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A stochastic surplus production model in continuous time

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Abstract

Surplus production modelling has a long history as a method for managing datalimited fish stocks. Recent advancements have cast surplus production models as state-space models that separate random variability of stock dynamics from error in observed indices of biomass. We present a stochastic surplus production model in continuous time (SPiCT), which in addition to stock dynamics also models the dynamics of the fisheries. This enables error in the catch process to be reflected in the uncertainty of estimated model parameters and management quantities. Benefits of the continuous-time state-space model formulation include the ability to provide estimates of exploitable biomass and fishing mortality at any point in time from data sampled at arbitrary and possibly irregular intervals. We show in a simulation that the ability to analyse subannual data can increase the effective sample size and improve estimation of reference points relative to discrete-time analysis of aggregated annual data. Finally, subannual data from five North Sea stocks are analysed with particular focus on using residual analysis to diagnose model insufficiencies and identify necessary model extensions such as robust estimation and incorporation of seasonality. We argue that including all known sources of uncertainty, propagation of that uncertainty to reference points and checking of model assumptions using residuals are critical prerequisites to rigorous fish stock management based on surplus production models.

Keywords Data-limited methods, fisheries management, maximum sustainable yield, Pella–Tomlinson model, seasonal population dynamics, stock assessment

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Introduction

Surplus production models are used to assess the biomass and exploitation level of marine populations in data-limited situations where age and size information are unavailable (Punt 2003). By modelling the temporal evolution of the aggregated biomass targeted by fishing, surplus production models do not require information about the age or size-structure of the stock. Surplus production models are therefore primarily applied to stocks where the only data available are observations of commercial catches together with some index of exploitable biomass such as commercial catch-perunit-effort (CPUE) or as derived from scientific survey data (Polacheck et al. 1993). Common applications of surplus production models include large migratory pelagic fishes such as tuna, sharks and billfishes (e.g. Brodziak and Ishimura 2011; Carvalho et al. 2014), but also crustaceans that are generally difficult to age (e.g. Smith and Addison 2003).

The population dynamics represented by surplus production models builds on principles of logistic or the more general theta-logistic growth (Pedersen et al. 2011) resulting in a dome-shaped relationship between biomass production population size. In the formulation of Schaefer (1954), the maximum production occurs when the population size is at half its carrying capacity. This restriction is avoided in the generalized form of Pella and Tomlinson (1969), where skewness of the production function allows maximum production at any biomass below the carrying capacity. Thus, a population is optimally exploited in terms of biomass if it is harvested to remain at the biomass level that results in the maximum production defined as the maximum sustainable yield (MSY).

It is widely recognized that the model structure of surplus production models is too simple to adequately describe the population dynamics of a real-world stock subject to variability in size-structure, species interactions, recruitment, catchability, selectivity, environmental conditions, etc. (Pella and Tomlinson 1969). To mitigate this, it is common to include a random error term in the equation governing the biomass dynamics as a proxy for unmodelled variability (process error). Similarly, it is often assumed that the biomass index is subject to error in sampling that causes the observed values to deviate from the true. This variability is incorporated by including an observation-error term in the equation describing how the index data relate to the biomass. Models including random terms are often referred to as stochastic.

While the basic deterministic formulation of surplus production models has remained unchanged for decades, the methods used for estimating model parameters from observations while acknowledging random variability have undergone major developments. Least-squares methods for fitting production curves (Pella and Tomlinson 1969) developed into more explicit assignment of noise in process-error or observation-error models (Polacheck et al. 1993). Key to the incorporation of random variability was the discrete-time formulation of Schnute (1977) achieved by approximating observations by annual averages. The majority of subsequent model extensions, with the exception of ASPIC (Prager 1994), adopted the discrete-time form including state-space models that simultaneously estimate both process and observation error using Bayesian (Meyer and Millar 1999) or frequentist approaches (Punt 2003). ASPIC, while free of discrete-time average approximations, lacks flexibility owing to its deterministic population dynamics. Regardless of temporal form, observation error in previous models refers to error in the index observation with the often unstated implicit assumption that catch observations are error free.

The increased flexibility of state-space models, which are the variants of surplus production models most commonly applied today, entails a need for informative data. Thus, data situations with short time series or limited contrast may require that some model parameters are constrained or fixed to obtain model identifiability. For example, the ratio of process to observation noise is often fixed (Ono et al. 2012), while estimating the shape of the production curve is commonly avoided by assuming the symmetric form of Schaefer (1954). Constraining estimation by fixing model parameters affects estimates of remaining free parameters, some of which are directly related to management quantities. Thus, assessing the sensitivity of results to assumed constraints is an important aspect of surplus production modelling, which if ignored may lead to poor management.

Stochastic surplus production models aim to account for autocorrelation inherent in time-series data of catches and biomass indices. This implies that the recursive, or so-called one-step-ahead, residuals of a model fit are assumed to be standard normally distributed and independent. While this is a well-known property of many time-series models (Ljung and Box 1978), assessing the validity of these assumptions is often ignored when fitting stochastic surplus production models. This is problematic as residual patterns can indicate violation of model assumptions, which potentially invalidates model estimates and associated confidence intervals thus conveying a misleading impression of stock status.

We present a stochastic surplus production model in continuous time (SPiCT), which incorporates dynamics in both biomass and fisheries and observation error of both catches and biomass indices. The model has a general state-space form that as special cases contain process and observation-error models (Polacheck et al. 1993; Prager 1994) as well as state-space models that assume error-free catches (Meyer and Millar 1999; Punt 2003: Ono et al. 2012). Seasonal extensions to the fisheries dynamics component of the statespace model are also developed facilitating the use of subannual data that contain seasonal patterns. Estimation performance, as a function of the number of available observations and under model misspecifications, is evaluated through simulation experiments. Differences in performance between discrete-time and continuous-time models are assessed both in a simulation experiment and by

analysing the South Atlantic albacore data set of Polacheck *et al.* (1993) and comparing with previously published results from discrete-time models. Finally, we analyse five North Sea stocks using quarterly resolved data with emphasis on the use of one-step-ahead residuals for detecting possible lack of model fit.

Methods

Model formulation

Generalized surplus production models in the form of Pella and Tomlinson (1969) can be written as

$$\frac{dB_t}{dt} = \frac{r}{n-1} B_t \left(1 - \left[\frac{B_t}{K} \right]^{n-1} \right) - F_t B_t \qquad (1)$$

where B_t is the exploitable stock biomass (ESB), F_t is the instantaneous fishing mortality rate, r is the intrinsic growth rate of the population, K is the carrying capacity, and n > 0 is a unitless parameter determining the shape of the production curve. In case n = 2 the stock production reduces to $rB_t(1 - B/K)$ (Schaefer 1954), while the limit of the stock production for $n \to 1$ is $rB_t \log (K/B_t)$ (Fox 1970). The term F_tB_t of Equation (1) represents the instantaneous catch while the remaining part of the right-hand-side of Equation (1) represents the instantaneous biomass surplus production of the stock following a theta-logistic growth function (Pedersen et al. 2011). The intrinsic growth rate r models density-independent growth and natural mortality. The carrying capacity, K, is a density-dependent growth penalty corresponding to the equilibrium B_t of an unexploited stock $(F_t = 0).$

The parametrization of Equation (1), while easy to interpret biologically, is difficult to estimate owing high correlation between r and K. Using geometric arguments related to the dome-shaped production curve Fletcher (1978) derived a more stable parametrization

$$\frac{dB_t}{dt} = \gamma m \frac{B_t}{K} - \gamma m \left(\frac{B_t}{K}\right)^n - F_t B_t \tag{2}$$

where $\gamma = n^{n/(n-1)}/(n-1)$ and

$$m = \frac{rK}{n^{n/(n-1)}}\tag{3}$$

The case n = 1 is a removable singularity corresponding to the model of Fox (1970). Estimating

m instead of r results in a more well-defined optimum because of the separate roles of m and K in defining the production curve with K representing the width of the biomass range and $m = \mathrm{MSY}^d$ representing the maximum sustainable yield (maximum attainable surplus production). The biomass and fishing mortality leading to maximum surplus production are $B_{\mathrm{MSY}}^d = n^{1/(1-n)}K$ and $F_{\mathrm{MSY}}^d = m/B_{\mathrm{MSY}}$. The superscript d indicates that these are deterministic reference points that do not account for random variability. If n=1, the reference points have the limits $B_{\mathrm{MSY}}^d = K/e$, $\mathrm{MSY}^d = rK/e$ and $F_{\mathrm{MSY}}^d = r$.

Equation (2) is a simplified and deterministic description of biomass dynamics. In reality, many additional factors (e.g. species interactions, environmental variability) influence biomass dynamics. In the absence of specific data pertaining to these processes, one can model their influence using a stochastic process noise term. Including process noise in Equation (2) results in

$$dB_t = \left(\gamma m \frac{B_t}{K} - \gamma m \left[\frac{B_t}{K} \right]^n - F_t B_t \right) dt + \sigma_B B_t dW_t$$
 (4)

where σ_B is the standard deviation of the process noise, and W_t is Brownian motion. Representing surplus production by a stochastic differential equation (SDE, Øksendal 2013) acknowledges the presence of random and unmodelled process variability, while retaining the property that the process is defined continuously in time and not restricted to specific discrete time points.

The process noise of Equation (4) is multiplicative owing to the presence of B_t in the noise term $\sigma_B B_t dW_t$. Multiplicative noise terms can, in terms of numerical implementation and model fitting, lead to instability problems. By Lamperti transforming the model (Iacus 2009), an additive noise term is obtained

$$dZ_{t} = \left(\frac{\gamma m}{K} - \frac{\gamma m}{K} \left[\frac{e^{Z_{t}}}{K}\right]^{n-1} - F_{t} - \frac{1}{2}\sigma_{B}^{2}\right) dt + \sigma_{B} dW_{t}$$
(5)

where $Z_t = \log (B_t)$. Interestingly, the standard deviation of the process noise, σ_B^2 , now appears in the deterministic part of Equation (5) and therefore influences the reference points related to MSY. For n > 1, Bordet and Rivest (2014) derived the following stochastic reference points

$$B_{\text{MSY}} = B_{\text{MSY}}^{d} \left(1 - \frac{1 + F_{\text{MSY}}^{d}(n-2)/2}{F_{\text{MSY}}^{d}(2 - F_{\text{MSY}}^{d})^{2}} \sigma_{B}^{2} \right)$$
(6)

$$F_{\text{MSY}} = F_{\text{MSY}}^d - \frac{(n-1)(1 - F_{\text{MSY}}^d)}{(2 - F_{\text{MSY}}^d)^2} \sigma_B^2 \qquad (7)$$

$$MSY = MSY^{d} \left(1 - \frac{n/2}{1 - (1 - F_{MSY}^{d})^{2}} \sigma_{B}^{2} \right)$$
 (8)

Determining whether the stochastic reference points hold for n < 1 is an area of ongoing research and outside the scope of this study. It is evident that $F_{\rm MSY} < F_{\rm MSY}^d$ when $\sigma_B^2 > 0$, which implies that the presence of process noise entails a conservative reduction in recommended fishing pressure relative to the deterministic case; however, the opposite behaviour is observed if Equation (7) is assumed to hold for n < 1. We therefore use the stochastic reference points when n > 1 and the deterministic reference points when $0 < n \le 1$.

Subject to constant F_t , B_t has a stationary distribution (Bordet and Rivest 2014) with an expected equilibrium biomass, which can be approximated by

$$E(B_{\infty}|F_{t}) = K \left(1 - \frac{(n-1)}{n} \frac{F_{t}}{F_{\text{MSY}}^{d}}\right)^{1/(n-1)}$$

$$\left(1 - \frac{n/2}{1 - \left(1 - nF_{\text{MSY}}^{d} + [n-1]F_{t}\right)^{2}} \sigma_{B}^{2}\right)$$
(9)

As expected $E(B_{\infty}|F_t)=K$ in the absence of fishing $(F_t=0)$ and process noise $(\sigma_B^2=0)$. In general, $E(B_{\infty}|F_t)$ can be interpreted as a fished equilibrium, which can serve as a biomass predictor if current conditions remain constant.

The majority of existing production models leave the process of commercial fishing, F_t , unmodelled. Discrete-time models (e.g. Punt 2003) commonly assume that $F_t = C_t/B_t$ where C_t is the observed aggregated catch in year t. This approach implicitly assumes that B_t and F_t represent annual averages of biomass and fishing mortality. Perhaps more importantly, it is assumed that the catch is observed without error. If present, observation error in the catch will therefore propagate directly to F_t and influence conclusions regarding the current fishing pressure. Furthermore, previous

models are only able to estimate F_t at times when a catch observation is available.

An alternative approach, which addresses the above issues, is to model F_t as a separate and unobserved process in the same sense that B_t is unobserved, which allows F_t to be estimated at any time even when a catch observation is unavailable. Our general model for F_t is the product of a random component G_t and a seasonal component S_t

$$F_t = S_t G_t \tag{10}$$

$$d \log G_t = \sigma_F dV_t \tag{11}$$

$$d\log G_t = \sigma_F dV_t \tag{11}$$

where dV_t is standard Brownian motion and σ_E is the standard deviation of the noise. If only annual data are available it is not possible to estimate within-year dynamics and therefore $S_t = 1$ and consequently $F_t = G_t$. Given subannual data, we suggest two models for seasonal variation in the fishing. The first model represents seasonal variation using a cyclic spline, while the second model uses a system of SDEs whose solutions oscillate periodically. In the case of the seasonal spline, F_t follows the model

$$F_t = \exp(D_{s(t)})G_t \tag{12}$$

where $D_{s(t)}$ is a cyclic B-spline with a period of 1 year with $s(t) \in [0; 1]$ being a mapping from t to the proportion of the current year that has passed. The possible annual variation allowed by the cyclic B-spline is determined by a chosen number of socalled knots. The number of knots must be smaller than or equal to the number of catch observations per year (e.g. quarterly catches can at most accommodate four temporally equidistant knots). The values of the cyclic B-spline is defined by the parameter vector φ of length equal to the number of knots minus one. In the case of annual data (one knot) the cyclic B-spline reduces to a constant $(D_{s(t)} = 1)$ and φ has zero length and is therefore not estimated. Note that the seasonal pattern represented by the spline remains constant in time. Thus, a spline-based model is not able to adapt to changes in amplitude and timing (phase) of the real seasonal fishing pattern. Such variations in the fishing pattern would, when fitted with a spline-based model, likely lead to autocorrelated catch residuals.

A way to overcome the potential problems of the spline-based seasonal model is to construct a model that uses the current state and the state a year ago to predict the future state. In discrete

time, such a seasonal correlation structure is readily implemented by including the state at appropriate temporal lags; however, in continuous time, obtaining such a correlation structure is more complicated. An approximation of this structure is to use a system of coupled SDEs, which perturb each other resulting in oscillating solutions. A simple form of such a system is

$$dU_{1,t} = -(\lambda U_{1,t} + \omega U_{2,t})dt + \sigma_U dX_{1,t}$$
 (13)

$$dU_{2,t} = -(\lambda U_{2,t} - \omega U_{1,t})dt + \sigma_U dX_{2,t}$$
 (14)

where $dX_{1,t}$ and $dX_{2,t}$ are standard Brownian motion, σ_U is the standard deviation of the noise, ω is a phase parameter set such that the period of the cycles is 1 year, and $\lambda > 0$ is a damping parameter that ensures that the process is stationary. The solution to the SDE system is known analytically to be oscillatory (Gardiner 1985); thus, $U_{1,t}$ can be used to represent seasonal variation in F_t by the following expression

$$F_t = \exp(U_{1,t})G_t \tag{15}$$

One can therefore think of $U_{2,t}$ as a slave state that is not used directly to calculate F_t as it only enters through its coupling with $U_{1,t}$. The coupled SDE model is more flexible than the spline-based model in that it is able to accommodate gradual shifts in amplitude and timing of the seasonal fishing pattern. This advantage comes at the cost of increased model complexity (as $U_{1,t}$ and $U_{2,t}$ are unobserved processes) and therefore potential difficulties in obtaining model convergence.

An observation of commercial catch is reported as the cumulative catch C_t over a time interval Δ_t . Given B_t and F_t the observed catch in log can therefore be written as an integral in continuous time (Prager 1994) plus noise

$$\log(C_t) = \log\left(\int_t^{t+\Delta_t} F_s B_s ds\right) + \epsilon_t \tag{16}$$

where the catch observation errors $\epsilon_t \sim N(0, \sigma_c^2)$ are independent and σ_C is the standard deviation of the catch observation error. The formulation of Equation (16) allows the noise of the F_t process to be separated from the observation noise of C_t . The model furthermore handles catches sampled at any time and aggregated over any interval length Δ_t , for example a quarter of a year. For model fitting, we assume to have $N_{\text{obs}C}$ observations of C_t together with the associated time intervals Δ_t over which C_t were accumulated.

In addition to catch observations, we assume to have observations of N_i series of indices of exploitable biomass $(I_{t,i} \text{ for } i=1,\ldots,N_i)$ with series i containing $N_{\text{obs}I,i}$ observations. A biomass index could be commercial or scientific catch-per-unit-of-effort data or other biomass indicators, for example as derived from acoustic surveys. Contrary to catch observations that are aggregated over a period of time, $I_{t,i}$ are regarded as 'snapshots' related to the time point t given as

$$\log(I_{t,i}) = \log(q_i B_t) + e_{t,i} \tag{17}$$

where $e_{t,i} \sim N(0, \sigma_{l,i}^2)$ are independent normal deviates and $\sigma_{l,i}$ is the standard deviation of the *i*th index observation error, and q_i is a catchability parameter for the *i*th index. In the common situation where only one series of biomass indices is available $(N_i = 1)$, the *i* index will be omitted for notational simplicity.

We also define the ratios between observation and process errors $\alpha = \sigma_I/\sigma_B$ and $\beta = \sigma_C/\sigma_F$. In cases where it is not possible to separate process and observation error, a common simplification is to assume process error of B_t and observation error of I_t to be equal (Ono *et al.* 2012; Thorson *et al.* 2013), that is to fix $\alpha = 1$. A similar relationship between the process error of F_t and the observation error of C_t could be envisioned, that is that $\beta = 1$, which we use when σ_C and σ_F cannot be estimated separately.

Extreme observations or outliers in index and catch are commonly encountered problem in fisheries data (Chen et al. 1994). Such outliers are poorly modelled when using the normal distribution for observation errors, which may lead to bias of parameter estimates. Common approaches to mitigate the influence of outliers include objective outlier detection and subsequent residual rescaling (Prager 2002), or robust estimation using fat-tailed error distributions (Chen et al. 2000). Here, we take a simple approach to robust estimation where the observation error follow the mixture distribution pN $(0, \sigma^2) + (1 - p)N(0, [w\sigma]^2)$, where p and w are parameters controlling the fatness of the tails. The parameters p and w should in principle be estimated from data; however, as we include robust estimation mainly for illustrative purposes, we fix p = 0.95 and w = 15 for simplicity.

In summary, SPiCT is a state-space model for surplus production containing unobserved processes for B_t and F_t and observed quantities C_t and

 $I_{t,i}$, which include observation noise. In addition to the usual parameters of deterministic production models, process- and observation variance parameters, σ_B , σ_F , $\sigma_{I,i}$ and σ_C , are estimated from data (if possible), while the unobserved processes B_t and F_t are treated as random effects. The spline-based representation of seasonal fisheries requires φ as an additional parameter vector, while the coupled SDE system requires the parameters λ and σ_U as well as the unobserved processes $U_{1,t}$ and $U_{2,t}$, which are also treated as random effects.

Including prior information

Auxiliary information can, if available, be incorporated in a Bayesian estimation framework using so-called informative priors, which are probability distributions that narrow the range of the model parameters they target. Including priors typically stabilizes model fitting and reduces uncertainty of estimated quantities. However, the opposite is true if information contained in the data contradicts prior information. In a data-limited situation where some model parameters may be difficult to estimate, vaguely informative priors may be specified to aid model convergence and identifiability. This approach can be regarded as a compromise between fixing parameters and estimating them unconstrained (Magnusson and Hilborn 2007). It is, of course, imperative that priors are only included if their specification relies on a solid foundation such as meta-analyses or independent data. Particular caution is required if informative priors are specified for n, r, m or K, as these are the main parameters determining management quantities.

Estimates and intervals

Both frequentist and Bayesian inference of model parameters are possible. In a frequentist framework, model parameters are estimated by maximizing the log-likelihood function (Pawitan 2013). In case of available *a priori* information, prior distributions are multiplied with the likelihood function to obtain the posterior distribution. Bayesian maximum *a posteriori* parameter estimates are thus located at the maximum of the posterior distribution (Box and Tiao 2011). While confidence intervals of the frequentist framework and credible intervals of the Bayesian framework have philosophical differences, they are in this study both approximated using the curvature of the objective

function at the optimum and are therefore both abbreviated CI.

Model checking

An important step in fitting stochastic models is the post hoc evaluation of the quality of the model fit as expressed by the residuals. Generally, for a state-space model the so-called one-step-ahead (OSA) residual r_t , at time t_i is

$$r_{t_j} = \frac{Y_{t_j} - E(Y_{t_j}|Y^{t_{j-1}})}{\sqrt{\operatorname{var}(Y_{t_j}|Y^{t_{j-1}})}}$$
(18)

where $E(Y_{t_j}|Y^{t_{j-1}})$ is the OSA prediction of theobservation Y_{t_i} given $Y^{t_{j-1}} = \{Y_{t_1}, ..., Y_{t_{j-1}}\}$ and $var(Y_{t_i}|Y^{t_{j-1}})$ is the variance of this prediction 1990). We (Harvey calculate $j \in \{2, ..., N_{\text{obs}}\}, \text{ where } N_{\text{obs}} = N_{\text{obs}C} + \sum_{i} N_{\text{obs}I,i}$ is the total number of available observations. The vector $Y^{t_{N_{obs}}}$, which comprises all catch and index observations, is ordered after the times the data points are observed. Thus, while indices are assumed to be observed as snapshots, catches are observed at the end of the interval they refer to (i.e. C_t is observed at $t + \Delta t$). Residual checks are performed separately for each data series of catches and indices to enable identification of the problematic model component.

If the estimating model is equal to the data-generating model the OSA residuals (onwards simply referred to as residuals) should be independent and standard normally distributed. Model deficiencies can therefore be indicated by checking whether the residuals display these properties. Possible violation of the independence assumption can be checked by plotting the empirical autocorrelation function or by using the test of Ljung and Box (1978). Apparent autocorrelation in the residuals indicates that the model lacks the structure to appropriately describe underlying dependencies in the data. As a consequence, the obtained parameter estimates and associated CIs cannot be trusted. Additionally, residuals should also be tested for normality, for example using the test of Shapiro and Wilk (1965), and bias (mean different from zero) using a standard t-test.

Implementation

The model is implemented using Template Model Builder (TMB, Kristensen *et al.* 2015), which is a recently developed estimation framework for R (R

Core Team, 2015). TMB is efficient in fitting models with many random effects and is therefore well-suited for estimation of state-space models. The temporal dimension of the continuous-time model is resolved numerically using an Euler scheme (Iacus 2009), which discretizes time into intervals of fixed length dt_{Euler} . The number of time intervals per year is therefore $1/dt_{Euler}$. To stabilize parameter estimation and calculation of the initial one-step-ahead residuals, we impose a wide normal distribution on the initial states $\log (B_t/K)$ and $\log (F_t)$ with mean $\log (0.8)$, on $\log (n)$ with mean $\log (2)$, and on $\log (\alpha)$ and $\log (\beta)$ with mean $\log (1)$. All distributions have a standard deviation of 10. These constraints on model parameters can also be interpreted as uninformative priors. Note, however, that these distributions are imposed purely for numerical reasons and should not have other impact than improving numerical stability and avoiding extreme non-sensical estimates.

R and TMB code for fitting the model is bundled in an R package included as part of the supplementary information. The package also includes routines for post-processing, model checking and plotting.

Examples

South Atlantic albacore

To facilitate a comparison with previously published surplus production models and estimation approaches, we fitted SPiCT to the South Atlantic albacore (Thunnus alalunga, scombridae) data set of Polacheck et al. (1993). The data set comprise $N_{\text{obs}C} = N_{\text{obs}I} = 23$ years of catch and index pairs. The data contained too little information to allow unconstrained estimation of parameters α , β and n. First, we therefore fixed $\alpha = 4$ similar to the estimates found by Meyer and Millar (1999), $\beta = 1$ and n = 2 resulting in a quadratic production curve (SPiCT1). Second, as an alternative to fixing parameters (SPiCT2), we applied vague normally distributed priors to α , β and n with mean parameters equal to the fixed values of the SPiCT1 case and standard deviations of 2 in the log domain (Table 1). In both SPiCT1 and SPiCT2, we set $dt_{\text{Euler}} = 1/16$ year. Finally, we fitted ASPIC version 7.02 (Prager 1994), which is a continuous-time observation-error model, to contrast the discrete-time observation-error and process-error models of Polacheck et al. (1993) and the statespace model of Meyer and Millar (1999).

Table 1 Stochastic surplus production model in continuous time (SPiCT) estimates of α , β and n using the South Atlantic albacore data set of Polacheck *et al.* (1993) with 95% CIs in parentheses.

Parameter	SPiCT1, fixed	SPiCT2, prior	SPiCT2, posterior
α	4.0	4.0 (0.073, 218)	12.3 (1.300, 117)
β	1.0	1.0 (0.018, 54.6)	0.12 (0.018, 0.78)
n	2.0	2.0 (0.037, 109)	0.75 (0.073, 7.75)

Confidence intervals were obtained using the bootstrap module of ASPIC with 1000 samples.

Similar to Polacheck *et al.* (1993) results are presented in terms of model parameters K, r and q, B_{1990} (the predicted biomass in 1990), B_{1990}/K (the biomass depletion relative to K in 1990), the estimated MSY, and the estimated optimal effort $E_{\rm MSY} = F_{\rm MSY}/q$. For SPiCT, the stochastic reference points are reported, the residuals were checked for autocorrelation and tested for bias and normality, and a short-term forecast under a constant F_t scenario was included to illustrate the ability of the model to forecast catch including uncertainty. To facilitate comparison of the estimation methods, we calculate estimated quantities relative to the estimates of the observation-error model of Polacheck *et al.* (1993).

Simulation study 1

The purpose of simulation study 1 study was to quantify the estimation performance of SPiCT in terms of estimation stability (proportion of converged runs), estimation precision (expressed by the coefficient of variation, CV, of estimates), the coverage of 95% CIs (proportion containing the true value) and the median bias of estimates. These quantities were evaluated for eight variants of SPiCT (Table S2), ASPIC version 7.02 (Prager 1994), with particular focus on the influence of fixing and misspecifying the parameters n, α , and β . which can be difficult to estimate. Subsequently, we also fitted the model of Meyer and Millar (1999) and compared with SPiCT using identical informative priors for the two models (see Data S1 for detailed description).

Simulation study 2

The purpose of simulation study 2 was to assess the difference in estimation performance between a continuous-time model fitted to quarterly data containing within-year seasonal variation and a discrete-time model ($dt_{\rm Euler}=1$ year) fitted to annual data obtained by aggregating the quarterly data. In practice, catch and biomass index data are often resolved by annual time steps; however for some stocks, data with a higher temporal resolution are available. While quarterly data contain four times the number of observations of the corresponding aggregated annual data, they do not necessarily contain four times the information. This is because increasing the sampling frequency of a process typically results in increased autocorrelation of data and is unlikely to substantially increase contrast compared to annual data.

For simulation, we used parameter values found by fitting SPiCT to the South Atlantic albacore data set of Polacheck *et al.* (1993) while fixing $\alpha=1$, $\beta=1$ and n=2. Seasonal variation in F_t was included using a spline-based model with $\varphi=(0.05,\,0.1,\,1.8)$ resulting in low values of F_t in quarters two and three, and high values of F_t in quarter four relative to quarter one. We simulated 30 years of biomass and fishing dynamics using a fine time step $(dt_{\rm Euler}=1/64~{\rm year})$ and collected quarterly $(N_{{\rm obs}C,Q}=N_{{\rm obs}I,Q}=120)$ and corresponding annual $(N_{{\rm obs}C,A}=N_{{\rm obs}I,A}=30)$ data sets of catches and biomass index.

We fitted SPiCT using $dt_{Euler} = 1/32$ year to the quarterly data with n = 2 and $\beta = 1$ fixed to their true values while σ_B and σ_I were estimated separately. The continuous-time model estimated φ to fit the seasonal variation in F_t using the spline-based approach. We also fitted the model using $dt_{\text{Euler}} = 1$ year (i.e. a discrete-time model) to the annual data with the same parametrization (excluding φ). The procedure was repeated 1000 times. To assess the difference in general utility of the models, the proportion of converged estimations for each model was calculated. Estimation performance was then summarized for the data sets for which both models converged in terms of median CV and coverage of 95% CIs of F_{MSY} , B_{MSY} , MSY, σ_B , and σ_I . As the CV scales with the inverse square root of the number of independent observations, we also calculated $I = (CV_A/CV_O)^2$ for each of the five quantities, to express the increase in effective sample size. The squared ratio of the CVs (J) is proportional to the ratio of the effective sample sizes (i.e. the number of independent observations) of quarterly data to annual data and therefore an indicator of information gain.

North Sea stocks

To illustrate the utility of SPiCT, we analysed subannual data from five North Sea fish stocks in the period 1975–2006: Cod (Gadus morhua, gadidae) with $N_{\rm obsC}=124$ and $N_{\rm obsI}=61$, whiting (Merlangius merlangus, gadidae) with $N_{\rm obsC}=124$ and $N_{\rm obsI}=60$, haddock (Melanogrammus aeglefinus, gadidae) with $N_{\rm obsC}=124$ and $N_{\rm obsI}=60$, herring (Clupea harengus, clupeidae) with $N_{\rm obsC}=124$ and $N_{\rm obsI}=60$ and Norway pout (Trisopterus esmarkii, gadidae) with $N_{\rm obsI}=119$ and $N_{\rm obsI}=57$.

Quarterly information on total catch was obtained from ICES (2005). As quarterly catches have not been compiled in the more recent years, and some of the included stocks have minor data issues requiring in-depth scrutiny that are beyond the scope of this study, our results for these stocks should be viewed solely as illustrative.

The fish stocks of the North Sea are subject to substantial fishing pressure and are therefore surveyed extensively by scientific vessels. We gathered survey data for the time period 1975-2006 from the ICES DATRAS database (ICES, 2012) and calculated indices of exploitable stock biomass by weighting age-structured survey catches with the ratio of age-specific commercial selectivity to survey selectivity. Specifically, the index $I_{\text{ESB}} = \sum_{a} Pc_a / Ps_a Ns_a Ws_a$, where Pc_a and Ps_a are commercial and survey selectivities, respectively, and Nsa and Wsa are numbers and mean weight at age a in the surveys, respectively (time index omitted for simplicity). Thus, calculating I_{ESB} requires knowledge or assumptions about selectivities. Here, we approximate the ratio Pc_a/Ps_a by the ratio Qc_a/Qs_a , where Qc_a is the proportion of the commercial catches at age a and Qs_a is the proportion of the survey catches at age a. For small sample sizes, this is a crude approximation in which case simple assumptions regarding selectivities may be more robust. If available, sampling CVs of the index can be used to weight index observations: however, this information was not part of the North Sea data set.

Owing to variable survey, frequency index time series contains annual, biannual and quarterly time steps between observations. With the continuous-time formulation of SPiCT it is possible to analyse the variably sampled data directly without the need to aggregate observations into fixed time steps.

We assessed to which extent models of different complexity were identifiable given the data by first estimating a baseline model with a spline-based seasonal representation while fixing $\alpha=1$ and n=2, which are standard assumptions (Ono *et al.* 2012; Thorson *et al.* 2013) and fixing $\beta=1$. All remaining model parameters $(K, m, q, \sigma_B=\sigma_I, \sigma_F=\sigma_C)$ were estimated. We then released the fixed parameters for estimation in the following order: α , β and n resulting in three models to be estimated in addition to the baseline. We set $dt_{\rm Euler}=1/32$ year for all model fits.

Residuals were analysed to diagnose model insufficiencies. If significant residual autocorrelation was detected in catches as an indication of potential shifts in the seasonal fishing pattern, we ran the more flexible coupled SDE seasonal model and re-examined residuals. If residuals deviated from normality possibly as a result of outlying observations we reanalysed data using robust observation-error distributions for either or both catch and index observations. Models were selected as the ones with best residual diagnostics (fewest violations in terms of autocorrelation, bias and non-normality). If models performed equally in terms of diagnostics, we selected models with highest complexity (most free model parameters) to give examples of estimated values of α , β and n. Aspects of formal statistical model selection are outside the scope of this study.

When fitting production models a constraint is often required to estimate absolute levels of biomass and fishing mortality (B_t and F_t , respectively). This was the case for cod and whiting for which a bound was imposed on B_t in the year (y), the maximum catch was observed by applying a vague prior distribution based on information in the catch data. For stocks that have been subjected to sustained and relatively high commercial exploitation for decades, it seems fair to assume that current biomass is not many orders of magnitude larger than the observed catches. As prior distribution for $\log (B_y)$, we therefore used $N(\mu_{B_u}, \sigma_{B_u}^2)$, with $\mu_{B_u} = \log(k \max C_t)$ and $\sigma_{B_u} = 3$. Note that a standard deviation of 3 in the log domain is a rather uninformative prior. The specific choice of k has minimal impact on the results as long as μ_{B_u} is in the same order of magnitude as the maximum catch $(1 \le k \le 9)$ and the prior is uninformative $(\sigma_{B_u} \ge 3)$. In this study, we set k = 3.

Results

South Atlantic albacore

Fitting SPiCT1 to the South Atlantic albacore data set of Polacheck *et al.* (1993) produced results comparable to those obtained using alternative approaches (Fig. 1). The point estimates of SPiCT1 were particularly similar to those obtained using observation-error estimators; however, the point estimates of alternative approaches were contained in the 95% CI of the SPiCT1 point estimates. SPiCT2 produced a wider CI of $E_{\rm MSY}$ while CIs of K and B_{1990} were narrower compared to CIs of SPiCT1. CIs of remaining parameters were largely similar between SPiCT1 and SPiCT2. Overall, among all methods estimates of MSY were nearly identical and associated with the narrowest CIs.

SPiCT1 produced 95% CIs of B_t and F_t that were wide relative to the 95% CIs of $B_t/B_{\rm MSY}$ and $F_t/F_{\rm MSY}$ (Fig. 2). The reduction in CIs of relative estimates was a result of high correlation between estimates of absolute levels and reference points. For SPiCT2, this correlation was less marked and reference point estimates were uncertain resulting in inflated CIs of relative levels of biomass and fishing mortality making stock status less clear.

In terms of parameter estimates, SPiCT2 obtained a significantly lower value of β than 1 (Table 1) resulting in reduced estimated catch observation error (Fig. 2). Overall, estimates of α , β and n were uncertain with standard deviations in log domain of 1 (Table 1). Model checking of both SPiCT1 and SPiCT2 did not indicate any violation of assumptions regarding independence, bias

and normality of residuals (Figures S1 and S2) for catch and index observations. Formal model selection would be a natural step to objectively choose between multiple models; however, this topic is beyond the scope of this study and is left to be explored in future work.

Predictions of B_t , $B_t/B_{\rm MSY}$, F_t , $F_t/F_{\rm MSY}$ and C_t can be obtained by propagating the model beyond the time span of the data (Fig. 2); however, uncertainty increases for each predicted year as a result of the lack of data. Furthermore, as the model is a highly simplified version of reality, it should not be relied on for generating long-term predictions (>2 years) even when residuals pass all tests.

Simulation study 1

Simultaneous estimation of α , β and n (model A) was less stable than estimation of models with fixed n = 2; however, while parameter estimates provided by model A were uncertain, coverage of the resulting CIs was generally close to the expected 0.95. Simultaneous estimation of α , β , and n using vague priors (model H) improved both model convergence rate, median CV and median bias, while CI coverage was unchanged relative to model A. Models with fixed n = 2 had lower CVs but also lower CI coverage for shorter data sets. CI coverage was generally reduced when misspecifying n, while misspecification of α and β influenced CI coverage less, in particular estimates of MSY and biomass relative to B_{MSY} were unaffected. Misspecified models were generally biased when estimating absolute quantities (F_{MSY}, B_{MSY}, B_t) and unbiased when estimating relative quantities

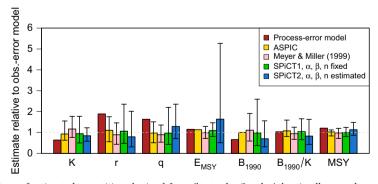


Figure 1 Comparison of estimated quantities obtained from fits to the South Atlantic albacore data set relative to estimates obtained using an observation-error model (Polacheck *et al.* 1993). Error bars are 95% CIs. Note that ASPIC does not report uncertainty on all quantities. Overall, estimates were similar with MSY and the relative estimate B_{1990}/K having lowest uncertainty. Note that estimating α , β and n (SPiCT2) lead to decreased uncertainty of K and K and K and K and K and K and K are while the uncertainty of K are substantially. [Color figure can be viewed at wileyonlinelibrary.com]

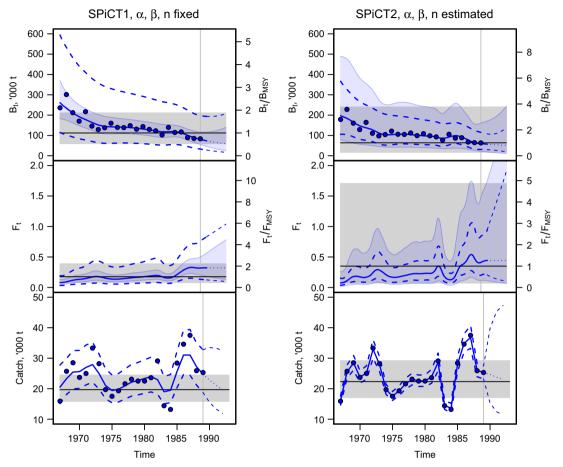


Figure 2 Fits to the South Atlantic albacore data set of Polacheck *et al.* (1993) with α , β and n fixed (SPiCT1) and α , β and n estimated (SPiCT2). Solid blue lines are estimated values, vertical grey lines indicate the time of the last observation beyond which dotted lines indicate forecasts, dashed lines are 95% CI bounds for absolute estimated values, shaded blue regions are 95% CIs for relative estimates (B_t/B_{MSY} or F_t/F_{MSY}), grey regions are 95% CIs for estimated absolute reference points (horizontal lines), solid circles are observations with index plotted as I_t/q . Notably, SPiCT1 estimated narrower CIs of reference points and relative levels while SPiCT2 estimated narrower CIs of absolute levels of B_t , F_t and catch. [Color figure can be viewed at wileyonlinelibrary.com]

 $(F_t/F_{\rm MSY}, B_t/B_{\rm MSY})$ and MSY. Despite its simpler model structure, ASPIC showed comparable performance to SPiCT for these three quantities.

Using informative priors SPiCT had a convergence rate close to one, while the model of Meyer and Millar (1999) had a convergence rate of 0.6 on average. Median CVs of both models decreased as the number of observations increased, however while CIs of SPiCT generally converged to the expected 95% the CIs of the Meyer and Millar (1999) model diverged for increasing number of observations. SPiCT estimates of absolute quantities were slightly biased (~5%) while relative quantities and MSY were unbiased. The Meyer and Millar (1999) model generally produced biased

estimates of absolute quantities while MSY and $B_{\rm last}/B_{\rm MSY}$ were largely unbiased.

See Data S1 for detailed description of results.

Simulation study 2

The convergence rate was 99.9% for both the discrete-time model fitted to the annual data and the continuous-time model fitted to quarterly data. The results of the runs for which both models converged (Table 2) did not show a significant difference in median CVs of estimated MSY, while the continuous-time model obtained significantly improved CVs for the remaining quantities. The improvement in CV of $B_{\rm MSY}$ corresponded to a

Table 2 Results of simulation 2 summarized by median CVs with *P*-values of Mann–Whitney (MW) rank-sum tests of identical CVs, implied information gain (*J*) as given by the squared CV ratio, and coverage of 95% CIs of estimates with *P*-values of equal-proportions test. Results are based on the continuous-time model using 30 years of quarterly observations ($N_{\text{obs}C} = N_{\text{obs}I} = 120$) and the discrete-time model using the corresponding aggregated annual observations ($N_{\text{obs}C} = N_{\text{obs}I} = 30$).

		F_{MSY}	B_{MSY}	MSY	$\sigma_{\mathcal{B}}$	σ_{l}
Median CV	Annual	0.516	0.852	0.469	0.605	0.216
	Quarterly	0.500	0.605	0.480	0.289	0.095
	MW, P-value	0.010	<10 ⁻³	0.817	<10 ⁻³	<10 ⁻³
	J	1.066	1.980	1.000	4.391	5.182
CI coverage	Annual	0.866	0.882	0.735	0.966	0.930
	Quarterly	0.866	0.875	0.870	0.949	0.960
	Prop., <i>P</i> -value	1.000	0.681	<10 ⁻³	0.076	0.004

Table 3 Results of the selected fits of stochastic surplus production model in continuous time (SPiCT) to the five North Sea stocks with 95% CIs given in parentheses. Residual analyses showed no significant violation of assumptions for Norway pout and whiting, while autocorrelation (AC) was detected for herring, and non-normality (NN) was detected for herring, haddock and cod.

Stock	Seasonal model	Robust error	α	β	n	Residual analysis
Herring	Spline	I_t	_	=	_	AC, NN
Norway Pout	Coupled SDE	_	0.59 (0.22, 1.58)	0.84 (0.44, 1.59)	1.27 (0.52, 3.07)	_
Haddock	Coupled SDE	_	1.53 (0.85, 2.77)	0.66 (0.42, 1.05)	_	NN
Cod	Spline	_	2.36 (1.31, 4.24)	0.27 (0.14, 0.52)	_	NN
Whiting	Spline	I_t	2.45 (1.03, 5.87)	1.32 (0.75, 2.3)	_	

doubling of the effective sample size, while CVs of process and observation-error standard deviations were reduced even further as a result of using quarterly data (Table 2). In terms of CI coverage, significant differences between the discrete-time and continuous-time model were found for MSY and σ_I with the most substantial improvement in coverage found for MSY.

North Sea stocks

Of the five selected model fits, α and β were estimated for all stocks except herring, while n was estimated for Norway pout only (Table 3).

The selected fit to Norway pout data detected a clear seasonal pattern in F_t (Fig. 3). Residual analysis of this fit showed no violations in terms of autocorrelation, bias or non-normality of residuals when using the coupled SDE seasonal model (Table 3). Fits to Norway pout data using the spline-based seasonal representation resulted in autocorrelated catch residuals; however, the more flexible correlation pattern allowed by the

coupled SDE seasonal model was able to eliminate this autocorrelation (Table S3). This illustrates how an observed pattern in residuals can aid in identifying a lacking model component. Similarly, the analysis of whiting resulted in no violations when using the spline-based seasonal model with robust observation error for index. This model was selected as a result of non-normality of index residuals (apparent outliers in index observations) detected in the baseline model (Table S3).

The selected model for haddock used the coupled SDE seasonal representation (Table 3) as a remedy to detected autocorrelation in catch residuals when using a spline-based seasonal model (Fig. 4). Non-normality detected in index residuals could not be remedied by neither a robust observation-error distribution nor a spline-based seasonal model (Table S3). In analysing the cod data, significant non-normality was detected in catch residuals. Normality could be achieved using a robust catch observation-error model, however, at a cost of introducing autocorrelation in catch residuals.

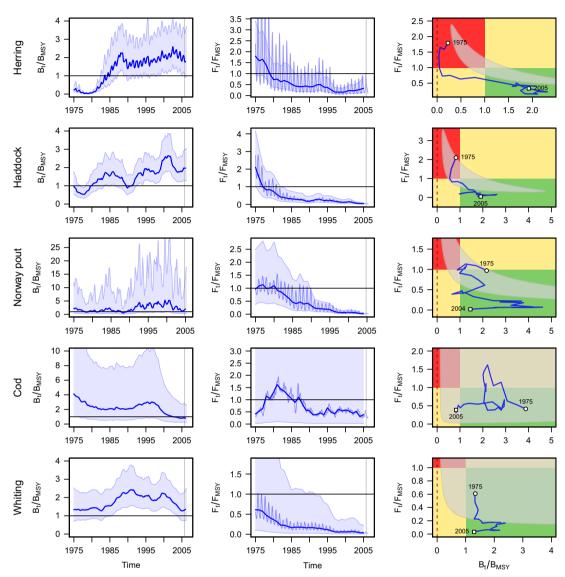


Figure 3 Estimated biomass and annually averaged fishing mortality (solid blue lines) of North Sea stocks relative to estimated reference points with 95% CI (blue shaded region). Fishing mortality including within-year variation (shaded blue lines in middle column) shows the estimated seasonal pattern, which is particularly marked for herring. Uncertainty of estimated reference points is represented by 95% confidence regions on the relative scale (grey-shaded region in right column). [Color figure can be viewed at wileyonlinelibrary.com]

which could not be eliminated using the coupled SDE seasonal model. Finally, for herring, a clear seasonal pattern in F_t was estimated (Fig. 3) resulting in seasonal fluctuations in B_t . Data covered contrasting periods of low and high levels of biomass resulting in reference point estimates with higher precision relative to those of the other stocks (Fig. 3, right column). While a robust observation-error distribution for the index eliminated non-normality of index residuals, non-normality of catch residuals and autocorrelation

could not be eliminated within the current model suite indicating that unmodelled variability in the herring data remains.

In identifying the process to observation noise relationships, two of four estimates of α were significantly different from the value one (Table 3), which is the commonly assumed value when estimation is not possible (Ono *et al.* 2012; Thorson *et al.* 2013). In contrast, only one of four estimates of β were significantly different from the value one. Similarly, the 95% CI of n for Norway

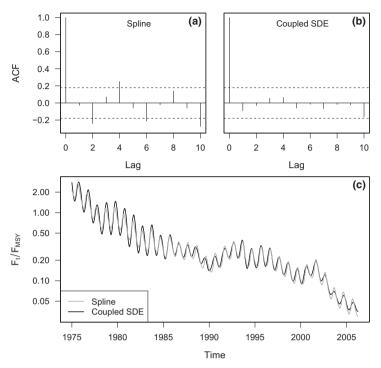


Figure 4 Significant lags are detected in the autocorrelation function (ACF) of North Sea haddock catch residuals using a spline-based seasonal model (a). This violation is remedied when using the coupled SDE seasonal model (b). The detected autocorrelation is caused by the constant amplitude of the spline-based model, which is insufficient compared to the adaptive representation of the coupled SDE seasonal model (c).

pout did not exclude the commonly assumed value of two indicating a lack of evidence to deviate from a symmetric production function (Schaefer 1954).

Discussion

The SPiCT presented here is a full state-space model in that both biomass and fishing dynamics are modelled as states, which are observed indirectly through biomass indices and commercial catches sampled with error. A wide range of previously published surplus production models are nested within SPiCT: Observation-error and process-error estimators (Polacheck et al. 1993) emerge if eliminating process or observation noise, respectively. State-space models that assume catches are observed without error (Meyer and Millar 1999; Punt 2003; Ono et al. 2012) are obtained by fixing the variance of the F_t -process to a large value and eliminating the observation noise on catches. Generally, discrete-time models (Meyer and Millar 1999; Punt 2003) are obtained by setting the temporal time step of the numerical

solver ($dt_{\rm Euler}$) to 1 year, while continuous-time models (Schaefer 1954; Pella and Tomlinson 1969; Prager 1994) arise when reducing $dt_{\rm Euler}$ to a value where parameter estimates do not change qualitatively if $dt_{\rm Euler}$ is refined further.

An advantage of the continuous-time formulation is the ability of the model to accommodate arbitrary and irregularly sampled data without a need for catch and index observations to match temporally. It is therefore straightforward to fit the model to data containing a mix of annual, biannual and quarterly data as demonstrated in our analysis of North Sea stocks. The additional information contained in quarterly observations relative to the corresponding annually aggregated data can be exploited to improve coverage of confidence intervals and reduce uncertainty of parameter estimates (Table 2). Furthermore, the substantial sample size provided by quarterly observations from North Sea stocks enabled estimation of process noise, observation noise and for one stock the production shape parameter (n), which are notoriously difficult to estimate using annual observations (Prager 2002; Ono et al. 2012).

Explicit modelling of the fishery dynamics (F_t) as a latent process in continuous time allows F_t to be estimated at arbitrary times without requiring temporal overlap with a catch observation. The F_t -process further allows catches to be predicted with uncertainty over any time interval by temporal integration of the product of the fishing and biomass processes. Catch predictions can then be compared with observed catches in an observation equation, which incorporates catch observation error [Equation (16)]. In this model formulation, the absence of the catch in the equation describing biomass dynamics avoids the risk of large catches leading to negative biomass estimates, which can be a problem when fitting discrete-time models. The F_t and C_t components of the model are novel relative to previous production models and are first steps towards a framework that more realistically represent the uncertainty of fishing and catch observation processes.

The current model formulation, where catches are given by the temporal integral of the product of fishing and biomass, implies that B_t represents the biomass of the exploitable part of the stock and that $B_{\rm MSY}$ is the level of exploitable stock biomass that maximizes production. It is therefore important when calculating the index of B_t from scientific survey data to only include the sizeclasses targeted by the fishery. It is furthermore important to distinguish ESB from other biomass representations when comparing B_t or B_{MSY} to absolute estimates from alternative models. If the commercial selectivity curve matches the maturity curve of the stock then ESB can be interpreted as the spawning stock biomass (SSB), which is commonly reported from, for example age-based models. Generally, the relative biomass, $B_t/B_{\rm MSY}$, is less sensitive to the choice of biomass representation making it a robust estimator of stock status.

Currently, the main mechanism driving the fishing dynamics is seasonal variations in effort, which is modelled by imposing an annual correlation structure on the fishing mortality. The analyses of North Sea stocks showed that estimation of seasonal variation in fishing is possible if subannual catch observations are available (Fig. 3). A potential extension of the fishing dynamics model could include economic components and allow estimation of bioeconomic quantities (Thorson *et al.* 2013), however, at a cost of increased data requirements.

In fisheries management, it is useful to evaluate the implications of management decisions on stock status and future catches. Such evaluations can be made by predicting catches using Equation (16) under different fishing scenarios or by fixing future catches and predicting the corresponding levels of fishing mortality and biomass including associated prediction intervals. However, as the model is a highly simplified version of reality, it is not suited for generating long-term predictions (>2 years).

While catch observations are aggregated over a time period, biomass index observations are assumed to be instantaneous snapshots. For survey-based indices, where data are often gathered within weeks, this assumption seems reasonable. However, if the data collection period spans months or perhaps the whole year as for commercial CPUE data the assumption can be questioned. In such cases, an alternative to using biomass index data is to use commercial effort data directly in the model as an indicator of F_t . Such an extension could be readily implemented within the presented modelling framework.

In statistical modelling, it is customary to conduct model checking by inspecting residuals for violation of independence and distributional assumptions. Deviating residual patterns can be used to diagnose model insufficiencies. Non-normal or biased residuals indicate the presence of an unmodelled trend in data, for example a temporal shift in catchability, or the presence of extreme outlying observations. If outliers are detected one may apply methods for outlier removal (Prager 2002) or alternatively, as illustrated in the analysis of North Sea stocks (Table 3), shift to a robust model formulation, for example by assuming a heavy-tailed distribution for the data series where outliers occur. Seasonal autocorrelation in residuals may be accounted for by including a seasonal component in the fisheries model. Here, we represented seasonal dynamics using either a cyclic spline [Equation (12)] or the more flexible oscillatory system of coupled SDEs [Equation (15)], which can adapt to shifts in timing and amplitude of the seasonal fishing pattern and reduce residual autocorrelation (Fig. 4).

Persisting residual patterns may be caused by violations of assumptions related to observations, that is differences in biomass index and fisheries data in terms of spatial coverage or uncorrected mismatch between commercial selectivity and survey selectivity. Residual patterns can also be a result of significant changes in biomass and catchability due to migration, shifts in technology or

fishing technique, trends in growth rate imposed by environmental changes, etc. Extending the biomass dynamic model, for example to a stage-based form could capture, more realistically, the selectivity of the fishery and the lagged dynamics induced by interactions between juveniles and adults through recruitment and maturation. Inclusion of environmental covariates such as temperature, nutrient or oxygen information may be used to induce longer-term trends in growth, which, if unmodelled, would result in biased estimates. Complete treatment of the course of action when model assumptions appear violated is outside the scope of this study. However, it is important to include results of residual analyses to enable an honest presentation of model results. If possible, it is furthermore preferable to compare results with those of alternative model classes (catch only, age structured, length based, etc.) in particular if residuals indicate a critical lack of fit.

Formal checking of statistical models is key in obtaining valid parameter estimates, reliable confidence intervals and useful quantities for model selection. In general, model checking has philosophical as well as practical aspects. Idealists would argue that a model producing residuals that violate assumptions lacks the model components required to completely describe data. However, as all models are simplifications of reality and therefore wrong, they can never be expected to fully explain observations. Indeed, surplus production models represent a highly simplified reality where many poorly understood mechanisms such as species interactions and environmental effects are modelled as random variability. Thus adopting a pragmatic approach to model checking allowing mild departures from assumptions may be required.

The flexibility of the presented model is obtained by explicit modelling of both the biomass process (B_t) and the fisheries process (F_t) . Estimation of the potentially large number of resulting random effects is facilitated by TMB via the Laplace approximation (Kristensen *et al.* 2015). In calculating confidence or credible intervals, TMB therefore relies on the assumption that the objective function is quadratic around the optimum, which in the frequentist framework is the asymptotic shape under certain regularity conditions (Wasserman 2013). Thus, in a data-limited scenario asymptotic assumptions may not hold resulting in decreased coverage of confidence intervals. In our simulation, the Laplace approximation provided

95% confidence intervals ranging from 0.8 to 0.9 coverage when 15 observations of catch and index were available to the expected 0.95 coverage at 120–240 observations (Figure S5). When analysing data containing a limited number of observations, it is thus advisable to verify confidence interval coverage using bootstrap or likelihood profiles (de Valpine and Hilborn 2005).

Comparing a deterministic model (ASPIC, Prager 1994) and a discrete-time model (Mever and Millar 1999) with SPiCT requires simplifying assumptions to become tractable. The presented comparison assumes that a realistic representation of population dynamics evolves continuously in time and involves some degree of random variability. Fitting SPiCT to data generated by such a system generally produced unbiased parameter estimates (Figure S6) with reliable confidence intervals (Figure S5), while intervals produced by alternative methods were unreliable in particular for larger data sets (60-240 observations). In a data-limited situation (15-30 observations), all models produced biased estimates of F_{MSY} and B_{MSY} . In contrast, estimates of stock status (F/F_{MSY}) and B/B_{MSY} and MSY performed well across models and are thus the quantities for which production models in general provide the most robust inference.

There are several well-known complications associated with fitting surplus production models: First, as the input data are often collected annually, it is common to have less than say 50 years of data available for estimation. Data scarcity can lead to problems with estimation instability and model identifiability. Second, reliable parameter estimation require sufficient contrast in data (Hilborn and Walters 1992). Lack of contrast can result in identifiability problems and high correlation among model parameters thus making the estimation unstable. Third, estimating both process and observation noise is notoriously difficult (Polacheck et al. 1993; de Valpine and Hilborn 2005) and may necessitate switching to either a processerror or observation-error model, or assume that the ratios of process error to observation error (α and β) are known (Ono et al. 2012; Thorson et al. 2013). Fourth, the three mentioned points apply to the case where n = 2 (Schaefer 1954). Thus, the increased nonlinearity of the generalized form (Pella and Tomlinson 1969) likely exacerbates the mentioned estimation complications.

We do not claim to have solved the above-mentioned problems. However, to mitigate potential

stability issues, we have used the improved parameterization of Fletcher (1978) and have used TMB for model estimation, which relies on analytical derivatives of the objective function to make estimation as stable as possible (Kristensen *et al.* 2015). Extensive simulation testing was facilitated by the efficiency of TMB enabling computing times of parameter estimates in the order of seconds. These aspects are key to obtaining stable fits of nonlinear models (Bolker *et al.* 2013).

Estimating the shape of the production curve (determined by n) is critical to management because the resulting reference points and associated stock status are sensitive to the value of n[Equations (6)–(8)]. Our simulation experiment showed stable performance and high coverage of confidence intervals when simultaneously estimating process noise, observation noise and n even for limited data (Figure S5). Additionally, the model was able to estimate n in one of five North Sea stocks (Table 3). These results are based on unconstrained estimation of n. Our simulation results demonstrated that estimation can be stabilized by translating meta-analyses (e.g. Thorson et al. 2012) or knowledge of stock dynamics into a prior distribution for n (Figure S3). Using a vague prior enabled n to be estimated for the South Atlantic albacore data set of Polacheck et al. (1993) resulting in differences with potential implications for management relative to results using fixed n = 2 (Fig. 2). The improved stability of the presented modelling framework enables stock assessment scientists to explore the possibility of estimating n rather than relying on the common assumption that n = 2, which may result in biased reference point estimates and poor management decisions (Maunder 2003).

Given an MSY-based approach to management, the modelling framework developed here focuses on honest reporting of model results through uncertainty quantification and model checking. By relaxing the common assumption that catches are known without error, a more realistic quantification of uncertainty on all reported quantities is obtained while residual diagnostics clarifies whether model assumptions are significantly violated. As a benefit of the continuous-time formulation quarterly resolved data may, if available, aid in reducing estimation uncertainty (Table 2) and allow seasonal predictions and management. Using the presented model, we have also demonstrated that while estimates of B_t , F_t , $B_{\rm MSY}$ and $F_{\rm MSY}$ are

often highly uncertain, the relative quantities $F_t/F_{\rm MSY}$ and in particular $B_t/B_{\rm MSY}$ may have considerably less uncertainty (Figure S4) and bias (Figure S6) and should therefore form the primary basis of management decisions.

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Supporting Information

- Additional Supporting Information may be found in the online version of this article:
 - Data S1. Supplementary methods and results.
 - Data S2. R code bundled as an R-package.

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