

Stock Reduction Analysis using catch at length data

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Abstract

Last thing to be written

1 Introduction

Modern stock assessments typically attempt to fit population dynamics models to catch at age and catch at length data, in hopes of extracting information from these data about age/size vulnerability and fishing mortality patterns (Methot and Wetzel, 2013; Hilborn and Walters, 1992). In cases where age data are lacking, models like MULTIFAN-CL attempt to obtain estimates of vulnerability and fishing mortality only from size distribution data (Fournier et al., 1998). Combined with a few assumptions regarding the structure and variability in length at age, this procedure can even be used to attempt to recover information about changes in body growth patterns if there is a strong age-class signal in the length frequency data (Fournier et al., 1998). Some assessment methods attempt to put aside the length frequency data, by converting these data to age compositions using age-from-length tables, perhaps using iterative methods to estimate proportions of fish at age for each length interval (Kimura and Chikuni, 1987). It is typical for assessment results from length-based assessment models to show substantial deviations between predicted and observed length distributions of catches, reflecting both sampling variation in the length composition data and incorrect assumptions about stability of growth and vulnerability patterns (Hilborn and Walters, 1992).

The vulnerability process is the combination of two processes: selectivity of the fishing gear and availability of the fished population in the area being fished (Beverton and Holt, 1957). Both processes can vary over time and therefore modify the resulting selectivity. Although selectivity process can often be directly measured through gear experiments, availability is generally harder to measure as it depends on the distribution of the exploited population. Fish movement, changes in fish distribution, and changes in fleet distribution, can all

25 cause availability changes.Changes in vulnerability are not uncommon (Sampson and Scott, 2012) but are
 26 usually difficult to track over time. This difficulty is associated with an inability to distinguish between changes
 27 in fishing mortality and changes in vulnerability in most age and length based stock assessment methods. For
 28 this reason, many assessment methods rely on ad hoc parametric vulnerability models that may or may not
 29 include changes over time (Maunder et al., 2014). If misspecified, such models might lead to severe bias in
 30 fishing mortality estimates, which could result in misleading management advice (Martell and Stewart, 2014).

31 Here we suggest an alternative approach to assessment modeling that begins by assuming that the assess-
 32 ment model should exactly reproduce the observed catch at length distribution. This approach follows the
 33 dynamics of an age structured stock reduction analysis (SRA) (Walters et al., 2006) which follows a “condi-
 34 tioned on catch” format, subtracting observed catches at age from modeled numbers at age in estimation of
 35 numbers at age over time. This assumption is analogous to the classical assumption in virtual population anal-
 36 ysis that reconstructed numbers at age should exactly match observed catch at age data (Hilborn and Walters,
 37 1992). The suggested approach may have two key advantages over statistical catch at age and/or catch at length
 38 models: (1) it does not require estimation of age or size vulnerability schedules, and (2) catch at length data are
 39 commonly available for every year, even when age composition sampling has not been conducted.

40 We demonstrate the performance of this model with a simulation-evaluation analysis and apply it to real
 41 fisheries data from the Peruvian jack mackerel (*Trachurus murphyi*) and Pacific Hake (*Merluccius productus*)
 42 fisheries.

43 2 Methods

44 In this section we describe the stock reduction analysis with catch at length data (Length-SRA), describe
 45 the simulation analysis and scenarios used to test the model and provide a description of the real data used to
 46 illustrate the model applicability.

47 2.1 Stock reduction analysis with catch at length data

48 The stock reduction analysis described here starts by calculating the proportions of individual at length
 49 given each age class ($P_{l|a}$ - eqs. T2.1-T2.5). The calculation of such proportions (eq. T2.1) relies on four main
 50 assumptions regarding the distribution of length at age: (1) The mean length at age follows a von Bertalanffy
 51 growth curve (eq.T2.4), (2) The length at age is normally distributed (eqs. T2.1 -T2.3), (3) The standard

deviation of the length at age distributions is defined (e.g. eq.T2.5) and (4) $P_{l|a}$ is constant for all lengths equal or greater than L (eq.T2.3).

The proportions of length at age is used to convert the length-based quantities into age based quantities which are used to propagate the age structured population dynamics forward (Table 2 - Population dynamics). We assume that recruitment follows a Beverton-Holt type recruitment curve (eq. T2.6), that harvesting occurs over a short, discrete season in each time step (year or shorter time period) and that natural survival rate is stable over time (eqs. T2.6-T2.10). The computation of numbers at age in the initial years is different from that in the remaining years (eq. T2.11). The calculations for this year rely on the parameter R_{init} , which assumes that the population is not at unfished equilibrium at the beginning of the time series.

The model estimates three main parameters: average unexploited recruitment R_0 , recruitment compensation ratio κ and recruitment in the initial year R_{init} (we assume that the data collection for the fisheries does not start when the population is at an unfished equilibrium state). In addition, the recruitment deviations w_t are estimated for all cohorts observed in the model, that is, the number of recruitment deviations is equal to the number of years in the time series plus the number of age classes greater than recruitment age.

The objective function (eq. T3.8) is composed of a negative log likelihood component, two penalties and a prior component for the recruitment compensation ratio κ . The negative log likelihood component minimizes the differences between the predicted and observed index of abundance (eq. T3.1). We assume that such differences are lognormally distributed (eqs. T3.3-T3.4) and use the conditional maximum likelihood estimator described by Walters and Ludwig (1994) to estimate the catchability coefficient (eq. T3.2). A lognormal penalty is added to the negative log-likelihood function to constrain annual recruitment residuals so that the estimates have mean of zero and fixed standard deviation σ_R (eq. T3.5). An additional penalty was implemented in order to prevent estimated values of exploitation rate at length greater than one ($U_{l,t} > 1$) (eq. T3.6). Lastly, a normal prior for $\log(\kappa)$ was included in the objective function (eq. T3.7).

2.2 Simulation evaluation

In order to perform a simulation evaluation of the Length-SRA under various scenarios we used the same model dynamics described in Table 2 - Population dynamics, as an operating model. However we modified the model population dynamics to control annual exploitation rate (eq. T2.17), time varying selectivity (eq. T2.19), and observation and process errors. Selectivity in the operating model was computed with the three parameter selectivity equation described by Thompson (1994) (eq. T2.19). The observation error in the operating model

Table 1: Indexes, variable definition and values used in simulation-evaluation

Symbol	Value	Description
l	$\{l_o, \dots, L\}$	Central point of length bin, $L = 15$
a	$\{a_o, \dots, A\}$	Age-class, $A = 10$
t	$\{1, \dots, T\}$	Annual time step, $T = 50$
a_o	21	First age or age of recruitment
l_o	1	Central point of first length bin
$init$	21	Annual time step in which data starts to be reported
Distribution of length given age		
L_{inf}	10	Maximum average length
k	0.3	Rate of approach to L_{inf}
t_o	-0.1	Theoretical time in which length of individuals is zero
cv_l	0.08	Coefficient of variation for length at age curve
$P_{l a}$		Matrix of proportions of length at age
Φ		Standard normal distribution
$z1_{a,l}$		Normalized z score for lower limit length bins
$z2_{a,l}$		Normalized z score for upper limit length bins
$b1_l$		Lower limit of length bins
$b2_l$		Upper limit of length bins
\bar{L}_a		Mean length at age
σ_{L_a}		Standard deviation of length at age
Population dynamics		
R_o	100	Average unfished recruitment
κ	10	Goodyear recruitment compensation ratio
S	0.7	Natural survival
σ_{rec}	0.4	standard deviation for recruitment deviations
w_t	$\mathcal{N}(0, \sigma_R)$	Recruitment deviations for years $\{init-A+1, \dots, T\}$
$N_{a,t}$		Numbers of fish at age and time
SB_t		Spawning biomass at time
mat_a		Proportion of mature individuals at age
a_{rec}, b_{rec}		Beverton & Holt stock recruitment parameters
R_{init}		Recruitment in the year data starts to be reported
$U_{a,init}$		Exploitation rate at age before data starts being reported.
$U_{a,t}$		Exploitation rate at age and time
$U_{l,t}$		Exploitation rate at length and time
$C_{l,t}$		Catch at length and time
$N_{l,t}$		Numbers at length and time
lx_a		Unfished survivorship at age
ϕ_e		Unfished average spawning biomass per recruit
Operating model		
$sel_{l,t}$		Fishing selectivity at length and time
g, a, b		Parameters for selectivity function
U_t		Annual maximum exploitation rate
$C_{l,t}$		Catch at length and time
$N_{l,t}$		Numbers at length and time
I_t		Index of abundance at time
VB_t		Biomass that is vulnerable to the survey at time
q	1.0	catchability coefficient

Table 2: population dynamics for Length-SRA and operating model

Distribution of length given age	
(T2.1)	$P_{l a} = \int_{z1_{a,l}}^{z2_{a,l}} \Phi(z) dz$
(T2.2)	$z1_{a,l} = \frac{b1_l - \bar{L}_a}{\sigma_{L_a}}$
(T2.3)	$z2_{a,l} = \begin{cases} \frac{b2_l - \bar{L}_a}{\sigma_{L_a}} & l < L \\ 1.0 & l = L \end{cases}$
(T2.4)	$\bar{L}_a = L_{inf} \cdot (1 - \exp^{(-k \cdot (a - t_o))})$
(T2.5)	$\sigma_{L_a} = \bar{L}_a \cdot cv_l$
Population dynamics	
(T2.6)	$N_{a,t>init} = \begin{cases} \frac{a_{rec} \cdot SB_{t-1}}{1 + b_{rec} \cdot SB_{t-1}} \cdot e^{w_t}, & a = a_o \\ N_{a-1,t-1} \cdot S \cdot (1 - U_{a-1,t-1}), & a_o < a < A \\ \frac{N_{a-1,t-1} \cdot S \cdot (1 - U_{a-1,t-1})}{1 - S \cdot (1 - U_{a,t})}, & a = A \end{cases}$
(T2.7)	$U_{a,t} = \sum_a (P_{l a} \cdot U_{l,t})$
(T2.8)	$U_{l,t} = \frac{C_{l,t}}{N_{l,t}}$
(T2.9)	$N_{l,t} = \sum_a (P_{l a} \cdot N_{a,t})$
(T2.10)	$SB_t = \sum_a (mat_a \cdot w_a \cdot N_{a,t})$
Initial year and incidence functions	
(T2.11)	$N_{a,init} = \begin{cases} R_{init} \cdot e^{w_{init}} & a = a_o \\ N_{a-1,init} \cdot S \cdot (1 - U_{a-1,init}) \cdot e^{w_{init}-a+1}, & a_o < a < A \\ \frac{N_{a-1,init} \cdot S \cdot (1 - U_{a-1,init})}{1 - S \cdot (1 - U_{a,init})} \cdot e^{w_{init}-a+1}, & a = A \end{cases}$
(T2.12)	$a_{rec} = \frac{\kappa}{\phi_e}$
(T2.13)	$b_{rec} = \frac{\kappa - 1}{R_o \cdot \phi_e}$
(T2.14)	$\phi_e = \sum_a lx_a$
(T2.15)	$lx_a = \begin{cases} 1, & a = 1 \\ lx_{a-1} \cdot S, & 1 < a < A \\ \frac{lx_{a-1} \cdot S}{1 - S}, & a = A \end{cases}$
Operating model	
(T2.16)	$N_{a,t=1} = lx_a * R_o$
(T2.17)	$U_{l,t} = U_t \cdot sel_{l,t}$
(T2.18)	$C_{l,t} = N_{l,t} \cdot U_{l,t} \cdot P_{l a}$
(T2.19)	$sel_{l,t} = \frac{1}{1 - g} \cdot \left(\frac{1 - g}{g} \right)^g \cdot \frac{e^{a \cdot g \cdot (b-l)}}{1 + e^{a \cdot (b-l)}}$
(T2.20)	$I_t = q \cdot VB_t \cdot e^{(\mathcal{N}(0, \sigma_t))}$

Table 3: Likelihood functions and penalties

Conditional Likelihood	
(T3.1)	$Z_t = \log(I_t) - \log(VB_t)$
(T3.2)	$q = e^{\bar{Z}}$
(T3.3)	$Z_{stat} = Z_t - \bar{Z}$
(T3.4)	$Z_{stat} \sim \mathcal{N}(0, \sigma_t)$
Penalties	
(T3.5)	$w_t \sim \mathcal{N}(0, \sigma_R)$
(T3.6)	$pen(U) = \frac{\sum^t \sum^l U_{l,t}}{T}$
Priors	
(T3.7)	$prior(\log(\kappa)) \sim \mathcal{N}(\log(\kappa), 0.9)$
Objective function	
(T3.8)	$Obj = negLL(Z_{stat}) + negLL(w_t) + pen(U) + prior(\log(\kappa))$

included lognormal error in the index of abundance and logistic multivariate error in the catch numbers at length. Recruitment deviations were assumed to be lognormally distributed.

We considered a total of six different scenarios in the simulation evaluation runs. Three different historical exploitation rate trajectories were used: contrast, one way trip and U ramp. In the contrast scenario the exploitation rate (U_t) starts low and increases beyond U_{msy} and then decreases until $U_t = U_{msy}$. In the one way trip scenario U increased through time until $U = 2 \cdot U_{msy}$. In the U ramp scenario, U_t increases steadily until $U_t = U_{msy}$ and remains constant thereafter. In addition to the exploitation rate scenarios, we considered two selectivity scenarios: constant and time varying selectivity. In the constant selectivity scenario, selectivity was assumed to follow a sigmoid shape. In the time varying selectivity scenario, the selectivity curve was assumed to vary every year, progressively changing from a dome shaped curve to sigmoid and back to dome shaped. The complete list of scenarios and the acronym used for them is presented in Table 4.

All simulation runs had 30 years of data and we used 100 simulation runs for each scenario. We evaluated the distribution of the % relative error ($\frac{estimated-simulated}{simulated}$) for the main parameter estimates (R_0 , R_{init} and κ) and for four derived quantities (Depletion: $\frac{B_t}{B_0}$, MSY , U_{MSY} and q).

2.3 Real data examples

Two species were chosen to illustrate the application of the Length-SRA to real datasets: Chilean jack mackerel and Pacific hake. Both species are believed to be subject to time varying selectivity.

Table 4: Simulation-estimation scenarios

Scenario Code	Selectivity	U trajectory
CC	constant	contrast
CO	constant	one way trip
CR	constant	U ramp
VC	time-varying	contrast
VO	time-varying	one way trip
VR	time-varying	U ramp

The Pacific hake fishery is believed to exhibit time varying selectivity due to cohort targeting and annual changes fleet spatial distribution. The population is know to have spasmodic recruitment, with high recruitment events occurring once or twice every decade (Ressler et al., 2008). Pacific hake tends to segregate by size during their annual migration(Ressler et al., 2008), allowing the fishing fleet to target the strong cohorts by changing the spatial distribution of fishing effort as the cohort ages.

The movement pattern of jack mackerel is not as well known, although fish appear to move between spawning and feeding areas (Gerlotto et al., 2012). Variability in selectivity patterns for the jack mackerel fishery are believed to be associated both with evolution of fleet capacity and gear utilization and with compression and expansion of the species range associated with abundance changes (Gerlotto et al., 2012).

3 Results

3.1 Simulation-evaluation

Even though the parameter estimates were not entirely unbiased, the amount of bias was relatively low, especially for the leading parameters R_0 and κ (Figure 1). The parameter R_0 was slightly overestimated or accurately estimated for the contrast and one-way-trip scenarios, with median % errors between 0% and 2.3%. In the U ramp scenarios, R_0 was underestimated with with median % errors of -10.8% (CR) and -7.4% (VR). The parameter R_{init} was underestimated for all scenarios with relatively high median % errors varying between -31.6% and -35.9%. Once again the U ramp scenarios produced the highest bias. (Figure 1)The κ parameter was the most accurately and precisely estimated with median % errors varying betwewen 0.0% and 1.8%.

In relation to the derived quantities (Figure 2), the Length-SRA model produced estimates of depletion with median % errors ranging between -13.6% and 7.5%, there was no clear pattern among scenarios that lead of overestimation of underestimation. In contrast, MSY was underestimated for all scenarios, with median % errors ranging between - 28.1% and -12.1%. The estimates for U_{MSY} showed very low (<2%) median % errors for all

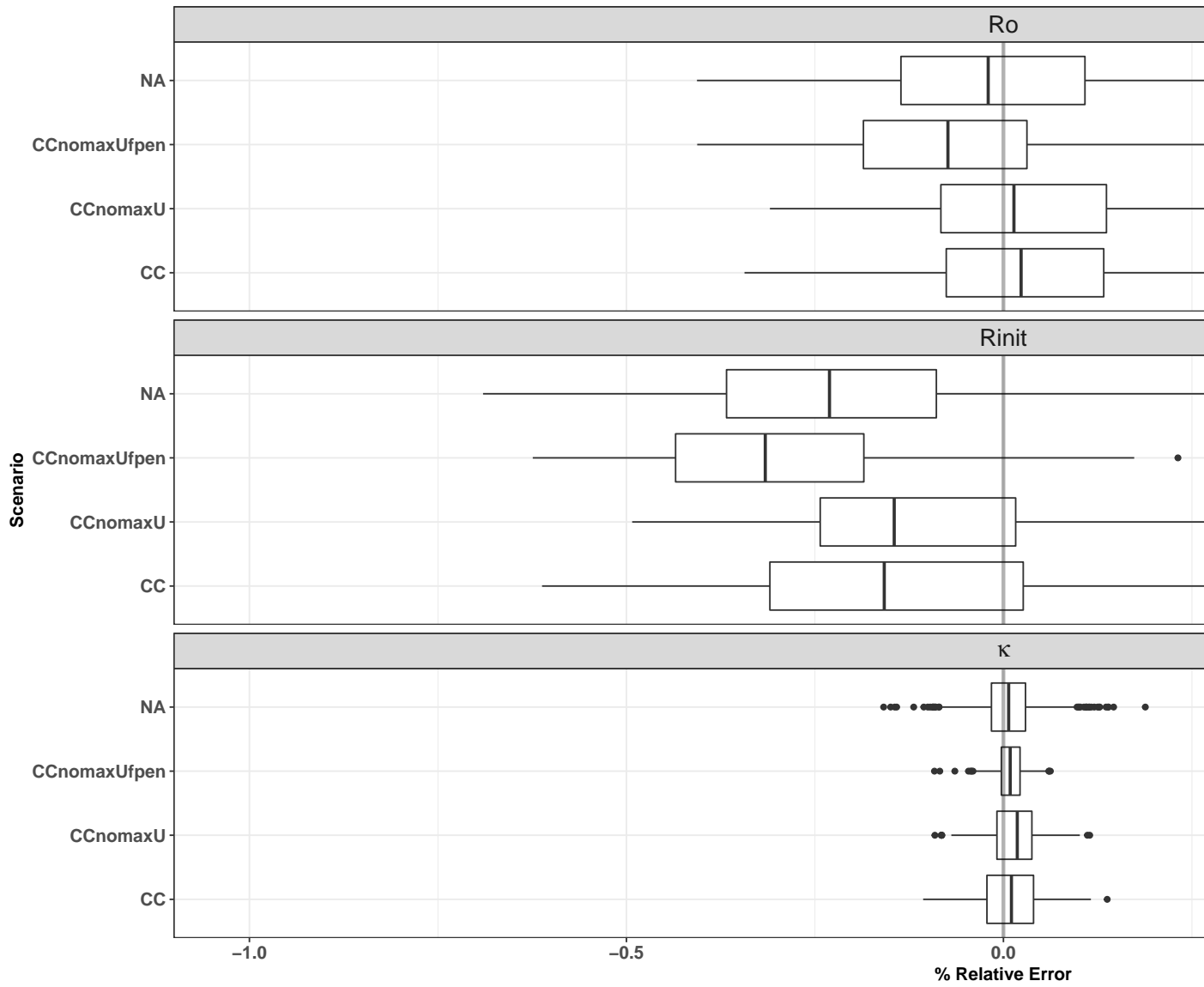


Figure 1: Relative error (%) for main parameters for all scenarios considered in the simulation-evaluation.

120 scenarios. The estimates of q were particularly positively biased for the the U ramp scenarios (CR and VR),
 121 this is likely to be associated with the underestimations of both R_0 and R_{init} for those scenarios.

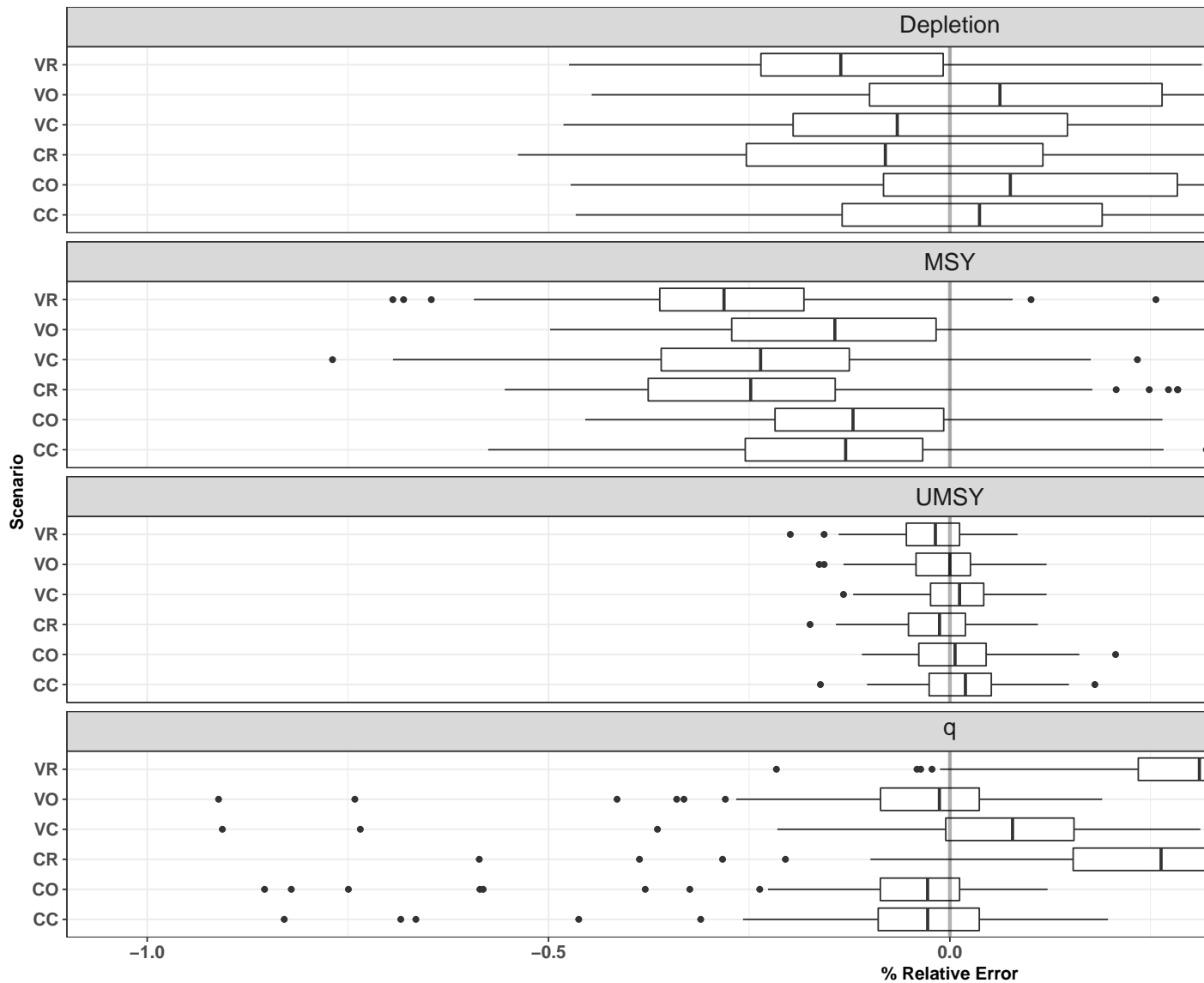


Figure 2: Relative error (%) for main parameters for all scenarios considered in the simulation-evaluation.

122 The simulation-evaluation exercise showed that the Length-SRA model is able to track selectivity changes
 123 over time relatively well (Figure 3).

124 3.2 Real data examples

125 show figures with real

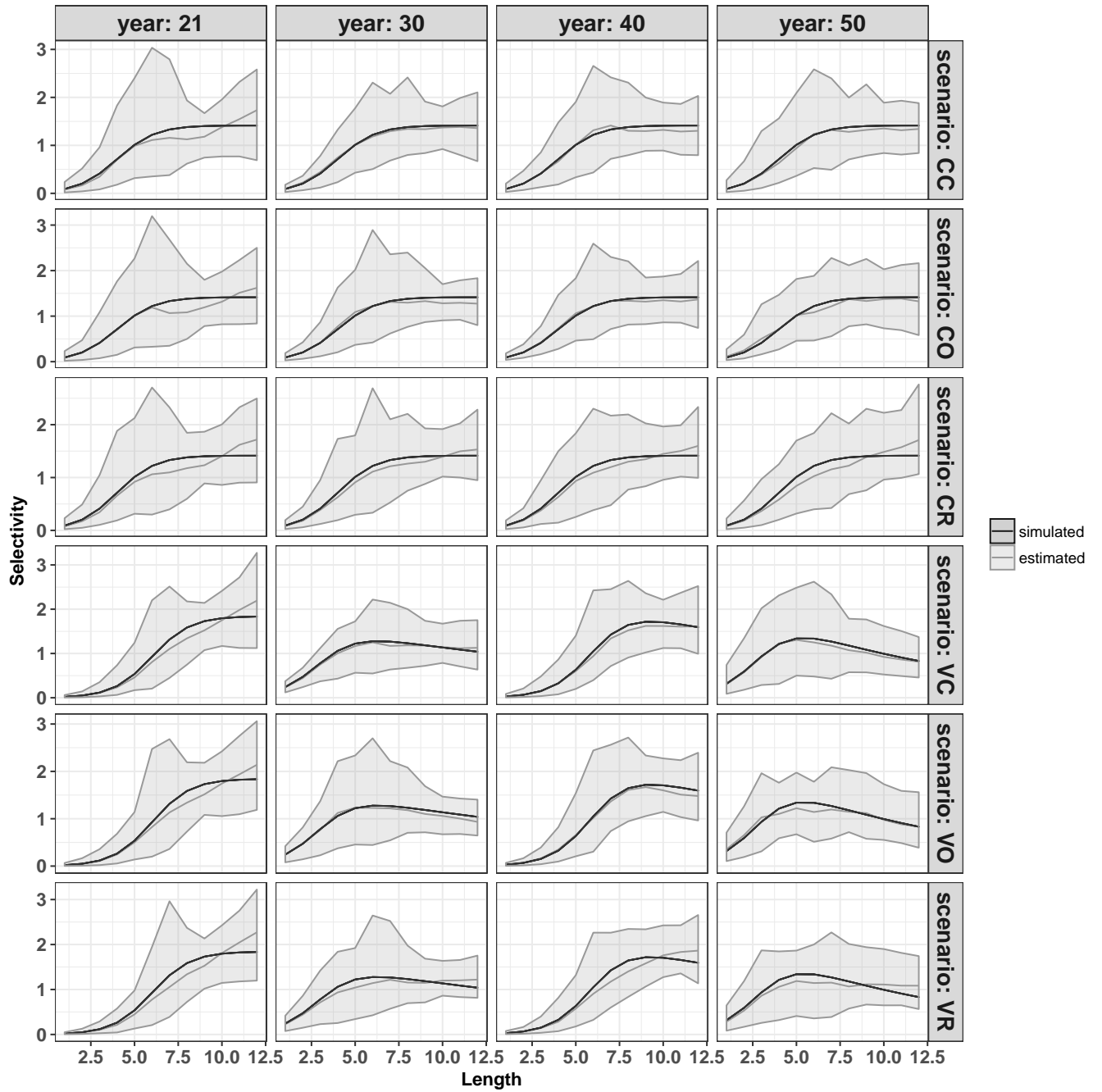


Figure 3: Simulated and realized selectivity estimates for a set of years within simulation-evaluation time series.

4 Discussion

We present a length-based stock reduction analysis that allows monitoring of time varying selectivity. In this model catch at length is assumed to be known without error and exploitation rate at length is derived directly from the estimates of numbers at length. This fact is important because it allows the mode to bypass the requirement for the estimation of a selectivity ogive, as is required in other length based models (Sullivan et al., 1990; Fournier et al., 1998).

Selectivity parameters can be particularly hard to estimate, especially it is changing over time and main conclusions: Length-SRA underestimates kappa productivity parameter Management quantities Depletion and MSY are underestimated - the model tends to produce conservative benchmarks.

UMSY estimates had low bias- good?

U ramp scenario yield very bad results - lack of information in the time series.

Model is able to track selectivity over time, - good

How does it compare to other length based models

Does the model work? Could it produce useful management advice? My nee dto evaluate in a closed loop frameworks to be certain.

assumption regarding Umsy $-l_i$ changes with selectivity $-l_i$

Time- varying growth might render the model less useful. Suggest estimating cohort-specific growth curves or implementing the density dependent functions shown in multifan-CL.

Further testing of this model in a closed-loop simulation set up would provide more insight into how

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