# Supplementary Materials for 

# Impacts of historical warming on marine fisheries production 

Christopher M. Free*, James T. Thorson, Malin L. Pinsky, Kiva L. Oken, John Wiedenmann, Olaf P. Jensen
*Corresponding author. Email: cfree14@gmail.com
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## Materials and Methods

## Study design

We used a Pella-Tomlinson surplus production model (17) with a multiplicative temperature influence term to measure the influence of ocean warming on the productivity of 235 marine fish and invertebrate stocks in the RAM Legacy Stock Assessment Database (19). We estimated the sea surface temperatures (SST) experienced by each stock by mapping the boundary of the stock (i.e., the spatial domain of the stock assessment) and calculating the mean annual SST within this boundary using the COBE SST dataset (18). To determine whether taxonomy (order or family), geography (FAO major fishing area or ecoregion), or stock assessment method structure the SST influences, we evaluated models with hierarchical SST influence based on each of these five groups. We used Akaike Information Criterion (AIC; (21)) to compare models and selected the model with the lowest AIC score as the "final" model. Next, we evaluated whether the final model SST influences were additionally determined by: (1) life history traits such as growth rate, maximum age, or depth preference; (2) behavioral traits such as reproductive guild, migratory habits, or spawning behavior; (3) stock characteristics such as trend in biomass or fishing pressure; and (4) thermal experience such as mean SST, SST trend, or latitude. Lastly, we used the final model to hindcast SST-dependent maximum sustainable yield from 1930-2010 over all stocks and among ecoregions. To ensure that the estimated distribution of SST influences was not due to chance alone, we compared the final model results to results from null models using simulated SST time series designed to decouple the observed SST and productivity time series. We also explored the sensitivity of our results to using the COBE SST dataset rather than the ERSST (53) or HadISST (54) datasets and to modeling SST influence as a random rather than fixed effect.

## 1. Data collection

### 1.1 Stock selection

We analyzed the non-salmon stocks in the RAM Legacy Stock Assessment Database (RAMLDB v3.8; (19)) with time series of total biomass (metric tons) and catch or landings (metric tons; catch preferred) longer than 20 years after trimming years poorly informed by catch and survey data (Table S5). We identified stocks and years to trim by visually inspecting the (1) surplus production and stock-recruit relationships and (2) biomass, recruitment, and catch time series for all candidate RAMLDB stocks (Appendix A). Stocks exhibiting smooth surplus production or stock-recruit relationships over the entire time series were excluded from the analysis. Years, largely at the beginning of the time series, exhibiting flat or smooth biomass or recruitment or perfectly linear catch were excluded from the analysis. Most stocks assessed using biomass dynamics models (largely tuna and marlin stocks) were excluded from the analysis because their dynamics were strongly driven by an assumed production function. However, we included 30 stocks assessed using biomass dynamics models that were visually judged to exhibit enough process variability for consideration in the analysis. Finally, we
excluded 28 stocks that prevented model convergence because they either (1) lacked periods of low exploitation and high biomass necessary to constrain carrying capacity or (2) exhibited population dynamics wildly divergent from stationary logistic population growth (Appendix B). The resulting 235 stocks represent a variety of taxa ( 213 bony fish, 15 crabs/shrimps/lobsters, 4 bivalves, 2 squids, 1 ray), life histories, and locations and approximately $33 \%$ of reported global catch (28 of 86 million metric tons in 2000; (1)).

### 1.2 Stock boundary delineation and SST time series

We estimated the sea surface temperatures (SST) experienced by each stock by mapping the boundary of the stock (i.e., the spatial domain of the stock assessment) and calculating the mean annual SST within this boundary using the COBE SST dataset (COBE v2; (18)). The COBE dataset provides monthly SST on a globally complete $1^{\circ} \times 1^{\circ}$ grid from 1850present based on an interpolation of in-situ and satellite-derived SST observations. We conducted sensitivity analyses using the ERSST (53) and HadISST (54) datasets to ensure that the results were not sensitive to the choice of SST dataset (Fig. S1; Appendix C). Stock boundaries were delineated by either (1) merging the statistical/management areas used to define the assessment area; (2) digitizing the assessment area directly from the stock assessment; or (3) clipping the managing country's exclusive economic zone or the managing agency's area of competence to the geographical reference points provided in the stock assessment. In the USA and Australia, we used information on the geographic distribution of each species (i.e., essential fish habitat and modelled distribution, respectively) to further constrain stock boundaries.

## 2. Modeling

### 2.1 Overview

We modeled temperature-dependent fisheries productivity in five stages. First, we modeled productivity without a temperature effect and used this "standard" model as a benchmark for parameterizing and evaluating models with a temperature effect. Second, we extended the standard model to include a multiplicative temperature influence term and used this model to evaluate whether temperature influences fisheries productivity. Third, we added hierarchical structure to the temperature term to test whether taxonomy, geography, or stock assessment method determines temperature influences. Fourth, we developed simulated temperature time series to confirm that our results were not an artifact of model structure. Finally, we evaluated how our results and conclusions would change if the temperature influences were modeled as fixed rather than random effects. We also detail how our attempt to model dome-shaped temperature dependence by estimating an additional thermal optima parameter proved impossible because of insufficient contrast in the temperature data.

### 2.2 Standard surplus production model

We modeled fisheries productivity using a Pella-Tomlinson surplus production model (17) with first-order autocorrelation in the residuals. Observed surplus production was calculated for each stock as the net change in total biomass in the absence of harvest:

$$
\begin{equation*}
S P_{i, t}=B_{i, t+1}-B_{i, t}+C_{i, t} \tag{Eq. 1}
\end{equation*}
$$

where $S P_{i, t}$ is the surplus production for stock $i$ over year $t, B_{i, t}$ and $B_{i, t+1}$ are the biomasses of stock $i$ in years $t$ and $t+1$, respectively, and $C_{i, t}$ is the catch for stock $i$ removed between years $t$ and $t+1$. By including the observed catch in the net change in biomass, surplus production accounts for the effect of fisheries removals on population growth (or decline). We used a Pella-Tomlinson model (17) because it contains a shape parameter ( $p$ ) that allows it to replicate either the Fox $(p \rightarrow 0)$ or Schaefer $(p=1)$ production models $(55,56)$ :

$$
\begin{equation*}
S P_{i, t}=\frac{r_{i}}{p} B_{i, t}\left(1-\left(\frac{B_{i, t}}{K_{i}}\right)^{p}\right)+\varepsilon_{i, t} \tag{Eq. 2}
\end{equation*}
$$

where $r_{i}$ is the intrinsic rate of growth for stock $i, K_{i}$ is the carrying capacity for stock $i$, and $\varepsilon_{i, t}$ is the residual for stock $i$ in year $t$. Residuals are assumed to follow a first-order autocorrelated (AR1) process:

$$
\begin{equation*}
\varepsilon_{i, t}=\rho_{i} \varepsilon_{i, t-1}+\sqrt{1-\rho_{i}^{2}} \delta_{i, t} \tag{Eq. 3}
\end{equation*}
$$

where $\rho_{i}$ is the first-order autocorrelation coefficient for stock $i, \varepsilon_{i, t}$ and $\varepsilon_{i, t-1}$ are the observed residuals around the production function for stock $i$ in years $t$ and $t-1$, respectively, and $\delta_{i, t}$ is a normally distributed random variable representing uncorrelated errors for stock $i$ in year $t$. We do not correct for the lower variance arising for $\varepsilon_{i, t}$ in the first year of data, and instead preclude doing so by estimating $\rho_{i}$ with unbounded support.

We used Akaike Information Criterion (AIC; (21)) to compare models with shape parameters $(p)$ that maximize productivity at $50 \%(p=1.00), 45 \%(p=0.55), 40 \%(p=0.20)$, and $37 \%(p=0.01)$ of carrying capacity and selected the model with the lowest AIC score as the best "standard" model. We evaluated these shape parameter values because $50 \%$ produces the symmetric Schaefer model, $40 \%$ is the meta-analytic mean for fish ( 57 ), and $37 \%$ is the asymptotic limit of this parameterization of the Pella-Tomlinson model.

In this model and in all the models described below, we (1) scaled biomass and production to each stock's maximum biomass to ease model fitting and (2) placed a likelihood penalty on carrying capacities greater than five times the observed maximum biomass to constrain unrealistic values. We fit all models using maximum likelihood estimation in the TMB package (Template Model Builder; (58)) in R (59). See Table S6 for a key to all model symbols.

### 2.3 Base SST-linked surplus production model

To evaluate the influence of temperature on fisheries productivity, we extended the best standard model to include a multiplicative temperature influence term:

$$
\begin{equation*}
S P_{i, t}=\frac{r_{i}}{p} B_{i, t}\left(1-\left(\frac{B_{i, t}}{K_{i}}\right)^{p}\right) * \exp \left(S S T_{i, t} * \theta_{i}\right)+\varepsilon_{i} \tag{Eq. 4}
\end{equation*}
$$

where $S S T_{i, t}$ is the sea surface temperature for stock $i$ in year $t$ (centered on the mean SST for stock $i$ to ease both model fitting and interpretation of the $\theta_{i}$ parameter) and $\theta_{i}$ is the influence of SST on the productivity of stock $i$. We estimated SST influences, $\theta_{i}$, as random effects:

$$
\begin{equation*}
\theta_{i} \sim N\left(\mu_{S S T}, \sigma_{S S T}^{2}\right) \tag{Eq. 5}
\end{equation*}
$$

where $\mu_{\text {SST }}$ and $\sigma_{\text {SST }}$ are the mean and standard deviation of the global distribution of SST influences $\left(\theta_{i}\right)$, respectively. $\theta_{i}<0$ means increasing SST reduces productivity at a given biomass and $\theta_{i}>0$ means increasing SST magnifies productivity at a given biomass.

We used AIC to compare models using SST averages from the COBE, ERSST, and HadISST datasets and selected the model with the lowest AIC score as the best "base" model.

### 2.4 Hierarchical SST-linked surplus production models

To determine whether taxonomy, geography, or stock assessment method structure SST influences, we used SST-linked surplus production models with hierarchical SST influence based on each of five groups in three categories (Table S1): (a) taxonomy (order and family); (b) geography (FAO major fishing area and marine ecoregion); and (c) stock assessment method (Table S7). Marine ecoregions were defined by intersecting Large Marine Ecosystems (LMEs; (60)) and High Seas Areas (HSAs; (61)). These models were identical to the base model except that SST influence was estimated as a nested hierarchical random effect:

$$
\begin{equation*}
\theta_{i, j} \sim N\left(\mu_{G, j}, \sigma_{G}^{2}\right) \tag{Eq. 6}
\end{equation*}
$$

where SST influences $\left(\theta_{i}\right)$ for stock $i$ in group $j$ were drawn from a normal distribution with a group-specific mean ( $\mu_{G, j}$ ) and group-level standard deviation ( $\sigma_{G}$ ). Group-specific means were drawn from a global normal distribution with mean ( $\mu_{S S T}$ ) and standard deviation ( $\sigma_{S S T}$ ):

$$
\begin{equation*}
\mu_{G, j} \sim N\left(\mu_{S S T}, \sigma_{S S T}^{2}\right) \tag{Eq. 7}
\end{equation*}
$$

We compared the group models to the base model using AIC and judged a group to be a significant driver of SST influence if its model exhibited an AIC score more than two points lower than the base model. The best or "final" SST-linked surplus production model was identified as the model producing the lowest AIC score.

### 2.5 Model validation

We tested whether the final SST-linked surplus production model described population dynamics better than the standard surplus production model by competing the models using AIC. We tested whether the results of the final model were an artifact of model structure by decoupling the SST and productivity time series using three null models with different simulated SST time series exhibiting: (1) the same mean, variance, autoregressive properties, and trend as the original time series; (2) the same mean, variance, and autoregressive properties as the original time series but without a trend; and (3) the same mean and variance as the original time series but without autocorrelation or a trend (Fig. S6; Appendix D). The SST simulations were performed using the $R$ package forecast (62). The null models were first evaluated using the true final model, which estimates first-order (AR1) autocorrelation in the residuals; however, with a weakened SST-productivity link and an AR1 process that explains a significant portion of the variability in production, the SST influences were estimated to be near zero and non-significant in all three null model scenarios. To weaken the strength of the AR1 process and better quantify the probability of measuring a significant SST influence by chance, we fixed the AR1 correlation coefficient to zero in all of the null models presented here.

### 2.6 Fixed effects sensitivity analysis

There are compelling arguments for estimating the influence of SST on productivity as either a fixed or random effect. On one hand, estimating SST influence as a fixed effect imposes no constraints on the magnitude and distribution of the influences and could more accurately identify influences that deviate from the patterns exhibited by other stocks in the dataset. On the other hand, estimating SST influence as a random effect could constrain poorly informed and unrealistically large influences. Thus, we evaluated the sensitivity of our results and conclusions to modeling SST influence as a random versus fixed effect. The fixed effects model was identical to the model described by Eq. 4 except that SST influence was estimated as a fixed effect. To measure the extent to which choice of modeling framework affects the results, we compared: (1) the distribution and magnitude of SST influences; (2) the importance of exploitation history, maximum age, and temperature trend in determining the influence of SST on productivity; and (3) the hindcast changes in MSY overall and among marine ecoregions.

The SST influences estimated by the random and fixed effects models were in high agreement on the direction of the influence and were generally correlated in magnitude (Fig. S9); however, and as expected, the fixed effects model estimated larger SST influences for many stocks and estimated SST influences at a higher rate of significance than the random effects model (Figs. S9 \& S10). The fixed effects model estimated absolute SST influences greater than 2.0 for eight stocks (Fig. S11), one of which appears accurate (e.g., Barents Sea capelin) while the others appear spurious. The drivers of SST influence estimates were consistent between the two models: (1) chronic overfishing increases the likelihood of negative impacts of warming on productivity; (2) faster-lived fish are more sensitive, positively and negatively, to warming than slower-lived fish; and (3) the position of a population within its species-specific thermal niche determines its response to warming (Fig. S12). However, the large SST influences estimated by the fixed effects model significantly changed the magnitude of the SST-driven losses in MSY from 1930-2010. The fixed effects model documented a 21.5\%
decline in MSY while the random effects model documented a $3.0 \%$ decline (Fig. S13). Hindcasts of ecoregion-scale changes in MSY were in high agreement on the direction of change and were generally correlated in magnitude (Fig. S14). However, the changes documented by the fixed effects model were larger than the random effects model with a few large departures, especially in the negative direction (Fig. S14).

We favored the random effects model because it improved estimates of SST influence for information-poor stocks, which exert considerable influence on hindcasts of SST-dependent MSY when estimated as fixed effects. The true loss in MSY of the evaluated stocks likely lies between 4.1\% (final random effects model) and 21.5\% (fixed effects model).

### 2.7 Dome-shaped temperature dependence

We attempted to fit two SST-linked surplus production models with dome-shaped temperature dependence - i.e., productivity increases as temperatures warm towards some thermal optimum but decreases once temperatures exceed this optimum - but were unable to achieve convergence with either model. The models attempt to estimate stock-specific (Eq. 8) and species-specific (Eq. 9) thermal optima, respectively:

$$
\begin{align*}
& S P_{i, t}=\frac{r_{i}}{p} B_{i, t}\left(1-\left(\frac{B_{i, t}}{K_{i}}\right)^{p}\right) * \exp \left(-\left(S S T_{i, t}-z_{i}\right)^{2} * \theta_{i}\right)+\varepsilon_{i, t}  \tag{Eq. 8}\\
& S P_{i, t}=\frac{r_{i}}{p} B_{i, t}\left(1-\left(\frac{B_{i, t}}{K_{i}}\right)^{p}\right) * \exp \left(-\left(S S T_{i, t}-z_{j}\right)^{2} * \theta_{i}\right)+\varepsilon_{i, t} \tag{Eq. 9}
\end{align*}
$$

where $S P_{i, t}, B_{i, t}, S S T_{i, t}, r_{i}, K_{i}, p$, and $\varepsilon_{i, t}$ are the same as in the SST-linked surplus production model with monotonic SST influence (Eq. 4), $z_{i}$ and $z_{j}$ are the thermal optima for stock $i$ and species $j$, respectively, and $\theta_{i}$, the SST influence is constrained to be larger than zero (i.e., to ensure that the dome is concave down). Species-specific thermal optima do not allow local adaptation by stocks within a species but increase sample size and estimation power. We suspect that the models failed to converge because the time series are too short ( 39.3 yr mean) and lack sufficient SST contrast ( $1.6^{\circ} \mathrm{C}$ breadth mean) to estimate thermal optima (Fig. S15).

## 3. Data analysis

### 3.1 Drivers of temperature influence

Because the influence of SST on productivity was estimated as a random effect, our estimates of SST influence cannot be considered independent and cannot undergo post-hoc analyses using formal statistical methods (i.e., formal hypothesis testing requires including explanatory variables inside the model, as we did with taxonomy and geography). Therefore, we graphically evaluated whether SST influence is determined by: (1) life history traits such as growth rate, maximum age, or depth preference; (2) behavioral traits such as reproductive
guild, migratory habits, or spawning behavior; (3) stock characteristics such as trend in biomass or fishing pressure; and (4) thermal experience such as mean SST, SST trend, or latitude. A list of evaluated explanatory variables and their sources is provided in Table S3. We could not include these drivers inside the model, as we did with taxonomy and geography, due to missing data for many of the evaluated explanatory variables (Table S3).

### 3.2 Hindcasting maximum sustainable yield

We used the final model's estimates of $p, r_{i}, K_{i}$, and $\theta_{i}$ to hindcast SST-dependent maximum sustainable yield (MSY) from 1930-2010 (Appendices E-G). We expanded the equation for MSY from the Pella-Tomlinson surplus production model:

$$
\begin{equation*}
M S Y=\frac{r * k}{(p+1)^{(p+1) / p}} \tag{Eq. 10}
\end{equation*}
$$

to include the SST influence term and calculated MSY for stock $i$ in year $t$ as:

$$
\begin{equation*}
M S Y_{i, t}=\frac{\left[\left(\exp \left(\widehat{\theta}_{i} * \overline{S S T}_{i, t}\right) * r_{i}\right]_{i} * k_{i}\right.}{(p+1)(p+1) / p} \tag{Eq. 11}
\end{equation*}
$$

where $\overline{S S T}_{i, t}$ is $S S T_{i, t}$ centered on the mean of the SST data used in model fitting and $\hat{\theta}_{i}$ is randomly drawn from a multivariate normal distribution described by the mean $\theta_{i}$ estimate and the $\theta_{i}$ covariance matrix. We bootstrapped $10,000 \mathrm{MSY}$ hindcasts for each stock to generate median MSY trends and confidence intervals. We assessed changes in MSY over the hindcast period using (1) Thiel-Sen regression slopes and (2) percent change in mean MSY from 1930-39 to 2001-2010. Theil-Sen regression, a form of robust regression, identifies the median slope of lines through all possible point pairs and is insensitive to outliers and endpoints in short time series. We limited the hindcast from 1930-2010 to minimize the extrapolation of MSY predictions to temperatures cooler or warmer than those used in model fitting (Fig. S24) and explored the sensitivity of measures of MSY change to the selection of hindcast window (Fig. S25).

### 3.3 Extrapolating vulnerability of global fish populations

We identified global fish populations in the FAO landings database (1) that are likely to be vulnerable to ocean warming as populations that (1) are overfished; (2) have experienced warming; and (3) are located at the warm end of their species-specific thermal niche.

We analyzed the 1,740 FAO fish stocks (FAO area-country-species triples) meeting the following criteria: marine wild capture fisheries for finfish and invertebrates with taxonomic identification resolved to the species-level and with catch time series $\geq 20 \mathrm{yrs}$ and $\geq 250 \mathrm{mt}$ of median annual catch after trimming years of zero catch from the beginning of the time series. We excluded: (1) stocks of invertebrate species that frequently lack the life history data required for the catch-only stock assessment model used to determine stock status (e.g.,
barnacles, corals, sea cucumbers, sea urchins, starfish, sponges); (2) stocks of highly migratory species whose population dynamics cannot be described by catch within a single country's exclusive economic zone (e.g., tuna, marlin, swordfish); and (3) stocks targeted by a distant water fleet whose catch time series are unlikely to be representative of total removals from that population (i.e., stocks whose FAO area and EEZ don't overlap were excluded).

We determined stock status using catch-MSY (63), the best performing individual catchonly stock assessment method (64). Catch-MSY is a stock reduction analysis that reconstructs historical abundance by simulating biomass trajectories that could produce the observed catch time series given priors on initial and final year depletion and stock dynamics such as carrying capacity, $K$, and intrinsic growth rate, $r$. The method establishes priors for $r$ based on population resilience (see below), $K$ based on maximum catch (e.g., between $C_{\text {max }}$ and $100^{*} C_{\max }$ ), and initial and final year depletion based on the ratio of initial and final year catch to the maximum catch. It then estimates "viable" pairs of $r$ and $K$ (i.e., pairs that do not allow the stock to collapse or exceed carrying capacity), generates biomass trends for each pair, and estimates $B / B_{\text {MSY }}$ as the median trend. We calculated the mean $\mathrm{B} / \mathrm{B}_{\text {MSY }}$ of each population over the last 25 years (19912015) and defined overfished as $B / B_{M S Y}<0.5$. The model converged for 1530 of the 1740 stocks.

In addition to a catch time series, cMSY requires an estimate of population resilience (i.e., the ability of a population to recover from disturbance) to establish a prior for the intrinsic growth rate. We derived resilience estimates for all species using a combination of FishBase (65), SeaLifeBase (66), and FishLife (67) life history information. We used the rfishbase package (68) to download Von Bertalanffy growth parameters, maximum size, and vulnerability and resilience from FishBase (FB, for finfish) and SeaLifeBase (SLB, for invertebrates). We used the FishLife package to estimate the Von Bertalanffy growth parameters for all finfish species. FishLife uses a multivariate model trained on FishBase to predict eight life history traits for $>32,000$ fish. We classified species into resilience categories (Table S8) using, in order of preference, resilience values: (1) reported on FB/SLB; (2) derived from the FishLife Von Bertalanffy growth parameter; (3) derived from the FB/SLB Von Bertalanffy growth parameter; (4) derived from the FB/SLB vulnerability metric; (5) derived from the FB/SLB Von Bertalanffy maximum age; (6) derived from the genus mode; or (7) derived from the family mode.

We estimated the temperature experienced by each population using the COBE sea surface temperature dataset (18) and the spatial boundary of the population (i.e., the intersection of the FAO region and exclusive economic zone). We calculated the mean annual temperature and the trend in annual temperature over the last 25 years (1991-2015). We estimated the position of each population in its species-specific thermal niche as the percentile of the population's mean temperature experience relative to the temperature experiences of other conspecific populations. We identified populations above the $80^{\text {th }}$ percentile as being at the warm end of their species-specific thermal niche.

## Supplementary Text

## Stock assessment models and treating output as "data"

## Overview of the assessment model types included in our analysis

Stock assessments are population models that combine different sources of information (catch, relative abundance, and life history) to estimate population size and harvest rates over time, as well as the reference points used in management. Different assessment models are used based on the available data for a stock. The four categories of commonly used assessment model types in our analysis were (Table S7): biomass dynamic models (BDM), virtual population analyses (VPA), statistical-catch-at-age models (SCAA), and integrated analyses (IA).

BDMs track changes in total biomass from one year to the next, ignoring size- or agestructure in the population. Changes in biomass are due to the observed catch and estimated surplus production in a given year (assumed to be a function of biomass in that year). BDMs require annual estimates of total catch and relative abundance (CPUE, catch-per-unit-effort) to estimate the parameters controlling production and biomass over time.

VPA and SCAA models, on the other hand, assume that population dynamics are agestructured, with recruitment events producing different-sized cohorts in the population over time. VPAs use observed catch-at-age (assumed known without error) and an assumed natural mortality rate $(M)$ to reconstruct historical cohort abundance using a backwards tuning procedure (tuned to available CPUE data), and total biomass each year is estimated using the estimated numerical abundance of each cohort in that year times an assumed weight-at-age. In contrast, SCAA models are forward projecting, and allow for catch data to be uncertain. Annual cohort sizes and other parameters are estimated statistically using maximum likelihood or Bayesian approaches and fit to the available catch-at-age and CPUE information. Total biomass is estimated using the estimated numerical abundance and an assumed weight-at-age. Like VPAs, SCAA models often use an assumed $M$ (either fixed over time or ages, or both), although $M$ can sometimes be an estimated parameter.

IA is a general assessment approach that aims to incorporate all the available data in as raw a form as possible into a single analysis via joint likelihood functions (69). The vast majority of assessments classified as IA in Table S7 (49 of 57) use the age-based Stock-Synthesis program (70), and are analogous to SCAA models in how annual estimates of biomass are produced. A key distinction between these IA models and those labeled SCAA here is that many of the inputs assumed fixed in SCAA may be estimated within the IA model, accounting for uncertainty in these quantities (e.g., weight-at-age, ageing error in catch-at-age data, etc.).

Problems with and recommendations for using stock assessment output as data
Maunder and Punt (69) outline five problems with using model output as data: (1) loss of information when converting data to model output; (2) inconsistent assumptions between
the meta-analysis model and stock assessment model; (3) challenges in identifying a statistical likelihood for model output when treated as data; (4) difficulties in representing precision of model output (i.e., due to covariance or non-normal distribution of model estimates); and (5) reduced ability to diagnose goodness of fit for the meta-analysis model.

Thorson et al. (47) provide six recommendations for conducting fisheries meta-analyses: (R1) choose appropriate model complexity and sample size; (R2) use multiple lines of evidence to support a hypothesis or interpretation; (R3) consider alternative hypotheses; (R4) strive for single-stage meta-analysis; (R5) account for experimental, parametric, and functional variability; and (R6) identify the desired type(s) of inference and proceed accordingly.

Brooks and Deroba (46) suggest that post-hoc analyses that use model output as data frequently fail to account for the assumptions, uncertainties, and biases of the original assessment models. They provide the following five recommendations: (R7) avoid using model output as data; (R8) collaborate with lead assessment scientists; (R9) conduct sensitivity analyses; (R10) use errors-in-variables methods; and (R11) use cross-validation methods.

## How we addressed these challenges

We mitigated the problems associated with using stock assessment output as data in fisheries meta-analyses by: (R1) using a simple model given our low sample size (235 time series); (R2) using multiple lines of evidence to corroborate our results including repeating the analysis with multiple SST datasets, parameterizations, modeling frameworks, and metrics of MSY change; (R6) clearly stating our intent to make global-, group- (e.g., family, ecoregion), and individual-scale (i.e., stock) inferences about the effect of SST on productivity. We also addressed the assumptions, uncertainties, and biases in using stock assessment output as data by trimming or removing time series produced with strict assumptions, removing trimmed assessments with short time series (<20 yr), using a mixed-effects model to share information between stocks and constrain poorly informed estimates, and explicitly examining the influence of assessment method on our results. The results were not affected by stock assessment method and were not sensitive to SST dataset, model parameterization, or metric of MSY change. Following (R1), we recommend that future region-specific studies validate our results using meta-analytic models that leverage the availability of higher resolution data at smaller spatial scales.

We were unable to: (R3) consider alternative environmental drivers of fisheries productivity given the lack of historic, globally complete data on environmental variables besides SST; (R4/R7) use a single-stage meta-analysis given the global-intent of our study and the lack of unprocessed data (i.e., survey data) at the global-scale; (R5/R10) incorporate reported measures of uncertainty given that this data is not included in the RAMLDB and not reported in a consistent manner in stock assessments; (R8) collaborate with lead assessment scientists for all 235 assessments represented in our study; (R9) conduct sensitivity analyses using alternative biomass estimates or confidence intervals for biomass estimates given that this data is not included in the RAMLDB and not reported in a consistent manner in stock
assessments; or (R11) use cross-validation methods to quantify uncertainty since withholding a testing dataset truncates our already short time series (40 yr mean). We know of only one published meta-analysis that used a single-stage analysis to estimate stock-recruit relationships for multiple populations simultaneously (71) and this analysis involved fewer than ten species.

## Supplemental Figures



Fig. S1. Mean global SST based on the COBE v2, ERSST v4, and HadISST v.1.1 datasets. Gray lines show monthly means and black lines show annual means. Horizontal dashed lines show mean ocean temperature from 1880-present. The discontinuity in the HadISST dataset in the 1980s is likely due to problems at the poles and dateline and problems with the bias correction algorithms and compromises the usefulness of the dataset for our purposes.


Fig. S2. Comparison of SST influence estimates from SST-linked surplus production models using the COBE v2, ERSST v4, and HadISST v1.1 datasets. In the top panels, points show mean estimates and error bars show $95 \%$ confidence intervals. Significant positive and negative SST influences are shown in blue and red, respectively. The shaded grey column indicates the 95\% confidence interval for the global mean of the SST influences. In the bottom panels, the diagonal line is the one-to-one line for pairwise comparisons of SST influence estimates from models using each SST dataset.


Fig. S3. Distribution of SST influences estimated by SST-linked surplus production models with hierarchy on SST influence. Hierarchical models are structured by (A) taxonomic order and (B) taxonomic family, (C) FAO major fishing area, (D) marine ecoregion, and (E) stock assessment method. Points show mean estimates and error bars show $95 \%$ confidence intervals. Significant positive and negative SST influences are shown in blue and red, respectively. The shaded grey column indicates the $95 \%$ confidence interval for the global mean of the SST influences.


Fig. S4. Correlation between SST influences estimated by the base model and five hierarchical
models. Hierarchical models are organized by taxonomy (order/family), geography (FAO area/marine ecoregion), or stock assessment method. Diagonal line is the one-to-one line.


Fig. S5. Mean of the SST influence distributions for stock assessment methods in the SSTlinked surplus production model with hierarchy on SST influence by stock assessment method. Points show mean estimates and error bars show $95 \%$ confidence intervals. None of the SST influence means were significantly different from zero and the model gained less support than the base model (see Table S1).


Fig. S6. Example (A) observed and (B-D) simulated SST time series (US West Coast, black rockfish). The simulated SST time series were used in the three null models.


Fig. S7. Distribution of SST influences estimated by the final model and three null models. Points show mean estimates and error bars show $95 \%$ confidence intervals. Significant positive and negative SST influences are shown in blue and red, respectively.


Fig. S8. Distribution of intrinsic rate of growth ( $r_{i}$ ), carrying capacity ( $\mathrm{K}_{\mathrm{i}}$ ), SST influence ( $\theta_{\mathrm{i}}$ ), residual process variability ( $\sigma_{\mathrm{p}, \mathrm{i}}$ ), and first-order (AR1) autocorrelation coefficient ( $\rho_{\mathrm{i}}$ ) estimates from the final model. Points show mean estimates and lines show $95 \%$ confidence intervals. Carrying capacity is a multiple of the maximum observed biomass (e.g., a carrying capacity of 1 , shown by the vertical dotted line, means that the carrying capacity is equivalent to the maximum observed biomass).


Fig. S9. Comparison of SST influences estimated by the fixed and random effects models.
Plots show (A) correlation between the random and fixed effects estimates and histograms of the (B) random and (C) fixed effects estimates.


Fig. S10. Distribution of SST influences estimated by the fixed and random effects models. Points show mean estimates and error bars show $95 \%$ confidence intervals. Significant positive and negative SST influences are shown in blue and red, respectively.


Fig. S11. Inspection of the stocks exhibiting SST influence estimates greater than 2.0. Blue points represent cooler than average years and red points represent warmer than average years. Black lines show the surplus production curves at each stock's average temperature. Blue and red lines show surplus production curves at temperatures progressively cooler and warmer than the average, respectively. Stocks with positive SST influences are more productive at warmer temperatures (red curves on top) and stocks with negative SST influences are more productive at cooler temperatures (blue curves on top). SST influences ( $\theta_{\mathrm{i}}$ ) are shown in the top-left corner of each plot and are colored to indicate the direction and significance of the SST influence (blue=positive, red=negative, bold=significant).


Fig. S12. SST influence as a function of (A) exploitation history, (B) maximum age, and (C) position of a population in its species-specific thermal niche. Panel (A) shows more and larger negative influences of warming for populations with histories of overfishing. Points represent individual populations and are colored by significance (blue=positive, red=negative, grey=nonsignificant). Solid lines show the $50^{\text {th }}$ percentile quantile regression fit and dashed lines show the $2.5 \%$ and $97.5 \%$ quantile regression fits. $F / F_{\text {MSY }}$ is the ratio of fishing mortality ( $F$ ) to the fishing mortality that produces maximum sustainable yield ( $F_{\text {MSY }}$ ): values greater than one indicate overfishing. Panel (B) shows larger and more significant influences of temperature for populations of species with faster life histories (i.e., shorter lifespan). Points and lines as in Panel (A). Panel (C) shows increasingly negative influences for populations at the warm end of their thermal niche for the two species with $\geq 10$ populations. Lines show Theil-Sen regression fits. Theil-Sen regression, a form of robust regression, identifies the median slope of lines through all possible point pairs and is insensitive to outliers and endpoints in small datasets.

Fixed effects


## Random effects



Fig. S13. Hindcast of SST-dependent maximum sustainable yield (MSY, mt=metric tons) using the fixed and random effects models. Solid lines indicate the median MSY estimates, shading indicates the $95 \%$ confidence intervals, and horizontal dashed lines indicate MSY at average temperature. Percent decline from 1930-39 to 2001-10 is shown in the top-right corner.


Fig. S14. Percent change in mean maximum sustainable yield (MSY) from 1930-39 to 2001-10 by ecoregion predicted by the fixed versus random effects models. Points are scaled to the MSY of the ecoregion and the number of stocks in the ecoregion is shown inside each point. The solid line indicates the one-to-one line.


Fig. S15. Histograms showing (A) breadth of SST experience, (B) length of time series, and the (C) start and (D) end year of time series for stocks used in the analysis. Mean values are indicated by the vertical dashed line (median values shown for start and end years).


Fig. S16. Map showing the global distribution of SST influences. Points were jittered to expose overlapping stock centroids. Dashed lines indicate FAO major fishing areas.


Fig. S17. Mean of the SST influence distributions for geographic or taxonomic groups in models with hierarchy on SST influence. Hierarchical models are structured by (A) marine ecoregion, (B) FAO major fishing area, (C) taxonomic family, and (D) taxonomic order. Points show mean estimates and error bars show $95 \%$ confidence intervals. Significant positive and negative SST influences are shown in blue and red, respectively. All but the taxonomic order model had more support than the base model.


Fig. S18. SST influence as a function of nine stock characteristics. SST influences are colored by significance (blue=positive, red=negative, grey=non-significant). Solid lines show the $50^{\text {th }}$ percentile quantile regression fit and dashed lines show the $2.5 \%$ and $97.5 \%$ quantile regression fits. Sample size is shown in the bottom-right corner if data were not available for all 235 stocks.


Fig. S19. SST influence as a function of nine life history traits. Life history traits are the: Brody growth coefficient (K), asymptotic maximum length ( $L_{\text {inf }}$ ), asymptotic maximum weight ( $\mathrm{W}_{\text {inf }}$ ), natural mortality (M), maximum age ( $T_{\max }$ ), age at maturity ( $T_{\text {mat }}$ ), length at maturity ( $L_{\text {mat }}$ ), trophic level, and median depth. SST influences are colored by significance (blue=positive, red=negative, grey=non-significant). Solid lines show the $50^{\text {th }}$ percentile quantile regression fit and dashed lines show the $2.5 \%$ and $97.5 \%$ quantile regression fits. Sample size is shown in the bottom-right corner if data were not available for all 235 stocks.


Fig. S20. Distribution of SST influences among (A) specific and (B) generic habitat types.
Brown and blue boxplot shading corresponds to demersal and pelagic habitats, respectively. Black numbers indicate total number of stocks for each habitat type. Blue and red numbers show the number of stocks with a positive and negative SST influence, respectively.


Fig. S21. Distribution of SST influences among reproductive strategies and migratory and spawning behaviors. Grey numbers indicate total number of stocks in each group.


Fig. S22. SST influence as a function of the mean temperature experienced by stocks of the same species for the seven species with $\geq 5$ stocks in the analysis. Lines show Theil-Sen regression fits. Theil-Sen regression, a form of robust regression, identifies the median slope of lines through all possible point pairs and is insensitive to outliers and endpoints in small datasets.


Fig. S23. SST influence as a function of the latitude of stocks of the same species for the seven species with $\geq 5$ stocks in the analysis. Lines show Theil-Sen regression fits. Theil-Sen regression, a form of robust regression, identifies the median slope of lines through all possible point pairs and is insensitive to outliers and endpoints in small datasets.


Fig. S24. The (A\&B) frequency of SST extrapolation by the hindcast model and (C) correlation between MSY estimates from the final model and data-rich stock assessments. In (A), each row shows the SST experience of an individual stock where black years were used in model development, grey years experienced temperatures also experienced during model years, and blue and red years experienced temperatures cooler and warmer than those experienced during model years, respectively. In (B), the blue and red shading show the percentage of years experiencing temperatures cooler and warmer than those experienced during model years, respectively. The hindcast model generally extrapolates for fewer than $15 \%$ (dashed line) of years between 1930-2010. In (C), the diagonal line is the one-to-one line.


Fig. S25. Sensitivity of hindcasted changes in MSY (mt=metric tons) to the determination of the hindcast window. Time series showing (A) mean global SST anomaly, (B) hindcast of SSTdependent maximum sustainable yield (MSY) for all stocks included in the analysis, (C) Thiel-Sen regression slope when evaluating MSY trends beginning in each year from 1850-1990 and ending in 2010, and (D) percent difference in MSY when comparing the mean MSY over the 10 years following each year from 1850-1990 and the mean MSY from 2001-2010. In (A), the grey shading indicates the hindcast window determined to minimize extrapolation to temperatures outside those included in the final model. In (B), the dark line shows a Thiel-Sen regression fit to the MSY time series in the hindcast window. In (C) and (D), the labeled points mark the measures of MSY change experienced over the hindcast window.


Fig. S26. Ecoregion-scale trends in maximum sustainable yield (MSY) related to ecoregion (A) latitude, (B) mean temperature, and (C) temperature trend.


Fig. S27. Comparison of ecoregion-scale changes in fisheries productivity estimated by Britten et al. (14) and the present study. Britten et al. (14) quantify the meta-analytic mean trend in recruitment potential (Rmax). Comparable values derived from the present study are: (A) change in scaled MSY (MSY divided by maximum MSY) per decade from 1930-2010; (B) percent difference in mean MSY from 1930-39 to 2001-2010; and (C) the meta-analytic mean of the SST influences of stocks in an ecoregion multiplied by the change in temperature from 1930-2010 in the ecoregion. In both studies, negative and positive values represent a negative and positive change, respectively. Blue and red points indicate ecoregions where both studies agree that change has positively and negatively impacted productivity, respectively. Grey points indicate ecoregions in which the studies disagree on the direction of productivity change. The present study describes SST influence for ten ecoregions not described in the Britten study (Bay of Biscay, Canary Current, Greenland Sea, Humboldt Current, Kuroshio Current, Labrador Sea, Mediterranean Sea, North Brazil Shelf, South Atlantic Ocean, and West Bering Sea) and the Britten study describes SST influence on one ecoregion not described in the present study (EastCentral Australian Shelf).


Fig. S28. Distribution of FAO stocks included in the vulnerability analysis. Each stock is a FAO area-country-species triple.


Fig. S29. Indicators of the vulnerability of populations to ocean warming. Points represent individual populations and are colored by the position of the population within its speciesspecific thermal niche (deep blue=coolest end of range; deep red=warmest end of range). Deep red populations in the bottom-right quadrant have experienced warming (positive temperature trend), overfishing ( $\mathrm{B}^{2} \mathrm{~B}_{\text {MSY }}<0.5$ ), and are located in the warm end of their thermal niche and are the most likely to be vulnerable to ocean warming.


Fig. S30. Number and total catch of populations vulnerable to warming by country (FAO region-country exclusive economic zone intersect). Points are scaled and colored based on the total catch of vulnerable populations in each country and the number of vulnerable populations in each country is shown inside the point. Zeros indicate countries without any vulnerable populations. Dashed lines indicate FAO major fishing areas.

## Supplemental Tables

Table S1. AIC of candidate surplus production models (PT=Pella-Tomlinson).

| Model | K | Likelihood | AIC | AAIC |
| :--- | :--- | :--- | :--- | ---: |
| Question 1. Does asymmetry matter? |  |  |  |  |
| PT model (MSY@45\%K) (standard model) | 940 | -19280.4 | -36680.9 | 0.0 |
| Schaefer model (MSY@50\%K) | 940 | -19278.9 | -36677.8 | 3.1 |
| PT model (MSY@40\%K) | 940 | -19271.3 | -36662.6 | 18.3 |
| PT model (MSY@37\%K) | 940 | -19261.1 | -36642.1 | 38.8 |
| Question 2. Does temperature matter? |  |  |  |  |
| COBE SST-linked PT model (base model) | 942 | -19300.4 | -36716.9 | 0.0 |
| HadISST SST-linked PT model | 942 | -19290.5 | -36697.0 | 19.9 |
| ERSST SST-linked PT model | 942 | -19289.2 | -36694.5 | 22.4 |
| PT model (MSY@45\%K) (standard model) | 940 | -19280.4 | -36680.9 | 36.0 |
| Question 3. Does group hierarchy matter? |  |  |  |  |
| SST-linked PT model w/ hierarchy by ecoregion (final model) | 943 | -19312.0 | -36738.0 | 0.0 |
| SST-linked PT model w/ hierarchy by FAO area | 943 | -19309.3 | -36732.7 | 5.3 |
| SST-linked PT model w/ hierarchy by assessment method (specific) | 943 | -19306.4 | -36726.7 | 11.3 |
| SST-linked PT model w/ hierarchy by family | 943 | -19305.3 | -36724.7 | 13.3 |
| SST-linked PT model (base model) | 942 | -19300.4 | -36716.9 | 21.1 |
| SST-linked PT model w/ hierarchy by order | 943 | -19300.6 | -36715.2 | 22.8 |
| SST-linked PT model w/ hierarchy by assessment method (generic) | 943 | -19300.4 | -36714.9 | 23.1 |
| Question 4. Null model tests |  |  |  |  |
| SST-linked PT model w/ hierarchy by LME (final model) | 943 | -19312.0 | -36738.0 | 0.0 |
| SST-linked PT model w/ hierarchy by LME - Null SST \#2 | 943 | -19285.1 | -36684.1 | 53.8 |
| SST-linked PT model w/ hierarchy by LME - Null SST \#1 | 943 | -19283.7 | -36681.3 | 56.7 |
| SST-linked PT model w/ hierarchy by LME - Null SST \#3 | 943 | -19280.5 | -36674.9 | 63.0 |

Table S2. Stocks whose productivity is significantly influenced by ocean warming (sorted from most positive to most negative temperature influence).

| Stock id | Species | Area | $\theta_{i}$ |
| :---: | :---: | :---: | :---: |
| GHAL4RST | Greenland halibut (Reinhardtius hippoglossoides) | Gulf of St. Lawrence | 0.51 |
| COD3Pn4RS | Atlantic cod (Gadus morhua) | Northern Gulf of St. Lawrence | 0.46 |
| HERR30 | Atlantic herring (Clupea harengus) | Bothnian Sea | 0.42 |
| SCALLGB | Sea scallop (Placopecten magellanicus) | Georges Bank | 0.40 |
| HERRRIGA | Atlantic herring (Clupea harengus) | Gulf of Riga East of Gotland | 0.31 |
| PANDALGOM | Northern shrimp (Pandalus borealis) | Gulf of Maine | 0.27 |
| WHAKEGBGOM | White hake (Urophycis tenuis) | Gulf of Maine / Georges Bank | 0.24 |
| SPANMACKSATLC | Spanish mackerel (Scomberomorus maculatus) | Southern Atlantic coast | 0.18 |
| BSBASSMATLC | Black sea bass (Centropristis striata) | Mid-Atlantic Coast | 0.16 |
| KINGKLIPSA | Kingklip (Genypterus capensis) | South Africa | -0.11 |
| ALBANATL | Albacore tuna (Thunnus alalunga) | Northern Atlantic | -0.20 |
| ARFLOUNDBSAI | Arrowtooth flounder (Atheresthes stomias) | Bering Sea and Aleutian Islands | -0.20 |
| PLAIC7d | European Plaice (Pleuronectes platessa) | Eastern English Channel | -0.25 |
| SOLEIIIa | Common sole (Solea solea) | Kattegat and Skagerrak | -0.27 |
| PLAICECHW | European Plaice (Pleuronectes platessa) | Western English Channel | -0.28 |
| CODVIIek | Atlantic cod (Gadus morhua) | Celtic Sea | -0.29 |
| HERRSIRS | Atlantic herring (Clupea harengus) | ICES VIIa-g-h-j | -0.32 |
| HERRNS | Atlantic herring (Clupea harengus) | North Sea | -0.33 |
| WHITNS-VIId | Whiting (Merlangius merlangus) | IV and VIId | -0.35 |
| HADNS-IIIa | Haddock (Melanogrammus aeglefinus) | Illa and North Sea | -0.36 |
| POLLNS-VI-IIIa | Saithe (Pollachius virens) | IIIa, VI and North Sea | -0.36 |
| SOLEIS | Common sole (Solea solea) | Irish Sea | -0.39 |
| SEELNSSA1 | Sand eel (Ammodytes marinus) | North Sea | -0.42 |
| CODNS | Atlantic cod (Gadus morhua) | North Sea | -0.44 |
| CODVIa | Atlantic cod (Gadus morhua) | West of Scotland | -0.45 |
| SEELNSSA2 | Sand eel (Ammodytes marinus) | North Sea | -0.47 |
| SEELNSSA3 | Sand eel (Ammodytes marinus) | North Sea | -0.50 |
| CODIS | Atlantic cod (Gadus morhua) | Irish Sea | -0.54 |

Table S3. Potential predictors of temperature influence and their sources (percentage of stocks with predictor available shown in parenthesis when coverage is incomplete).

| Variable | Source |
| :---: | :---: |
| SST experience |  |
| SST average ( ${ }^{\circ} \mathrm{C}$ ) | COBE SST + stock boundary database (1930-2010) |
| SST trend ( ${ }^{\circ} \mathrm{C} / \mathrm{yr}$ ) | COBE SST + stock boundary database (1930-2010) |
| Latitude (absolute value) | Centroid of the stock area (stock boundary database) |
| Stock characteristics |  |
| Biomass average (MT) | RAM Legacy Database |
| Scaled biomass trend (scaled MT/yr) | RAM Legacy Database |
| Stock area (sq. km) | Stock boundary database |
| Time series length (year) | RAM Legacy Database |
| $\mathrm{B} / \mathrm{B}_{\text {ms }}$ average | RAM Legacy Database (52\%) |
| F/Fmsy average | RAM Legacy Database (57\%) |
| Geography |  |
| Marine ecoregion | Containing the centroid of the stock area |
| FAO Major Fishing Area | Containing the centroid of the stock area |
| Life history traits |  |
| Taxonomy (family/order) | RAM Legacy Database (corrected for errors) |
| Natural mortality rate ( $\mathrm{M}, 1 / \mathrm{yr}$ ) | FishLife (finfish, 100\%), SeaLifeBase (inverts, 19\%) |
| Brody growth coefficient (K) | FishLife (finfish, 100\%), SeaLifeBase (inverts, 100\%) |
| Asymptotic maximum length ( Linf cm ) | FishLife (finfish, 100\%), SeaLifeBase (inverts, 38\%) |
| Asymptotic maximum mass ( $\mathrm{W}_{\text {inf }}, \mathrm{kg}$ ) | FishLife (finfish, 100\%), SeaLifeBase (inverts, 24\%) |
| Length at maturity ( $L_{\text {mat }}, \mathrm{cm}$ ) | FishLife (finfish, 100\%), SeaLifeBase (inverts, 0\%) |
| Age at maturity ( $\mathrm{T}_{\text {mat }}, \mathrm{yr}$ ) | FishLife (finfish, 100\%), SeaLifeBase (inverts, 0\%) |
| Maximum age ( $\mathrm{T}_{\text {max, }} \mathrm{yr}$ ) | FishLife (finfish, 100\%), SeaLifeBase (inverts, 19\%) |
| Trophic level | FishBase (finfish, 93\%), SeaLifeBase (inverts, 19\%) |
| Habitat (e.g., demersal, pelagic, etc.) | FishBase (finfish, 99\%), SeaLifeBase (inverts, 95\%) |
| Depth (m) | FishBase (finfish, 95\%), SeaLifeBase (inverts, 0\%) |
| Behavioral traits |  |
| Migratory behavior (e.g., catadromous, etc.) | Fishbase (finfish, 69\%), SeaLifeBase (inverts, 0\%) |
| Reproductive mode (i.e., dioecism or protogyny) | Fishbase (finfish, 96\%), SeaLifeBase (inverts, 95\%) |
| Reproductive guild 1 (e.g., bearers, guarders, etc.) | Fishbase (finfish, 91\%), SeaLifeBase (inverts, 71\%) |
| Reproductive guild 2 (e.g., nesters, brooders, etc.) | Fishbase (finfish, 82\%), SeaLifeBase (inverts, 67\%) |
| Spawning ground (e.g., coastal, shelf, etc.) | Fishbase (finfish, 62\%), SeaLifeBase (inverts, 0\%) |
| Spawning frequency | Fishbase (finfish, 55\%), SeaLifeBase (inverts, 5\%) |

Table S4. Hindcasted changes in SST-dependent maximum sustainable yield (MSY) from 1930-2010 among ecoregions (LME=large marine ecosystem; HSA=high seas area; sorted by ascending percent difference).

| Type | Ecoregion | \# of stocks | $\begin{array}{r} \text { Total MSY } \\ (1000 \mathrm{~s} \text { of } \mathrm{mt}) \\ \hline \end{array}$ | $\begin{array}{r} \text { Mean } \\ \text { sst }\left({ }^{\circ} \mathrm{C}\right) \\ \hline \end{array}$ | SST trend ( ${ }^{\circ} \mathrm{C} /$ decade) | $\begin{array}{r} \text { SST } \\ \text { influence }\left(\boldsymbol{\theta}_{\mathrm{i}}\right) \\ \hline \end{array}$ | MSY change |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  |  |  |  |  | mt / decade | \% difference |
| LME | Sea of Japan | 1 | 30.9 | 12.3 | 0.134 | -0.14 | -1119.7 | -34.7 |
| LME | North Sea | 9 | 2454.6 | 9.8 | 0.082 | -0.36 | -78384.5 | -34.6 |
| LME | Iberian Coastal | 2 | 3.1 | 16.2 | 0.085 | -0.14 | -59.8 | -19.2 |
| LME | Kuroshio Current | 6 | 3720.8 | 22.1 | 0.121 | -0.15 | -67403.0 | -17.4 |
| LME | Celtic-Biscay Shelf | 15 | 294.0 | 12.5 | 0.064 | -0.25 | -3278.4 | -15.2 |
| LME | East China Sea | 5 | 1907.5 | 21.5 | 0.142 | -0.05 | -17185.3 | -8.3 |
| LME | Benguela Current | 3 | 176.6 | 19.6 | 0.079 | -0.05 | -1120.1 | -6.0 |
| HSA | South Atlantic Ocean | 1 | 25.2 | 14.4 | 0.075 | -0.07 | -180.2 | -5.3 |
| LME | Southeast U.S. Continental Shelf | 9 | 20.4 | 25.4 | -0.115 | 0.09 | -176.0 | -5.0 |
| HSA | North Atlantic Ocean | 6 | 328.8 | 20.4 | 0.023 | -0.11 | -1440.5 | -4.7 |
| LME | Faroe Plateau | 3 | 86.7 | 9.2 | 0.018 | -0.07 | 48.2 | -3.5 |
| LME | Iceland Shelf and Sea | 5 | 1261.0 | 4.6 | 0.031 | -0.04 | -554.0 | -3.0 |
| LME | Agulhas Current | 5 | 761.2 | 25.0 | 0.076 | -0.04 | -2503.2 | -3.0 |
| LME | Gulf of Alaska | 20 | 356.2 | 9.0 | 0.014 | -0.01 | -586.8 | -2.1 |
| LME | East Bering Sea | 14 | 3297.4 | 4.7 | 0.038 | -0.02 | -5306.8 | -2.1 |
| HSA | Labrador Sea | 2 | 138.5 | 4.2 | 0.042 | -0.05 | 66.7 | -2.0 |
| LME | Greenland Sea | 1 | 1029.3 | 1.1 | 0.031 | -0.01 | -108.1 | -0.2 |
| LME | Gulf of Mexico | 3 | 2.0 | 25.9 | 0.000 | 0.01 | -0.7 | -0.2 |
| LME | Humboldt Current | 16 | 10178.6 | 15.1 | 0.077 | -0.01 | 28.6 | -0.2 |
| LME | North Brazil Shelf | 1 | 17.3 | 27.7 | 0.022 | 0.00 | -1.1 | -0.1 |
| LME | California Current | 29 | 286.5 | 16.5 | 0.034 | 0.02 | -14.5 | -0.1 |
| LME | Patagonian Shelf | 2 | 349.2 | 10.3 | 0.059 | 0.01 | 270.7 | 0.3 |
| LME | Mediterranean Sea | 2 | 40.1 | 19.7 | 0.025 | 0.01 | 11.3 | 0.6 |
| HSA | Bay of Biscay | 1 | 7.2 | 15.0 | 0.084 | 0.01 | 6.2 | 1.0 |
| LME | Norwegian Sea | 1 | 1262.7 | 6.8 | 0.040 | 0.01 | 614.4 | 1.1 |


| LME | Barents Sea | 5 | 2633.8 | 1.1 | 0.079 | 0.03 | 3967.5 | 1.7 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| LME | West Bering Sea | 1 | 143.3 | 4.3 | 0.037 | 0.03 | 35.7 | 1.8 |
| LME | Scotian Shelf | 2 | 41.0 | 7.2 | 0.074 | 0.04 | 94.0 | 1.9 |
| HSA | North Pacific Ocean | 4 | 149.6 | 22.1 | 0.036 | -0.01 | 304.4 | 2.0 |
| LME | New Zealand Shelf | 9 | 5.8 | 15.1 | 0.049 | 0.03 | 18.5 | 3.0 |
| LME | Canary Current | 2 | 712.9 | 21.0 | 0.066 | 0.04 | 2228.8 | 4.0 |
| HSA | South Pacific Ocean | 5 | 86.7 | 18.1 | 0.046 | 0.10 | 387.7 | 4.2 |
| LME | South West Australian Shelf | 5 | 2.8 | 17.0 | 0.074 | 0.07 | 16.2 | 4.2 |
| LME | Southeast Australian Shelf | 8 | 9.2 | 14.6 | 0.088 | 0.09 | 60.2 | 4.7 |
| LME | Northeast U.S. Continental Shelf | 15 | 878.0 | 11.5 | 0.064 | 0.20 | 4096.5 | 6.5 |
| HSA | Indian Ocean | 4 | 452.4 | 15.3 | 0.066 | 0.09 | 4374.6 | 7.3 |
| LME | Baltic Sea | 5 | 828.7 | 7.5 | 0.088 | 0.20 | 11597.4 | 11.2 |
| LME | Labrador - Newfoundland | 8 | 292.0 | 4.5 | 0.017 | 0.33 | 2802.3 | 14.2 |

Table S5. RAM Legacy Database stocks used in analysis (TB = total biomass).

| Condition | \# of stocks |
| :--- | ---: |
| All RAMLDB stocks | 1058 |
| Not Pacific salmon stocks | 685 |
| Only stocks with TB/catch in metric tons | 350 |
| Only stocks with TB/catch time series $\geq 20$ years | 300 |
| Removed 23 stocks with strong SP/SR relationships | 277 |
| Removed 9 stocks without 20 years of data after trimming | 268 |
| Removed 5 stocks without SST data (e.g., Seto Sea not covered by COBE) | 263 |
| Removed 28 stocks preventing model convergence | 235 |

Table S6. Model symbols and their definitions.

| Type | Symbol | Definition |
| :--- | :--- | :--- |
| Data | $\mathrm{C}_{\mathrm{i}, \mathrm{t}}$ | Catch for stock $i$ in year $t$ |
| Data | $\mathrm{SP}_{\mathrm{i}, \mathrm{t}}$ | Surplus production for stock $i$ in year $t$ |
| Data | $\mathrm{Bi}_{\mathrm{i}, \mathrm{t}}$ | Total biomass for stock $i$ in year $t$ |
| Data | $\mathrm{SST}_{\mathrm{i}, \mathrm{t}}$ | Sea surface temperature (SST) experienced by stock $i$ in year $t$ |
| Data | $\mathrm{G}_{\mathrm{i}}$ | Group (taxonomic, geographic, or stock assessment model) for stock $i$ |
| Derived | $\varepsilon_{\mathrm{i}, \mathrm{t}}$ | Residual process variability for stock $i$ in year $t$ |
| Parameter | $\mathrm{r}_{\mathrm{i}}$ | Intrinsic rate of growth for stock $i$ |
| Parameter | $\mathrm{K}_{\mathrm{i}}$ | Carrying capacity for stock $i$ |
| Parameter | $\theta_{\mathrm{i}}$ | Influence of SST on productivity for stock $i$ |
| Parameter | $\mu_{\mathrm{SST}}$ | Mean of the distribution of SST influences $\left(\theta_{\mathrm{i}}\right)$ |
| Parameter | $\sigma_{S S T}$ | Standard deviation of the distribution of SST influences $\left(\theta_{\mathrm{i}}\right)$ |
| Parameter | $\mu_{\mathrm{G}, \mathrm{j}}$ | Mean of the distribution of SST influences $\left(\theta_{\mathrm{i}}\right)$ for group $j$ |
| Parameter | $\sigma_{\mathrm{G}}$ | Standard deviation of the group-specific distributions of SST influences $\left(\theta_{\mathrm{i}}\right)$ |
| Parameter | $\sigma_{p, i}$ | Standard deviation of the residual process variability for stock $i$ |
| Parameter | $\rho_{\mathrm{i}}$ | First-order (AR1) autocorrelation coefficient for stock $i$ |
| Constant | p | Shape parameter: fixed at 1.00, 0.55, 0.20, or 0.01 |
| Index | t | Year |
| Index | i | Stock |
| Index | j | Group (taxonomic or geographic) |

Table S7. Stock assessment methods represented in the data.

| Assessment model | Number | Countries |
| :--- | ---: | :--- |
| Biomass dynamics model (n=30) |  |  |
| BSPM: Bayesian surplus production model | 10 | Canada, Tuna-RFMO |
| ASPIC: Surplus production model | 6 | Tuna-RFMO, USA |
| Delay difference model | 5 | Canada, USA |
| ASPM: Age-structured surplus production model | 4 | South Africa |
| DPM: Dynamic production model | 2 | West Africa |
| qR: Surplus production model | 2 | Australia |
| LPM: Logistic production model | 1 | Canada |
| Integrated analysis (n=57) |  |  |
| SS3: Stock Synthesis v3.0 model | 26 | Australia, Europe, Tuna-RFMO, USA |
| SS2: Stock Synthesis v2.0 model | 22 | Australia, USA |
| SMS: Stochastic multi-species model | 3 | Europe |
| CASAL: C++ Algorithmic Stock Assessment Laboratory | 2 | New Zealand |
| IA: Integrated analysis | 1 | USA |
| JJM: Joint jack mackerel | 1 | Chile |
| SS1: Stock Synthesis v1.0 model | 1 | USA |
| SYM: Stochastic yield model | 1 | USA |
| Statistical catch-at-age model (n=55) | 20 | Europe, South Africa, USA |
| AD-CAM: AD-Model Builder statistical catch-at-age model | 8 | Canada, Europe, Tuna-RFMO, USA |
| SCA: Statistical catch-at-age model | 6 | USA |
| ASAP: Age Structured Assessment Program | 6 | USA |
| BAM: Beaufort assessment model | 5 | Europe |
| ICA: Integrated catch-at-age analysis | 4 | Canada, Europe |
| TSA: State-space catch-at-age time series analysis | 2 | Tuna-RFMO |
| MULTIFAN-CL: Length-based, age/spatially-structured model | 2 | Europe |
| SAM: State-space assessment model |  |  |
|  |  |  |
|  |  |  |


| CSA: Catch-survey analysis (like a state space approach) | 1 | USA |
| :---: | :---: | :---: |
| SCALE: A statistical catch-at-length model | 1 | USA |
| Statistical catch-at-length model ( $n=3$ ) |  |  |
| AD-CAL: AD-Model Builder catch-at-length model | 2 | USA |
| LBA: Length-based analysis | 1 | USA |
| Survey index ( $n=5$ ) |  |  |
| Temporal indices derived from scientific survey data | 3 | USA |
| SURBA: Survey-based stock assessment method | 2 | Canada |
| Unknown ( $n=38$ ) |  |  |
| Unknown | 27 | Canada, Chile, Europe, Peru, South Africa, USA |
| MSLM: Multi-stock length-based model | 7 | New Zealand |
| SnapEst: SnapEst age- and length-based model | 2 | Australia |
| CapTool: Spreadsheet assessment model used for capelin | 1 | Europe |
| RYM: Replacement yield model | 1 | South Africa |
| Virtual population analysis ( $n=47$ ) |  |  |
| XSA: Extended survivor analysis | 26 | Argentina, Canada, Europe, Tuna-RFMO |
| VPA: Virtual population analysis | 16 | Argentina, Canada, Europe, Japan, Russia |
| ADAPT: Adaptive framework-virtual population analysis | 1 | Europe |
| B-ADAPT: ADAPT approach with year effects in a catch multiplier | 1 | Europe |
| FLXSA: FLR variant of extended survivor analysis | 1 | Europe |
| NFT-ADAPT: VPA/ADPAT version 2.3.2 NOAA Fisheries | 1 | Europe |
| SPA-ADAPT: Sequential population analysis / ADAPT | 1 | Canada |

Table S8. AFS and FishBase guidelines for using life history traits to classify the resilience of fish stocks to exploitation and the $r$ priors used by catch-MSY for each resilience category.*

| Resilience | r prior | Von B <br> $\mathrm{K}(1 / \mathbf{y r})$ | Maximum <br> age $(\mathbf{y r})$ | Age at <br> maturity (yr) |
| :--- | :--- | :--- | :--- | :--- |
| High | $[0.6,1.5]$ | $>0.3$ | $1-3$ | $<1$ |
| Medium | $[0.2,1.0]$ | $0.16-0.30$ | $4-10$ | $2-4$ |
| Low | $[0.05,0.5]$ | $0.05-0.15$ | $11-30$ | $5-10$ |
| Very low | $[0.015,0.1]$ | $<0.05$ | $>30$ | $>10$ |
| Unknown | $[0.2,1.0]$ | ----- | ----- | ----- |

* Resilience categories were assigned using, in order of preference, resilience values: (1) reported on FB/SLB; (2) derived from the FishLife Von Bertalanffy growth parameter; (3) derived from the FB/SLB Von Bertalanffy growth parameter; (4) derived from the FB/SLB vulnerability metric; (5) derived from the FB/SLB Von Bertalanffy maximum age; (6) derived from the genus mode; or (7) derived from the family mode.


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