

Age-structured model for Alaska herring stocks

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Abstract

1 Introduction

2 Model deconstruction

This section is intended to serve three purposes: 1) to document the model structure given the code in `model.tpl`, 2) to detail proposed changes to the model code to improve overall numerical stability, and 3) provide statistical approach that is amenable to maximum likelihood and Bayesian methods.

2.1 Model Structure

Table 1 begins with the objective function that is being minimized. There are four components defined in (T1.1), where three of the four components are scaled by user defined coefficients λ . The first component is the commercial age-composition data (QC), the second is the spawnig biomass age-composition (QS), the third is egg deposition data (WQE), and finally the fourth component is a penalty on the recruitment deviations from the underlying Ricker stock-recruitment model (QR).

For the commercial age-composition data, observation errors in the age-proportions are assumed to be normal (T1.2), where the predicted proportion-at-age (T1.3) is a function of the numbers-at-age (T1.4) and selectivity (T1.5). Note that in (T1.4) that the function is not continous. In this case the selectivity curve is rescaled to have a maximum value of 1.0. The `max` operation in the denominator of this function breaks the derivative chain in AD Model Builder and can result in numerical problems during parameter optimization associated with corrupt derivative information.

The same normal error distribution is also assumed for the age-proportions in the spawner catch composition samples (T1.6). In this case, the age proportions are based on the mature numbers-at-age at the time of spawning, where the fishhries removals are first subtracted from the mature numbers-at-age (T1.7). Note that this further assumes that all removals (i.e.,

fisheries selectivity) only harvests sexually mature fish. This assumption is inconsistent with (T1.4), where a different logistic curve is assumed for fisheries selectivity.

The catch-at-age data is internally derived in the model (T1.9) conditional on the numbers-at-age and the estimated selectivity. The model further assumes the total catch (in short tons) is measured without error. This is also referred to as conditioning the model on catch.

The residual sum of squares for the egg deposition survey is given by (T1.10). In this case the observation errors are assumed to be lognormal, and each year's observation is weighted by the inverse variance of sampling observation errors (φ_i).

Table 1: Decomposition of the objective function based on the source code provided in `model.tp1`. The objective function f is what AD Model Builder is trying to minimize. Note that $\dot{\cdot}$ represents mature state variables (e.g., mature weight-at-age \dot{w}_j)

Objective function		
$f = \lambda_C QC + \lambda_S QS + WQE + \lambda_R QR$		(T1.1)
Commercial catch proportion-at-age		
$QC = \sum_i \sum_j (\hat{Q}_{i,j} - Q_{i,j})^2$	$\hat{Q}_{i,j}$ observed proportions-at-age	(T1.2)
$Q_{i,j} = \frac{V_{i,j}}{\sum_j V_{i,j}}$	$Q_{i,j}$ predicted proportion-at-age	(T1.3)
$V_{i,j} = N_{i,j} \frac{S_{i,j}}{\max(S_{i,j})}$	$V_{i,j}$ vulnerable numbers-at-age	(T1.4)
$S_{i,j} = \frac{1}{1 + \exp(-g_i(j - a_i))}$	$S_{i,j}$ Selectivity in year i for age j	(T1.5)
Spawn proportion-at-age		
$QS = \sum_i \sum_j (\hat{O}_{i,j} - O_{i,j})^2$	$\hat{O}_{i,j}$ observed proportion-at-age	(T1.6)
$O_{i,j} = \frac{\dot{N}_{i,j}}{\sum_j \dot{N}_{i,j}}$	$O_{i,j}$ predicted proportion mature-at-age	(T1.7)
$\dot{N}_{i,j} = N_{i,j} M_{i,j} - C_{i,j}$	$\dot{N}_{i,j}$ Number-at-age at spawning	(T1.8)
$C_{i,j} = \frac{\hat{c}_i Q_{i,j}}{\sum_j Q_{i,j} w_j}$	\hat{c}_i observed catch, w_j weight-at-age	(T1.9)
Egg deposition survey		
$WQE = \sum_i \varphi_i [\ln(\hat{y}_i) - \ln(y_i)]^2$	\hat{y}_i observed egg deposition, φ_i weight	(T1.10)
$y_i = 0.5 \sum_j O_{i,j} (\rho_i \dot{w}_{i,j} - \alpha_i)$	ρ_i and α_i fecundity-weight regression	(T1.11)
Penalized recruitment deviations		
$QR = \sum_i^{(I-k)} \left[\ln(N_{i+k,k}) - \ln(f(\dot{N}_{i,j})) \right]^2$	$N_{i+k,k}$ number of age k recruits	(T1.12)
$f(\dot{N}_{i,j}) = a \dot{B}_i \exp(-b \dot{B}_i)$	Ricker stock-recruitment	(T1.13)
$\dot{B}_i = \sum_j \dot{N}_{i,j} \dot{w}_j$	\dot{B}_i mature biomass at spawning	(T1.14)

3 Methods

3.1 Input Data

3.2 Population dynamics

Estimated parameters for the population dynamics model include the initial abundance of ages 3-9+ for the initial year, abundance of age-3 recruits each year, and the natural mortality rate. In the original parameterization of the model, these initial recruitments and the vector of initial numbers-at-age result in creating $(N + A - 1)$ scaling parameters. To reduce the potential confounding with other global scaling parameters, updates to the model code include estimation of two recruitment scaling parameters, and two vectors of deviates that represent deviations from the mean. This modification reduces the potential for parameter confounding among the many parameters that affect global scaling (i.e., catchability coefficients, natural mortality rates).

Table 2: Notation and equations for population dynamics model.

Model parameters	
$\theta = \{\ln(M), \ln(\bar{R}), \ln(\ddot{R}), \ln(\alpha), \ln(\beta), \vec{\delta}\}$	(T2.1)
Initial States ($i = 1980$)	
$\iota_j = \begin{cases} \exp(-M * (j - \min(j))) & \text{where } 3 \leq j \leq 7 \\ \frac{\exp(-M * (j - \min(j)))}{1 - \exp(-M)} & \text{where } j = 8 \end{cases}$	(T2.2)
$N_{i,j} = \ddot{R} \exp(\delta_{i-j}) \iota_j$	$i = 1980, \forall j$ (T2.3)
$N_{i,j} = \bar{R} \exp(\delta_i)$	$\forall i, j = 3$ (T2.4)
Dynamic States ($i > 1980$)	
$Q_{i,j} = \frac{N_{i,j} S_{i,j}}{\sum_j N_{i,j} S_{i,j}}$	(T2.5)
$\bar{w}_i = \sum_j w_j Q_{i,j}$	(T2.6)
$C_{i,j} = \frac{\hat{c}_i Q_{i,j}}{\bar{w}_i}$	where \hat{c}_i is the observed catch (mt) (T2.7)
$\dot{N}_{i,j} = N_{i,j} \exp(-F_{i,j})$	(T2.8)
$N_{i+1,j+1} = \dot{N}_{i,j} \exp(-M_{i,j})$	(T2.9)

Table 3: Mathematical notation, symbols and descriptions.

Symbol	Description
<u>Index</u>	
g	group
h	sex
i	year
j	time step (years)
k	gear or fleet
l	index for length class
m	index for maturity state
o	index for shell condition.
<u>Leading Model Parameters</u>	
M	Instantaneous natural mortality rate
\bar{R}	Average recruitment
\dot{R}	Initial recruitment
α_r	Mode of size-at-recruitment
β_r	Shape parameter for size-at-recruitment
R_0	Unfished average recruitment
κ	Recruitment compensation ratio
<u>Size schedule information</u>	
$w_{h,l}$	Mean weight-at-length l
$m_{h,l}$	Average proportion mature-at-length l
<u>Per recruit incidence functions</u>	
ϕ_B	Spawning biomass per recruit
ϕ_{Q_k}	Yield per recruit for fishery k
ϕ_{Y_k}	Retained catch per recruit for fishery k
ϕ_{D_k}	Discarded catch per recruit for fishery k
<u>Selectivity parameters</u>	
$a_{h,k,l}$	Length at 50% selectivity in length interval l
$\sigma_{s_{h,k}}$	Standard deviation in length-at-selectivity
$r_{h,k,l}$	Length at 50% retention
$\sigma_{y_{h,k}}$	Standard deviation in length-at-retention
$\xi_{h,k}$	Discard mortality rate for gear k and sex h

References