequency data (although the 1:5:20 weighted model was almost able to reproduce the population of very large cockles in 1983, Appendix 9). Further, models weighted towards the length frequency data failed to converge, produced parameter values on their respective bounds (Table 3), and sometimes produced unstable, "saw tooth" patterns in the biomass. We selected a lognormal model with a weighting of 5 toward the survey length frequency distributions as the base case (1:1:5; Table 3).


Figure 9: Model fits to survey biomass estimates under increasing weighting toward commercial and survey length frequency data (weighting ratios biomass : commercial LF : survey LF)

## Lognormal error model base case diagnostics

As for the normal error model, natural mortality ( $M$ ) and mean recruitment (Rcoff) were correlated in the fitting process, though not as badly as in the normal error model ( 0.607 Appendix 5 b). Conversely, mean recruitment was highly correlated with a large proportion of the annual recruitment parameters (Appendix 5b) and there was higher correlation among recruitment parameters than in the normal error model (Appendix 5a).

The posterior distributions for the three selectivity parameters ( $\nu^{1 r}, \nu^{15}, \phi^{\text {diff }}$ ) were broad, suggesting that these parameters were not well estimated (Appendix $10 \mathrm{f}, \mathrm{g}, \mathrm{h}$ ). There was a high degree of auto correlation in MCMC chains (Appendix 11) and, as with the normal error model, convergence was not attained for several parameters, including some important ones (Appendix 7b). We conclude that this model, too, provides a poor description of the observed data, and our confidence that the estimated parameter values have biological meaning is, therefore, low.

## 11. Discussion:

### 11.1 General

Both normal and lognormal error models had problems rationalising the observed biomass and length frequency data; fits were obtained to one series at the expense of the fit to the other. There seems to be a fundamental conflict in the observed data, and this may point to the existence of an "unseen" or unaccounted mortality factor impacting upon the cockle population, or high variability of growth or mortality between years. Landings were relatively low and stable over the period of rapid decline in biomass in the 1980s, whereas reported landings increased rapidly as biomass increased in the early 1990s (Figures $6 \& 8$ ). Within both models, the only means of "shedding" or "gaining" the required biomass over such short times are through, respectively, implausibly high rates of natural mortality and improbably fast growth. That the normal error model was slightly better at finding a compromise between the observational data sets is not surprising because it has the capacity to remove biomass from the population rapidly by allowing a large proportion of harvestable cockles to shrink.

Using our base cases models, we estimate the current recruited biomass of cockles on Snake Bank to be slightly over $50 \%$ of virgin recruited biomass. However, the apparent incompatibility of data sets, the favouring of implausibly high rates of mortality and growth, and the lack of convergence inferred from MCMC chains are reasonably serious shortcomings of the models, and we have little confidence in any estimates of current stock status.

### 11.2 Future development of the model

We think that both normal and lognormal error models are over-parameterised; there seems to be insufficient information in the observed data to estimate all the parameters. Both models would benefit from the inclusion of independent data on growth (because growth and mortality are inextricably linked and the model frequently favoured implausible estimates for both). A notch tagging study is underway on Snake Bank and some recaptures have been made. Further tagging and recapture events should provide sufficient marginal increment data to include a new likelihood term for these data in the model. This should help constrain the model to more realistic values for natural mortality as well as growth.

Broad posterior distributions for the selectivity parameters in the lognormal error model suggest that these parameters were not well estimated although the base case lognormal model fits to the commercial length frequency distribution appear much better than those for the normal error model. More data are being collected on the length frequency distribution of the commercial catch in 2003 and this may help.

Since the underlaying model is expressed in terms of numbers at length, we suggest the population likelihood term would be better formulated as population numbers instead of its current formulation based on biomass.

Biomass is calculated in the current model using a length weight relationship without associated error. In fact, the length weight relationships estimated at different times have been different, and there is quite a lot of scatter around each. This may be a source of bias in the estimation and should be investigated.

We have coded an additional term for density dependent growth in the normal error model but because we were exploring other growth models and fits using density-dependent growth took a long time to converge, we have not explored this in detail. The rapid loss of biomass early in the time series and the maintenance of a relatively high biomass despite heavy fishing later in the time series are both intuitively consistent with some density dependent effect. We think this should be explored further in any model development.

Finally, if the reported landings used to drive the model are grossly inaccurate, modelling will be futile unless the actual catches can be estimated. Anecdotes suggest that landings may have been under-reported in some earlier years but, conversely, there is evidence that cockles have been gathered commercially from parts of the harbour other than Snake Bank. Thus, there is the potential for extractions to have been over-reported as well as under-reported. If refining the biological aspects of the model cannot remove the current inconsistencies among data sets and provide for robust parameter estimates, then the potential for mis-reported landings will have to be explored or modelling abandoned (in favour of a simpler "CAY" approach, perhaps).

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## 11. Conclusions:

1. A stochastic, dynamic, length-based, observation-error, time-series model for cockles on Snake Bank, Whangarei Harbour, is described. The model is driven by reported landings, and fitted using Bayesian techniques to estimates of absolute biomass and length frequency distributions from surveys, to length frequency samples from the commercial catch, and to seasonal length frequency distributions from part of the bank between 1992 and 1995.
2. Two alternative growth models are explored; one has normal error structure and allows negative growth whereas the other has log-normal error structure and does not allow negative growth. Within each error structure, a variety of weighting scenarios were explored, giving more or less relative weight to the different data sets. The selected "base cases" preferentially weight to survey length frequency distributions.
3. Both models seem over-prescribed and tests on MCMC chains suggested the models did not converge on estimates for several parameters, some of them important ones. There was also high correlation among some of the estimated parameters; in the growth parameters in the normal error model, and in the recruitment vectors in the lognormal error model.
4. To explain rapid decreases and increases in observed biomass, both models favour implausibly high rates of natural mortality and growth. Neither model could resolve the apparent incompatibility of the biomass and length frequency data sets, and the results were very sensitive to the relative weights accorded each data set.
5. Both base case models suggest that current recruited biomass is slightly higher than $50 \%$ of virgin recruited biomass. However, because neither model provides a reasonable description of the observed data, and may not have converged properly for several parameters, we are not confident that these estimates have biological meaning.

## 12. Publications:

None.

## 13. Data Storage:

Length and weight data from surveys are held on a secure, backed-up server at NIWA, Auckland and have been copied to the Empress database, beach.

## Appendix 1: Non estimable model parameters.

| RO | base number of annual recruits ( $=\exp (\ln (R 0)$ ) |
| :---: | :---: |
| $\beta_{k}^{R}$ | proportion of recruits that enter the $k$ th length class |
| $\tau_{t}^{R}$ | a switch based whether animal is recruited in period $t\left(\tau_{t}^{R}=1\right)$ or not ( $\tau_{t}^{R}=0$ ) |
| $\sigma_{\varepsilon}$ | assumed standard deviation of recruitment deviations in logarithmic space |
| $N_{k, t}$ | number of cockles in the $k$ th length class in period $t$ |
| $R_{k, t}$ | recruits to the model in the $k$ th length class in period $t$ |
| $\Delta l_{k}$ | expected seasonal growth increment for cockle in the $k$ th length class |
| $\sigma^{\Delta J_{k}}$ | standard deviation of the growth increment for cockle in the $k$ th length class |
| G | growth transition matrix |
| $B_{1}$ | biomass of cockles available to the fishery in period $t$ |
| $l_{k}$ | length of a cockle at the midpoint of the $k$ th length class |
| $w_{k}$ | average weight of a cockle at $l_{k}$ |
| $m_{k}$ | proportion of mature cockles at $l_{k}$ |
| $S_{t}$ | biomass of mature cockle in period $t$ |
| $C_{t}$ | total observed catch in period $t$ |
| $U_{1}$ | exploitation rate in period $t$ |
| $U^{\max }$ | maximum permitted exploitation rate |
| $S F_{k, t}$ | finite rate of survival from fishing for cockles in the $k$ th length class in period $t$ |
| $\hat{J}_{t}$ | predicted research diver survey index |
| $J_{1}$ | observed research diver survey index |
| $V_{k}^{r}$ | relative selectivity of research divers for cockles in the $k$ th length class |
| $\lambda_{k}^{r}$ | proportion of mixing the two normal curve for the research survey selectivity |
| $V_{k, 1}^{s}$ | relative selectivity of the commercial fishery for cockles in the $k$ th length class, in period $t$ |
| $\lambda_{k, t}^{s}$ | proportion of mixing two normal curve for the commercial fishery selectivity |
| $\phi_{t}$ | size of maximum selectivity for catch sampling selectivity function in period $t$ |
| $\sigma_{t}^{J}$ | the standard error of the estimate of research survey index in period $t$ |
| $\omega^{J}$ | relative weight assigned to the research diver survey index data set |
| $K_{t}^{s}$ | a relative weight for length frequency data from commercial catch sampling in period $t$ |
| $\hat{p}_{k, t}^{s}$ | predicted proportion-at-length in the $k$ th length class in period $t$ in commercial catch sampling |
| $p_{k, t}^{s}$ | observed proportion-at-length in the $k$ th length class in period $t$ in commercial catch sampling |
| $\omega^{s}$ | relative weight assigned to the commercial catch sampling length frequency data |
| $\kappa_{t}^{r}$ | a relative weight for length frequency data from research surveys in period $t$ |
| $\hat{p}_{k, t}^{r}$ | predicted proportion-at-length in the $k$ th length class in period $t$ in research surveys |
| $p_{k, t}^{r}$ | observed proportion-at-length in the $k$ th length class in period $t$ in research surveys |
| $\varpi^{r}$ | relative weight assigned to the research survey length frequency data |
| $L$ | likelihood |
| $\mu_{M}$ | mean of the prior distribution for $M$ |
| $\sigma_{M}$ | standard deviation of the prior distribution for $M$ |

Appendix 2: Priors and parameter bounds.

| Parameter | prior | mean | cv | lower bound | upper bound |
| :---: | :---: | :---: | :---: | :---: | :---: |
| $\ln (R 0)$ | uniform | - | - | 1.000 | 50.000 |
| $l^{R}$ | uniform | - | - | 2.000 | 20.000 |
| $\left(\sigma^{R}\right)^{2}$ | uniform | - | - | 0.200 | 100.000 |
| M | normal | 0.3 | 0.8 | 0.050 | 0.800 |
| $L_{\infty}$ | uniform | - | - | 20.000 | 50.000 |
| or gl0 | uniform | - | - | 1.000 | 20.000 |
| $K$ | uniform | - | - | 0.010 | 2.000 |
| or g30 | uniform | - | - | 0.000 | 2.000 |
| $\alpha$ | uniform | - | - | 0.001 | 1.000 |
| $\sigma_{M I N}$ | uniform | - | - | 0.001 | 5.000 |
| $v^{1 r}$ | uniform | - | - | 0.001 | 50.000 |
| $v^{1 s}$ | uniform | - | - | 0.001 | 50.000 |
| $\phi^{\text {diff }}$ | uniform | - | - | 0.000 | 1.000 |
| $\widetilde{\sigma}$ | uniform | - | - | 0.010 | 100.000 |
| $\varepsilon_{t}$ | normal | 0.0 | 0.4 | -2.300 | 2.300 |

Appendix 3: Normal error model fits to the survey and commercial length frequency data under equal weighting.


Fits to Seasonal samplling data










Fits to commercial shed sampling data





Appendix 4: Normal error model fits to the survey and commercial length frequency data under high weighting toward the survey length frequency data (weighting ratios biomass : com LF : surv LF 1:5:20).

FIte to Survey Length Frequency data
115220












Fits to Seasonal sampiling data









Fits to commercial shed sampling data




Appendix 5a: Normal error model base case (1:5:20) parameter values and correlations from autodifferentiation fitting process.


Appendix 5b: Lognormal error model base case (1:1:5) parameter values and correlations from autodifferentiation fitting process.

|  | value | cy | Rcont | algmatide | m | 0100 | g10 | Seloctuls | Seluctidith | Sevectich | Recslase Ex | Recstaply | Growthey |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Acolf | 19.043 | 0.004 | 1.000 |  |  |  |  |  |  |  |  |  |  |
| ergrametida | 0.107 | 0.021 | 0.018 | 1.000 |  |  |  |  |  |  |  |  |  |
| M | 0.657 | 0.040 | Hewn nemin | 0.025 | 1.000 |  |  |  |  |  |  |  |  |
| 900 | 0.539 | 0.046 | 0.094 | -0.002 | 0.262 | 1.000 |  |  |  |  |  |  |  |
| 910 | 5.911 | 0.024 | 0.085 | 0.087 | 0.448 | -0.046 | 1.000 |  |  |  |  |  |  |
| selectus | 1.571 | 0.558 | -0,040 | 0.000 | -0,086 | -0,081 | -0.005 | 1.000 |  |  |  |  |  |
| Selectitimis | 0.047 | 0.037 | -0.003 | -0.001 | -0.009 | 0.009 | -0.001 | -0.002 | 1.000 |  |  |  |  |
| seloctlif: | 0.230 | 0.451 | 0.011 | 0.002 | -0.018 | -0.103 | 0.003 | 0.000 | -0.357 | 1.000 |  |  |  |
| Recsixs Ex | 3.117 | 73.055 | 0.000 | 0.000 | 0.000 | 0.000 | 0.000 | 0.000 | 0.000 | 0.000 | 1,000 |  |  |
| Recsizever | 0.226 | 107.842 | 0.000 | 0.000 | 0.000 | 0.000 | 0.000 | 0.000 | 0.000 | 0.000 | -0.755 | 1.000 |  |
| Cromitcy | 1.000 | 0.000 | 0.000 | 0.000 | 0.001 | -0.002 | 0.002 | 0.000 | 0.000 | 0.000 | 0.000 | 0.000 | 1.000 |
| Cstalmin | 3.862 | 0.086 | -0, 484 | -0.012 | -0.446 | -0.179 | 0.888 | 0.001 | -0.024 | -0.079 | 0.000 | 0.000 | 0.001 |
| Epe 1802 | 0.510 | 0.336 | -0.740 | -0.008 | 0.020 | 0.084 | -0.002 | -0.000 | -0.004 | -0.023 | 0.000 | 0.000 | 0.000 |
| Eps 1509 | 0.008 | 7.082 | Wmacsomid | 0.018 | 0.282 | -0.061 | 0.301 | -0.003 | 0.002 | 0.020 | 0.000 | 0.000 | 0.001 |
| Epe 9 9e4 | 0.104 | 1.286 | -0,386 | -0.015 | 0.178 | 0.145 | 0.076 | -0.019 | -0,010 | -0.027 | 0.000 | 0.000 | 0.000 |
| Eps 1005 | -0.432 | -0.451 | -0. 185 | 0.019 | 0.053 | -0.054 | 0.079 | 0.005 | 0.007 | 0.010 | 0.000 | 0.000 | 0.000 |
| Epe 1908 | $-0.390$ | -0.733 | -0.087 | -0.008 | 0.096 | 0.047 | 0.051 | -0.007 | -0.002 | -0.004 | 0.000 | 0.000 | 0.000 |
| Epe 1907 | -0.564 | -0.441 | -0.147 | -0.004 | 0.083 | 0.048 | 0.047 | -0.008 | -0.002 | -0,006 | 0.000 | 0.000 | 0.000 |
| Epe 1060 | -0.624 | -0.414 | -0.173 | 0.012 | 0.003 | 0.001 | 0.005 | -0.001 | -0.002 | -0.008 | 0.000 | 0.000 | 0.000 |
| Epe 1000 | 0.060 | 4.841 | -0.196 | 0.003 | -0.127 | -0.132 | -0.041 | 0.010 | -0.013 | -0.001 | 0.000 | 0.000 | 0.000 |
| Epe 1090 | 0.332 | 0.503 | -0.285 | -0,022 | 0.076 | 0.136 | 0.056 | -0,001 | 0.007 | -0.012 | 0.000 | 0.000 | 0.000 |
| Epe 1901 | -0.722 | -0.247 | -0 204 | 0.036 | -0.005 | 0.038 | -0.064 | -0.023 | 0.008 | 0.006 | 0.000 | 0.000 | -0.001 |
| Epe 1092 | 0.498 | 0.145 | -0.814 | -0.028 | -0,317 | -0.010 | -0.196 | 0.003 | -0.004 | -0.016 | 0.000 | 0.000 | -0.001 |
| Epe 1909 | 0.950 | 0.098 | -0.497 | -0.015 | -0.033 | -0.078 | -0.049 | -0.011 | 0.051 | 0.006 | 0.000 | 0.000 | 0.000 |
| Epe 1904 | 0.030 | 5.445 | -0.260 | 0.014 | 0.107 | 0.156 | 0.170 | -0.024 | -0.051 | -0.018 | 0.000 | 0.000 | 0.000 |
| Epe 1905 | 0.007 | 11.040 | -0.393 | -0.009 | 0.237 | 0.130 | -0.003 | 0.016 | -0.003 | -0.009 | 0.000 | 0.000 | 0.000 |
| Epe 1808 | -0.026 | -0.285 | -0,607 | 0.008 | 0.044 | -0.045 | 0.257 | 0.001 | 0.027 | -0.006 | 0.000 | 0.000 | 0.001 |
| Epe 1007 | 0.716 | 0.120 | -0.729 | -0.017 | -0.421 | 0.012 | -0.305 | -0.047 | -0.011 | -0.013 | 0.000 | 0.000 | -0.001 |
| Epe 1008 | 0.855 | 0.097 | -0.511 | -0,012 | 0.149 | 0.081 | 0.200 | 0.024 | 0.020 | -0.006 | 0.000 | 0.000 | 0.001 |
| Ept 1980 | -0.014 | -0.299 | -0.035 | 0.025 | 0.336 | 0.044 | 0.252 | -0.017 | -0.007 | -0.004 | 0.000 | 0,000 | 0,000 |
| Epe 2000 | -0.850 | -0.184 | -0.279 | 0.019 | -0.051 | 0.009 | -0.110 | -0.007 | 0.000 | 0.000 | 0.000 | 0.000 | 0.000 |
| Epe 2009 | 0.384 | 0.192 | -0.659 | -0.033 | -0.041 | 0.101 | -0.043 | -0.017 | 0.003 | -0.015 | 0.000 | 0.000 | 0.000 |
| Eps 2002 | -1.261 | -0.174 | -0.110 | 0.077 | 0.120 | -0.011 | 0.229 | 0.016 | 0.002 | 0.010 | 0.000 | 0,000 | 0.001 |
|  |  | Epa 1802 | $\begin{gathered} \text { Epe } 1082 \\ 1.000 \end{gathered}$ | Epe 1800 | Epe 1984 | Epe 1505 | Epe 1500 | Epa 1907 | Ept 198 | Epe 1800 | Epe 1980 | Epe 1507 | Epe 1002 |
|  |  | Epr 1903 | -0.725 | 1.000 |  |  |  |  |  |  |  |  |  |
|  |  | Epe 1004 |  | -0.475 | 1.000 |  |  |  |  |  |  |  |  |
|  |  | Epe 1 ges | 0.275 | -0.155 | -0.481 | 1.000 |  |  |  |  |  |  |  |
|  |  | Epa 1005 | 0.189 | -0.100 | 0.181 | -0.005 | 1.000 |  |  |  |  |  |  |
|  |  | Epp 1007 | 0.257 | -0.150 | 0.230 | 0.004 | -0,406 | 1.000 |  |  |  |  |  |
|  |  | Epa 1 18e | 0.230 | -0.158 | 0.178 | 0.022 | -0.289 | -0.404 | 1.000 |  |  |  |  |
|  |  | Ept 1000 | 0.154 | -0.138 | 0.073 | 0.048 | 0.070 | 0.062 | -0,068 | 1000 |  |  |  |
|  |  | Epe 1090 | 0.422 | -0.269 | 0.288 | 0.119 | 0.103 | 0.128 | 0.082 | -0.759 | 1.000 |  |  |
|  |  | Epe 1009 | 0.287 | -0211 | 0.178 | 0.085 | 0.051 | 0.072 | 0.067 | 0.123 | -0.240 | 1.000 |  |
|  |  | Epe 1092 |  | -0.682 | 0.488 | 0.207 | 0.122 | 0.184 | 0.203 | 0.289 | 0.315 | -0.065 | 1.000 |
|  |  | Epp 1003 | [9m0980 | -0.448 | 0.394 | 0.182 | 0.115 | 0.150 | 0.148 | 0.100 | 0.259 | 0.231 | 0.509 |
|  |  | Epf 1904 | 0.374 | -0.211 | 0.278 | 0.095 | 0.084 | 0.108 | 0.085 | 0.040 | 0.180 | 0.086 | 0.280 |
|  |  | Epe 1095 | $\cdots$ | -0.468 | 0.524 | 0.211 | 0.162 | 0.210 | 0.173 | 0.087 | 0.329 | 0.219 | 0.550 |
|  |  | Eps 1096 | $0.780: 1$ | -0456 | 0.479 | 0.261 | 0.147 | 0.200 | 0.181 | 0.113 | 0.332 | 0.222 | $\left[{ }^{2 \times 8} 0.0040^{3 / 8}\right.$ |
|  |  | Epa 1097 | - 0,684 | . 0.647 | 0.378 | 0.149 | 0.088 | 0.140 | 0.157 | 0.158 | 0.256 | 0.216 | $\underline{L x} 50,728$ con |
|  |  | Epe 1098 | L. $+0.07455_{480}$ | -0.446 | 0.512 | 0.217 | 0.157 | 0.206 | 0.173 | 0.105 | 0.329 | 0.197 | 0.571 |
|  |  | Eps 1090 | 0.290 | -0.072 | 0.230 | 0.123 | 0.005 | 0.101 | 0.089 | 0.002 | 0.147 | 0.085 | 0.122 |
|  |  | Epe 2000 | 0.357 | -0.279 | 0.220 | 0.004 | 0.061 | 0.088 | 0.083 | 0.063 | 0.143 | 0.112 | 0.321 |
|  |  | Eps 2001 | [-4xame | .0.601 | 0.541 | 0.222 | 0.155 | 0.213 | 0.193 | 0.133 | 0.356 | 0.242 | \% |
|  |  | Epe 2002 | 0.184 | -0.051 | 0.120 | 0.088 | 0.046 | 0.058 | 0.046 | 0.013 | 0.092 | 0.051 | 0.101 |
|  |  | Epe 1094 | Epe 1094 1.000 | Eps 1906 | Epm 1998 | Epx 1007 | Ept 1008 | Epe 1009 | Epe 2000 | Ept 2000 | Ept 2002 |  |  |
|  |  | Epe 1295 | -0.092 | 1.000 |  |  |  |  |  |  |  |  |  |
|  |  | Ep 1006 | 0.364 | 0.355 | 1.000 |  |  |  |  |  |  |  |  |
|  |  | Epe 1007 | 0.184 | 0.454 | 0.390 | 1.000 |  |  |  |  |  |  |  |
|  |  | Epe 1080 | 0.321 | 0.575 |  | 0.184 | 1.000 |  |  |  |  |  |  |
|  |  | Epe 1009 | 0.138 | 0.279 | 0.296 | 0.178 | -0.118 | 1.000 |  |  |  |  |  |
|  |  | Epe 2000 | 0.111 | 0.267 | 0.251 | 0.265 | 0.341 | -0.325 | 1.000 |  |  |  |  |
|  |  | Epe 2001 | 0.315 |  |  | 4.40,422 | 2x\% 01818 | 0.223 | -0.029 | 1.000 |  |  |  |
|  |  | Epe 2002 | 0.095 | 0.143 | 0.227 | 0.027 | 0.178 | 0.139 | 0.105 | -0.006 | 1.000 |  |  |

Appendix 6: Normal model posterior distributions and traces (chain length $\mathbf{1 0} 000$ sampled from 1000000 MCMC steps)


a.

Log of recruitment $\boldsymbol{\operatorname { l n }}(\boldsymbol{R} 0)$


b. Length at first recruitment $l^{R}$



- trace
c. Natural Mortality $M$

d. Brody L infinity $L_{\infty}$


## Appendix 6 cont:



e. Brody K


- trace
f. shape parameter for the left hand limb of research survey selectivity $v^{1 r}$

g. shape parameter for the left hand limb commercial survey selectivity $v^{1 s}$

h. Annual difference in commercial selectivity mean $\phi^{\text {diff }}$

Appendix 7: Geweke convergence diagnostics for MCMC chains. The P-values were generated using two tail Z-tests of the null hypothesis $\left(\mathrm{H}_{0}\right)$ that the first $10 \%$ of the chain and the last $50 \%$ of the chain come from the same population. It is common practice to conclude that there is evidence against convergence when $\mathrm{P}<0.05$.
a. Normal error model base case

| Parameter | Z-Score | p -value | pass/fail |
| :---: | :---: | :---: | :---: |
| $\ln (R 0)$ | 3.461501 | 0.001 | fail |
| $l^{R}$ | -0.644644 | 0.519 | pass |
| M | 5.084667 | 0.000 | fail |
| $L_{\infty}$ or g30 | 0.147982 | 0.882 | pass |
| $K$ or gl0 | 5.818667 | 0.000 | fail |
| $v^{\text {Ir }}$ | -6.766396 | 0.000 | fail |
| $v^{1 s}$ | 5.674292 | 0.000 | fail |
| $\phi^{*}$ | 9.94504 | 0.000 | fail |
| E 1982 | 8.607849 | 0.000 | fail |
| $\varepsilon 1983$ | 3.088832 | 0.002 | fail |
| ع 1984 | -2.311027 | 0.021 | fail |
| E 1985 | -10.22158 | 0.000 | fail |
| ع 1986 | 7.360203 | 0.000 | fail |
| $\varepsilon \quad 1987$ | 8.584229 | 0.000 | fail |
| $\varepsilon \quad 1988$ | -0.443228 | 0.658 | pass |
| ع 1989 | -3.691601 | 0.000 | tail |
| $\varepsilon \quad 1990$ | 0.993183 | 0.321 | pass |
| E 1991 | -3.327398 | 0.001 | fail |
| ع 1992 | -5.932129 | 0.000 | tail |
| ع 1993 | 6.295206 | 0.000 | fail |
| E 1994 | -0.650703 | 0.515 | pass |
| E 1995 | -2.022365 | 0.043 | fail |
| E 1996 | -3.077236 | 0.002 | fail |
| ع 1997 | -5.132583 | 0.000 | fail |
| ع 1998 | -7.25622 | 0.000 | fail |
| ع 1999 | 10.96383 | 0.000 | fail |
| ع 2000 | -9.377436 | 0.000 | fail |
| $\varepsilon \quad 2001$ | 5.344583 | 0.000 | fail |
| $\varepsilon \quad 2002$ | -10.81582 | 0.000 | fail |
| B0 | 2.221315 | 0.026 | fail |
| Borec | -5.420128 | 0.000 | fail |
| B2002rec | 2.262185 | 0.024 | fail |
| B2002red/B0rec | 4.72718 | 0.000 | fail |

b. Lognormal error model base case

| Parameter | Z-Score | $p$-value | pass/fai |
| :---: | :---: | :---: | :---: |
| $\ln (\mathrm{RO})$ | 17.79603 | 0.000 | fail |
| $l^{R}$ | -1.430893 | 0.152 | pass |
| M | 13.20311 | 0.000 | fail |
| g30 | 0.798906 | 0.424 | pass |
| g10 | 1.771632 | 0.076 | fail |
| $\nu^{\text {tr }}$ | 6.256203 | 0.000 | fail |
| $\nu^{15}$ | -29.98975 | 0.000 | fail |
| $\phi^{*}$ | 36.7378 | 0.000 | fail |
| $\varepsilon \quad 1982$ | -9.855305 | 0.000 | fail |
| $\varepsilon \quad 1983$ | 4.51492 | 0.000 | fail |
| $\varepsilon \quad 1984$ | 0.02641 | 0.979 | pass |
| $\varepsilon \quad 1985$ | -10.44307 | 0.000 | fail |
| $\varepsilon \quad 1986$ | 6.697309 | 0.000 | fail |
| $\varepsilon \quad 1987$ | 11.42401 | 0.000 | fail |
| $\varepsilon \quad 1988$ | -16.4652 | 0.000 | fail |
| \& 1989 | 1.697791 | 0.090 | pass |
| $\varepsilon \quad 1990$ | -5.082495 | 0.000 | fail |
| $\varepsilon \quad 1991$ | -6.898729 | 0.000 | fail |
| $\varepsilon \quad 1992$ | -11.70173 | 0.000 | fail |
| $\varepsilon \quad 1993$ | -7.161302 | 0.000 | fail |
| E 1994 | -4.996716 | 0.000 | fail |
| $\varepsilon \quad 1995$ | -3.424094 | 0.001 | fail |
| $\varepsilon \quad 1996$ | -13.76289 | 0.000 | fail |
| \& 1997 | -14.17461 | 0.000 | fail |
| $\varepsilon 1998$ | -5.43286 | 0.000 | fail |
| $\varepsilon \quad 1999$ | 5.623511 | 0.000 | fail |
| $\varepsilon \quad 2000$ | -17.41529 | 0.000 | fail |
| $\varepsilon \quad 2001$ | -2.424601 | 0.015 | fail |
| E 2002 | -10.09547 | 0.000 | fail |
| B0 | 9.756857 | 0.000 | fail |
| B0rec | -11.17416 | 0.000 | fail |
| B2002rec | 5.061952 | 0.000 | fail |
| B2002rec/B0rec | 10.69674 | 0.000 | fail |

Appendix 8: Lognormal error model fits to the survey and commercial length frequency data under equal weighting.


Fits to Seasonal sampling data










Fits to commercial shed sampling data





Appendix 9: Lognormal error model fits to the survey and commercial length frequency data under high weighting toward the survey length frequency data (weighting ratios biomass : commercial LF : survey LF 1:5:20).


Fits to Seasonal sampling data









Fits to commerclal shed sampiling data





Appendix 10: Lognormal model posterior distributions and traces (chain length 10000 sampled from 1000000 MCMC steps).


a. Log of recruitment $\ln (R 0)$



b. Length at first recruitment $l^{R}$


c. $\quad$ Natural Mortality $M$

d. g 10


e. g 30


f. shape parameter for the left hand limb of research survey selectivity $\boldsymbol{v}^{1 r}$

g. shape parameter for the left hand limb commercial survey selectivity $v^{1 s}$

h. Annual difference in commercial selectivity mean $\phi^{\text {diff }}$

