

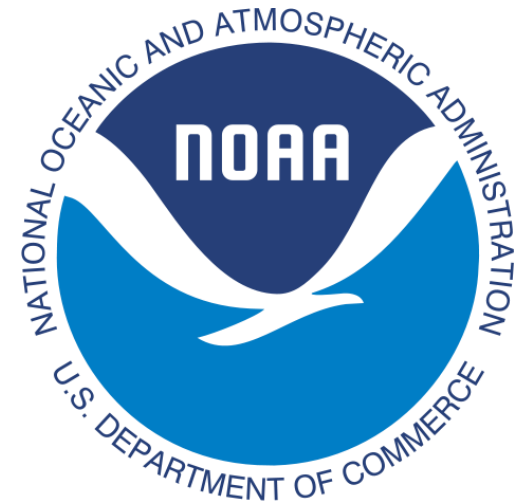
JABBA: Just Another Bayesian Biomass Assessment for Indian Ocean Blue shark

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agriculture,
forestry & fisheries

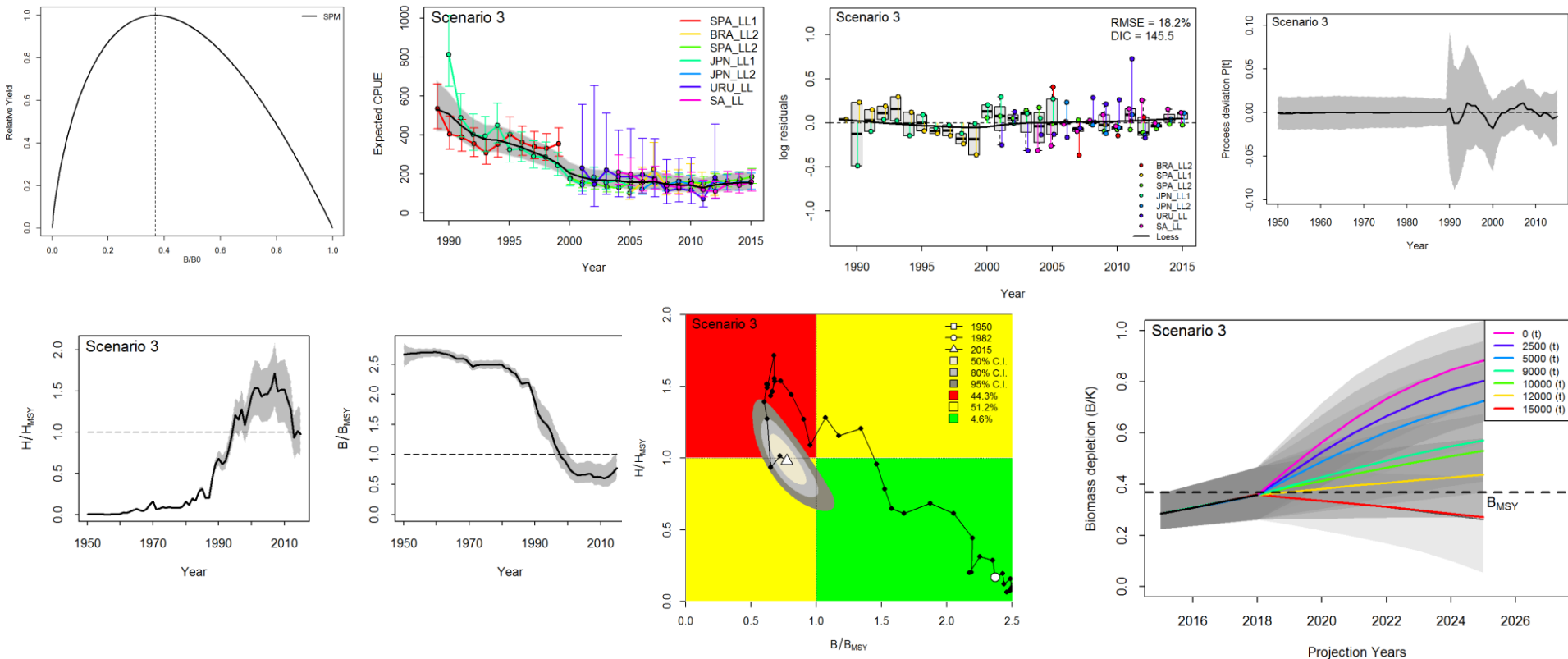
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JABBA R/JAGS interface

AIM: Improve estimation properties of Bayesian state-space surplus production models by building on previous formulations by Pella and Tomlinson (1969), Wang *et al.* (2014) and Fletcher (1978; c.f. Thorson *et al.* 2012) within a user-friendly R interface.



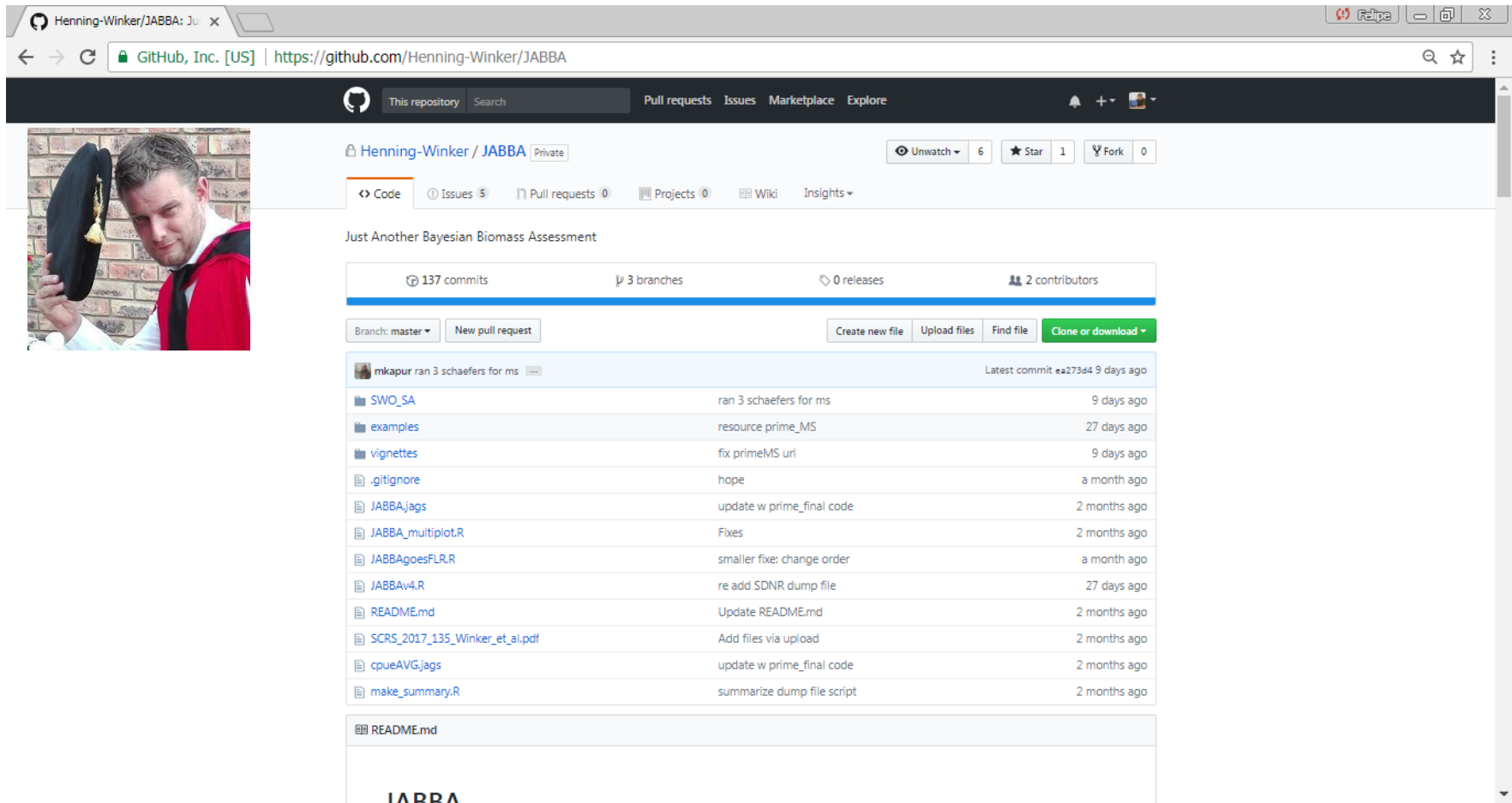
JABBA is a further development of the models applied in the 2015 ICCAT South Atlantic blue shark assessment and 2017, North Pacific blue shark assessment, the 2017 ICCAT Mediterranean Albacore assessment and the 2017 ICCAT North and South Atlantic shortfin mako shark assessments

JABBA (Just Another Bayesian Biomass Assessment): A generalized Bayesian State-Space Surplus Production Model

The inbuilt options include:

- Integrated state-space tool for averaging multiple CPUE series (+SE) for optional use in assessments
- Automatic fitting of multiple CPUE time series and associated standard errors
- Fox, Schaefer or Pella Tomlinson production function (optional as input B_{msy}/K)
- Flexible r prior specification: (1) range or (2) mean + CV of lognormal distribution
- Flexible initial depletion prior specification: (1) mean + CV lognormal or (2) mean + CV of beta distributions
- Kobe-type biplot plotting functions
- Improved Residual and MCMC diagnostics
- Optional estimation additional observation variance for individual or grouped CPUE time series
- Easy implementation of time-block changes in selectivity
- Forecasting of stock status under alternative TACs

JABBA (Just Another Bayesian Biomass Assessment): A generalized Bayesian State-Space Surplus Production Model



The screenshot shows the GitHub repository page for 'Henning-Winker / JABBA'. The repository is private and has 6 forks, 1 star, and 0 pull requests. The repository description is 'Just Another Bayesian Biomass Assessment'. The repository statistics show 137 commits, 3 branches, 0 releases, and 2 contributors. The repository is on the 'master' branch. The file list includes:

File	Commit Message	Time Ago
SWO_SA	ran 3 schaefers for ms	9 days ago
examples	resource prime_MS	27 days ago
vignettes	fix primeMS url	9 days ago
.gitignore	hope	a month ago
JABBA.jags	update w prime_final code	2 months ago
JABBA_multiplot.R	Fixes	2 months ago
JABBAgoesFLR.R	smaller fixer: change order	a month ago
JABBAv4.R	re add SDNR dump file	27 days ago
README.md	Update README.md	2 months ago
SCRS_2017_135_Winker_et_al.pdf	Add files via upload	2 months ago
cpueAVG.jags	update w prime_final code	2 months ago
make_summary.R	summarize dump file script	2 months ago

The repository also includes a README.md file. The repository is created by 'mkapur' and the latest commit is 'ea273d4' from 9 days ago.



JABBA Model Formulation

ADVANTAGE: Links surplus production models more directly to convention age-structured model formulations (e.g. SS3; Methot and Wetzel 2013).

Surplus production function of the generalized three parameter SPM by Pella and Tomlinson (1969)

$$SP_t = \frac{r}{m-1} B_{t-1} \left(1 - \left(\frac{B_{t-1}}{K} \right)^{m-1} \right) \quad (1)$$

where r is the intrinsic rate of population increase at time t , K is the unfished biomass and m is a shape parameter that determines at which B/K ratio maximum surplus production is attained.

- If $m = 2$, the model reduces to a Schaefer form, with the surplus production $g(B_t)$ attaining MSY at exactly $K/2$
- If $0 < m < 2$, $g(B_t)$ attains MSY at depletion levels smaller than $K/2$ and vice versa
- The Pella-Tomlinson model reduces to a Fox model if m approaches one ($m=1$) resulting in maximum surplus production at $\sim 0.37K$

JABBA Model Formulation

B_{msy} is given by:

$$B_{MSY} = Km^{\frac{-1}{m-1}} \quad (2)$$

and the corresponding harvest rate at MSY (H_{MSY}) is:

$$H_{MSY} = \frac{r}{m-1} \left(1 - \frac{1}{m} \right) \quad (3)$$

where the harvest rate H is defined here as the ratio of:

$$H = \frac{C}{B} \quad (4)$$

where C denotes the catch. Correspondingly H_{MSY} can be expressed by:

$$H_{MSY} = \frac{MSY}{B_{MSY}} \quad (5)$$

JABBA Model Formulation

Combing and re-arranging equation (3) and (5), it follows that r in equation (1) can be expressed as:

$$H_{MSY} = \frac{r}{m-1} \left(1 - \frac{1}{m} \right) \longleftrightarrow H_{MSY} = \frac{MSY}{B_{MSY}}$$

$$(6) \quad r = \frac{MSY}{B_{MSY}} \frac{m-1}{1-m^{-1}} \quad \text{or} \quad r = H_{MSY} \frac{m-1}{1-m^{-1}} \quad (7)$$

This allows re-formulating the production function of the Pella-Tomlinson equation as a function of H_{MSY} , such that:

$$SP_t = \frac{H_{MSY}}{(1-m^{-1})} B_{t-1} \left(1 - \left(\frac{B_{t-1}}{K} \right)^{m-1} \right) \quad (8)$$

where m can be directly translated into B_{MSY}/K and thus determines the biomass depletion level where MSY is achieved (Thorson *et al.* 2012a), using the following relationship:

$$\frac{B_{MSY}}{K} = m \left(-\frac{1}{m-1} \right) \quad (9)$$

JABBA Model Formulation

Because prior formulations for most SPM-based assessments are specified for r , we provide the following equation to easily convert r estimates (or prior means) into H_{MSY} for any given shape parameter input m :

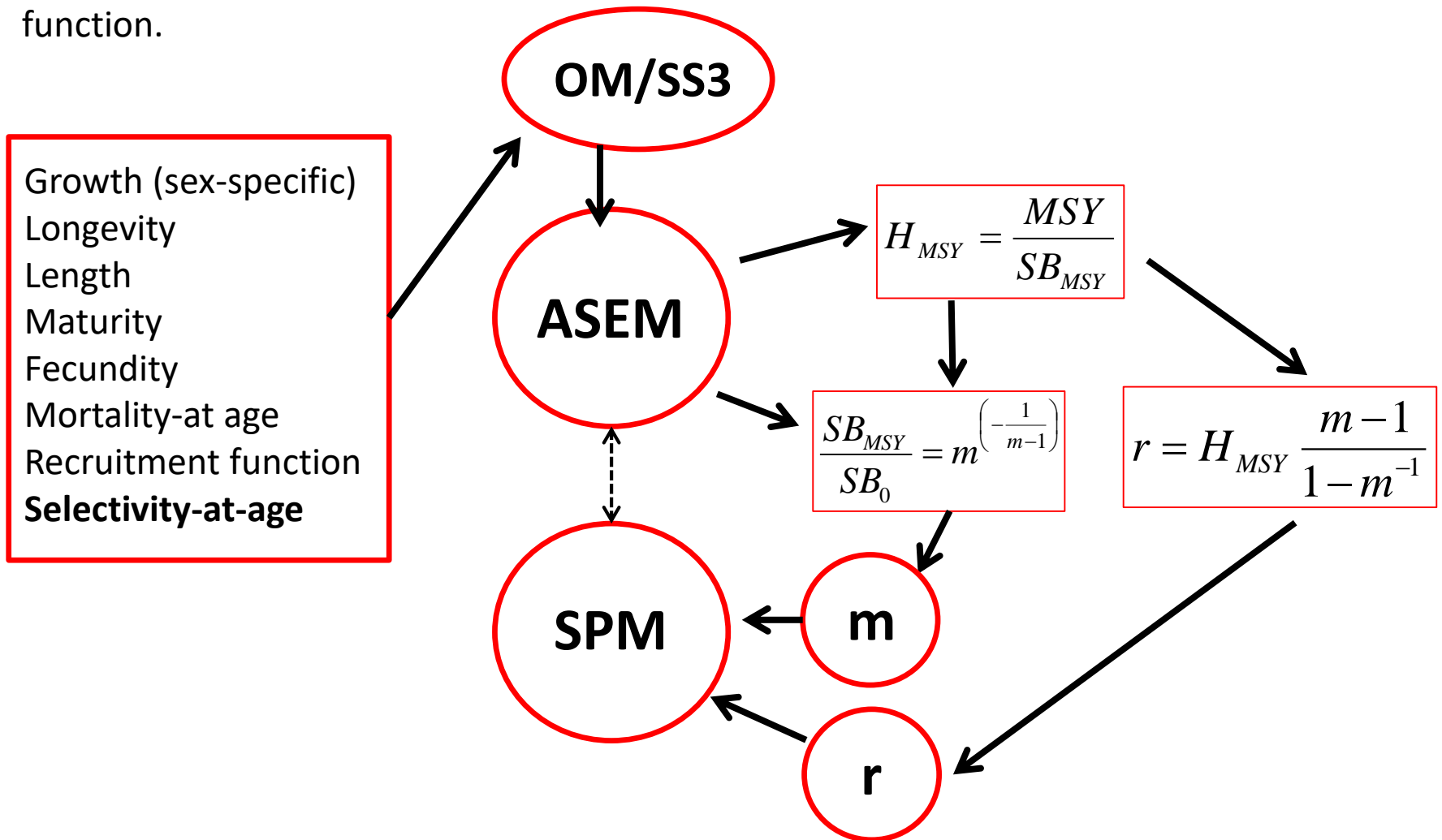
$$H_{MSY} = r \frac{(m - 1)}{(1 - m^{-1})} \quad (10)$$

Equations (5) - (10) illustrate the direct link between the Pella-Tomlinson SPM and the age-structured, which emphasizes the potential for deriving informative priors for r and m from spawning biomass- and yield-per-recruit analysis with integrated spawning recruitment relationships by generating deviates of $H_{MSY} = MSY/B_{MSY}$ and B_{MSY}/K , respectively (Maunder 2003; Thorson *et al.* 2012a; Wang *et al.* 2014).

DETOUR: Linking age-structure and surplus production models:

Potential MSE (MP) and Cross-validation application

Equations (5) - (7) illustrate the direct link between the Pella-Tomlinson SPM and an age-structured model, which can be used to translate SS3 estimates of the ratios of $MSY/BMSY$ and B_{MSY}/SB_0 and MSY/B_{MSY} into values of H_{MSY} and the shape parameter m of the production function.



JABBA Process-Equation

Bayesian State-Space formulation

We formulated the JABBA process equation by building on the Bayesian state-space estimation framework proposed by Meyer and Millar (1999). The biomass B_y in year y is expressed as proportion of K (i.e. $P_y = B_y / K$) to improve the efficiency of the estimation algorithm.

The model is formulated to accommodate multiple CPUE for fisheries f . The initial biomass in the first year of the time series was scaled by introducing model parameter φ to estimate the ratio of the spawning biomass in the first year to K (Carvalho *et al.* 2014). The stochastic form of the process equation is given by:

$$P_y = \begin{cases} \varphi e^{\eta_y} & y = 1 \\ \left(P_{y-1} + \frac{r}{(m-1)} P_{y-1} (1 - P_{y-1}^m) - \frac{\sum_f C_{f,y-1}}{K} \right) e^{\eta_y} & y = 2, 3, \dots, n \end{cases} \quad (11)$$

Where η_y is the process error, with $\eta_y \sim N(0, \sigma_\eta^2)$, $C_{f,y-1}$ is the catch in year y by fishery f

JABBA Observation Equation

The corresponding biomass for year y is:

$$B_y = P_y K \quad (12)$$

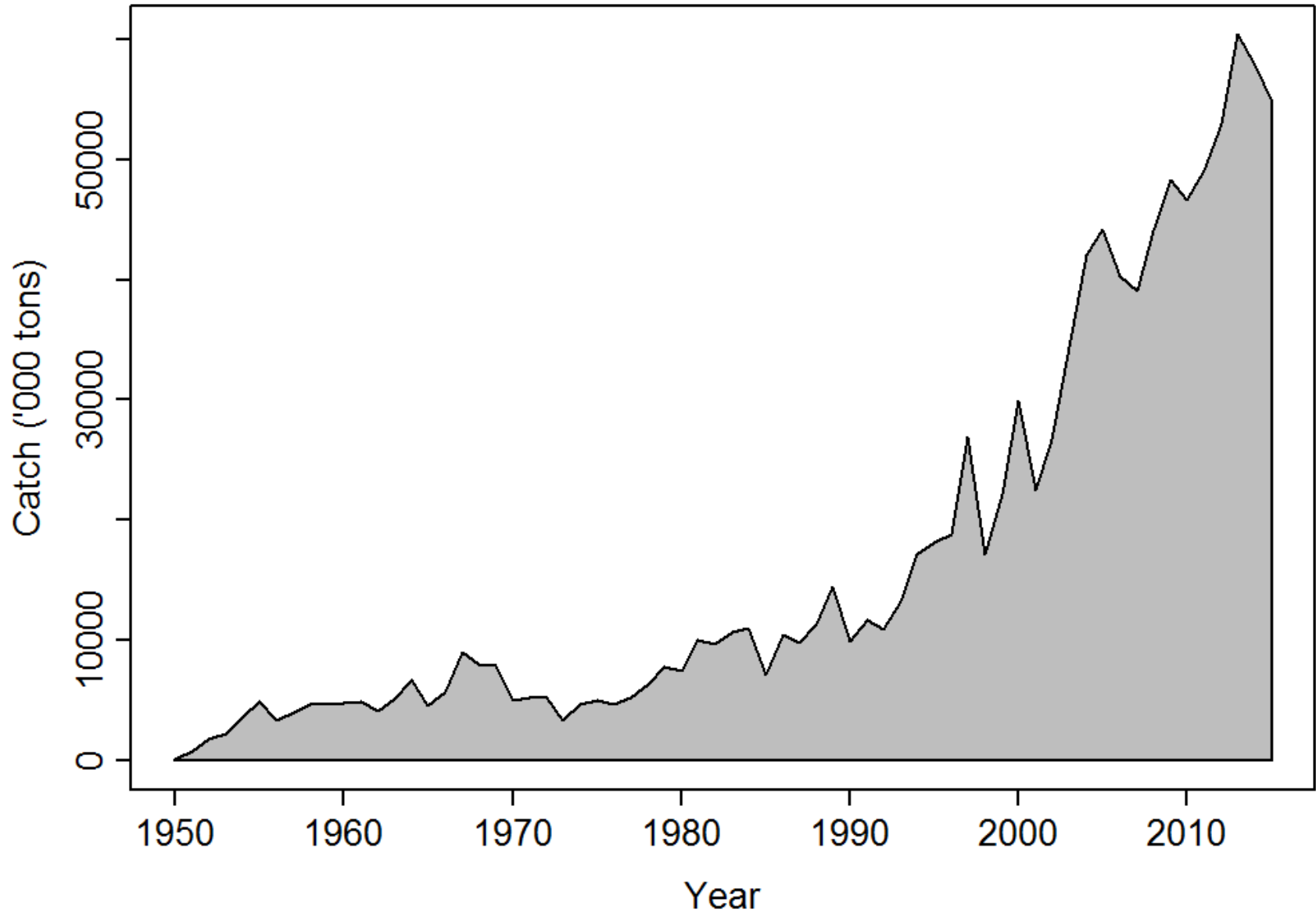
The observation equation is given by:

$$I_{f,y} = q_f B_{f,y} e^{\varepsilon_y} \quad y = 1, 2, \dots, n. \quad (13)$$

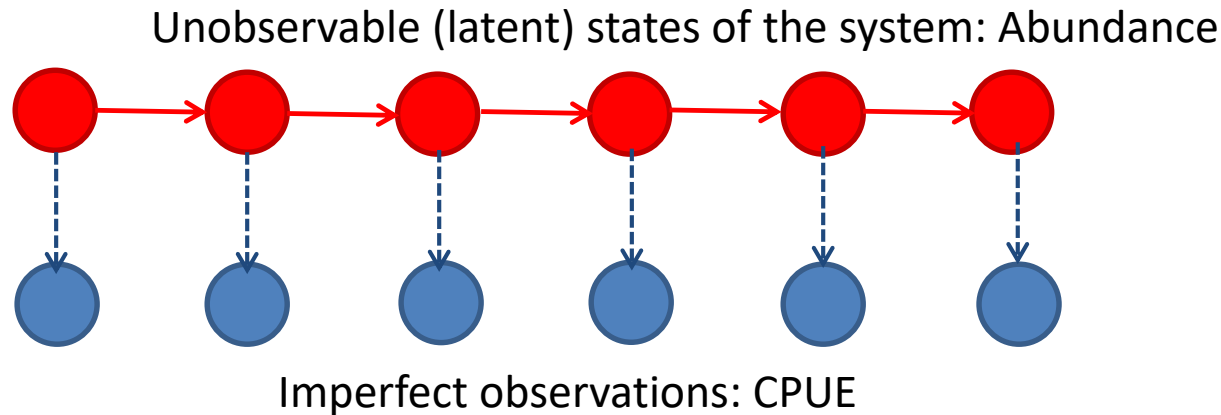
where, q_f is the estimable catchability coefficient associated with the abundance index for fishery f and ε_y is the observation error, with

$$\sigma_{\varepsilon,y,f}^2 = \hat{\sigma}_{SE,y,f}^2 + \sigma_{Add,f}^2 \quad \varepsilon_{y,f} \sim N(0, \sigma_{\varepsilon,y,f}^2) \quad (14)$$

Indian Ocean Blue shark (GAM)



Bayesian State-Space CPUE averaging tool



Process, State or Evolutionary equation

$$z_t = z_{t-1} \exp(r + \tau_t)$$

State variable

Underlying trend

Process error

$$\tau_t \sim N(0, \sigma_\tau^2)$$

Process variance

Observation equation

$$y_{y,i=1} = q_i z_i \exp(\varepsilon_{t,i})$$

Observation

Scaling coefficient

Observation error

$$\varepsilon_t \sim N(0, \sigma_\varepsilon^2)$$

Observation variance

JABBA runs

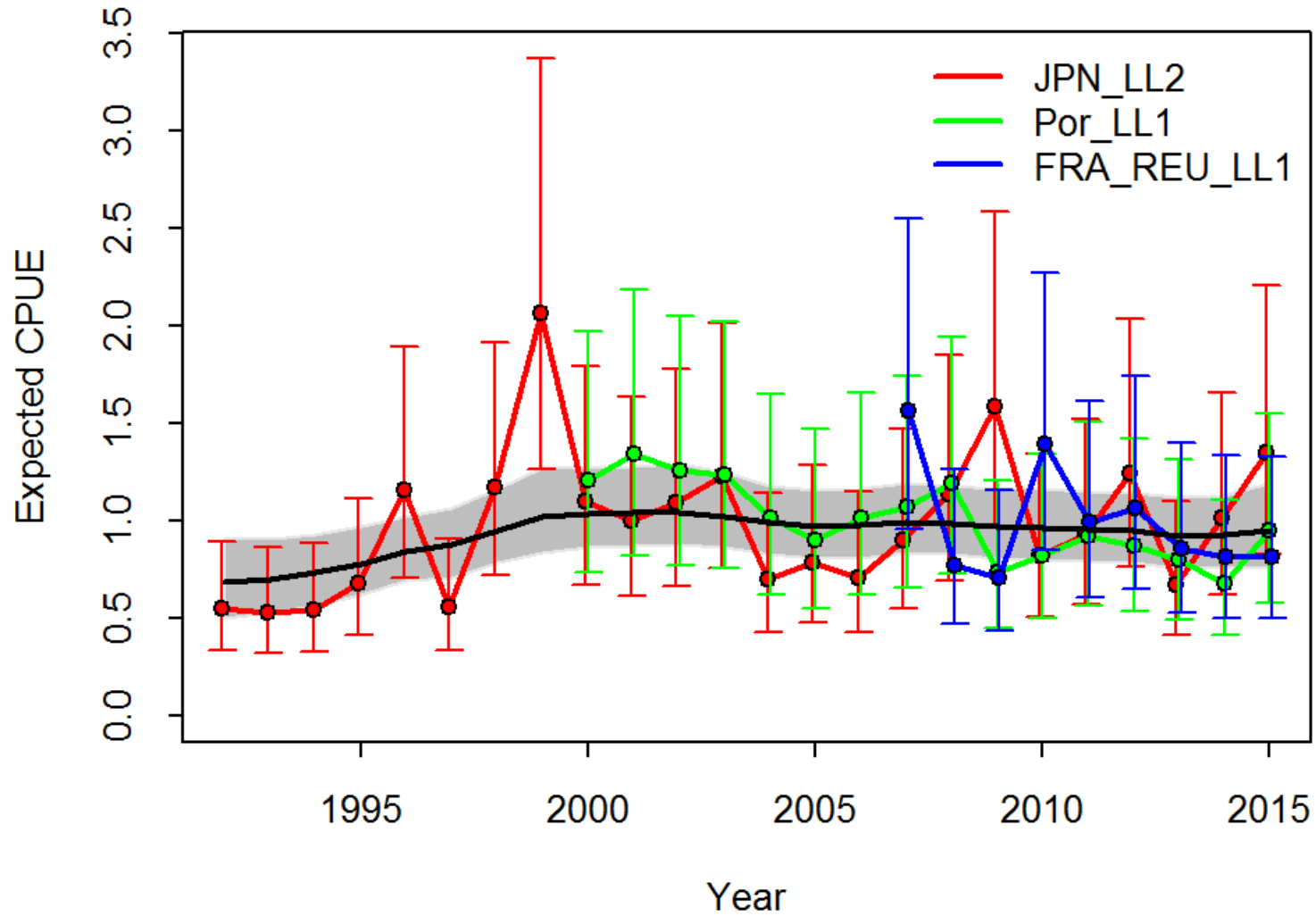
Prior formulation: Summary

Parameter	Distribution	min/ μ	max/CV
K	lognormal	600000	2
r	lognormal	0.267	0.07
$\varphi = B_{1950}/K$	lognormal	1	0.25
σ_{η}^2	inverse-gamma	4	0.01
$\sigma_{ADD,i}^2$	inverse-gamma	0.001	0.001
m	fixed	2	(Schaefer)
q	uniform	0.000000001	1

Minimum observation error CV was fixed at 0.25, so that: $\sigma_{\varepsilon,y,f}^2 = 0.25_{Add,f}^2 + \sigma_{Add,f}^2$

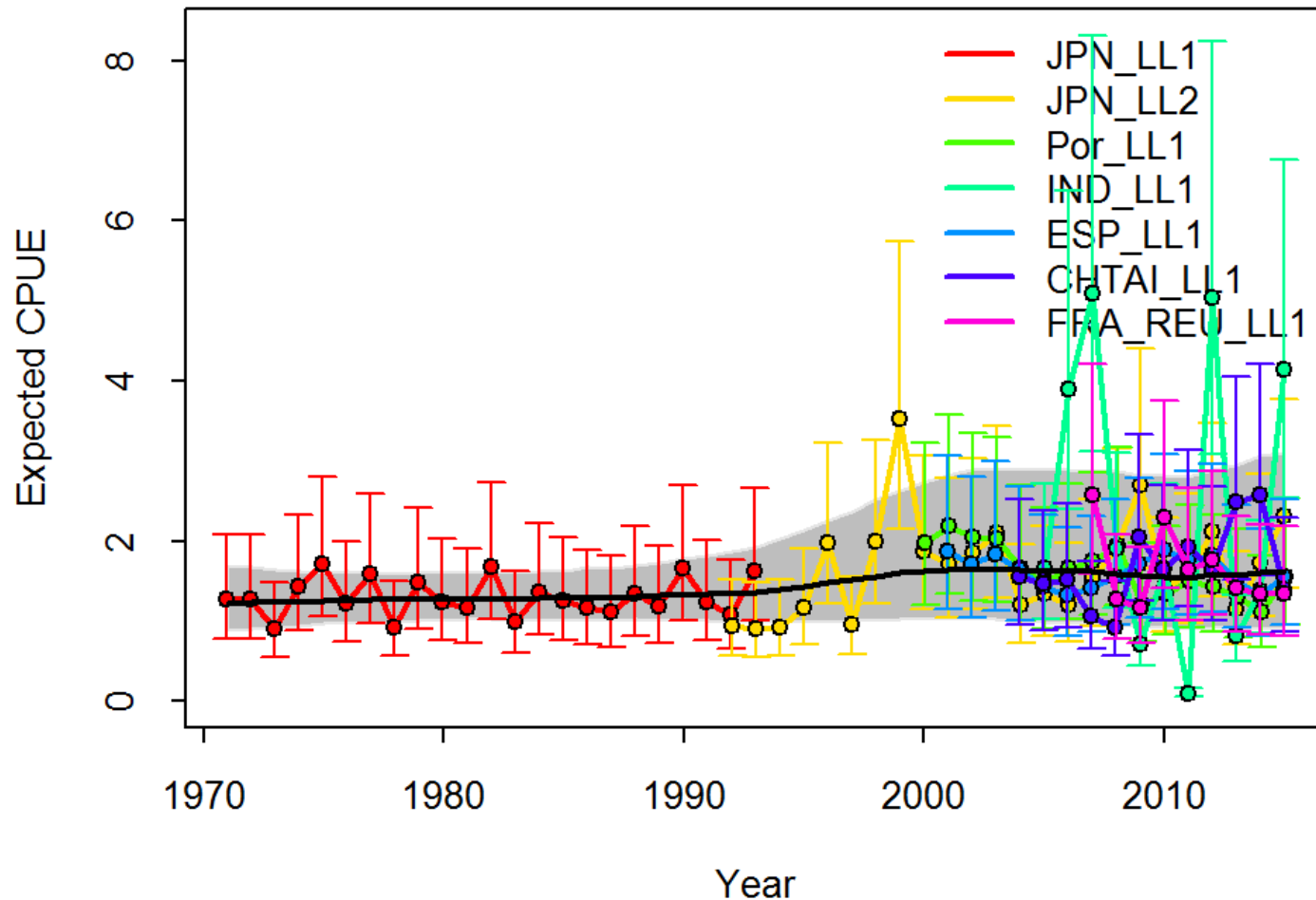
Base-case model

Japan late + EU-Portugal + EU-Spain + EU-France



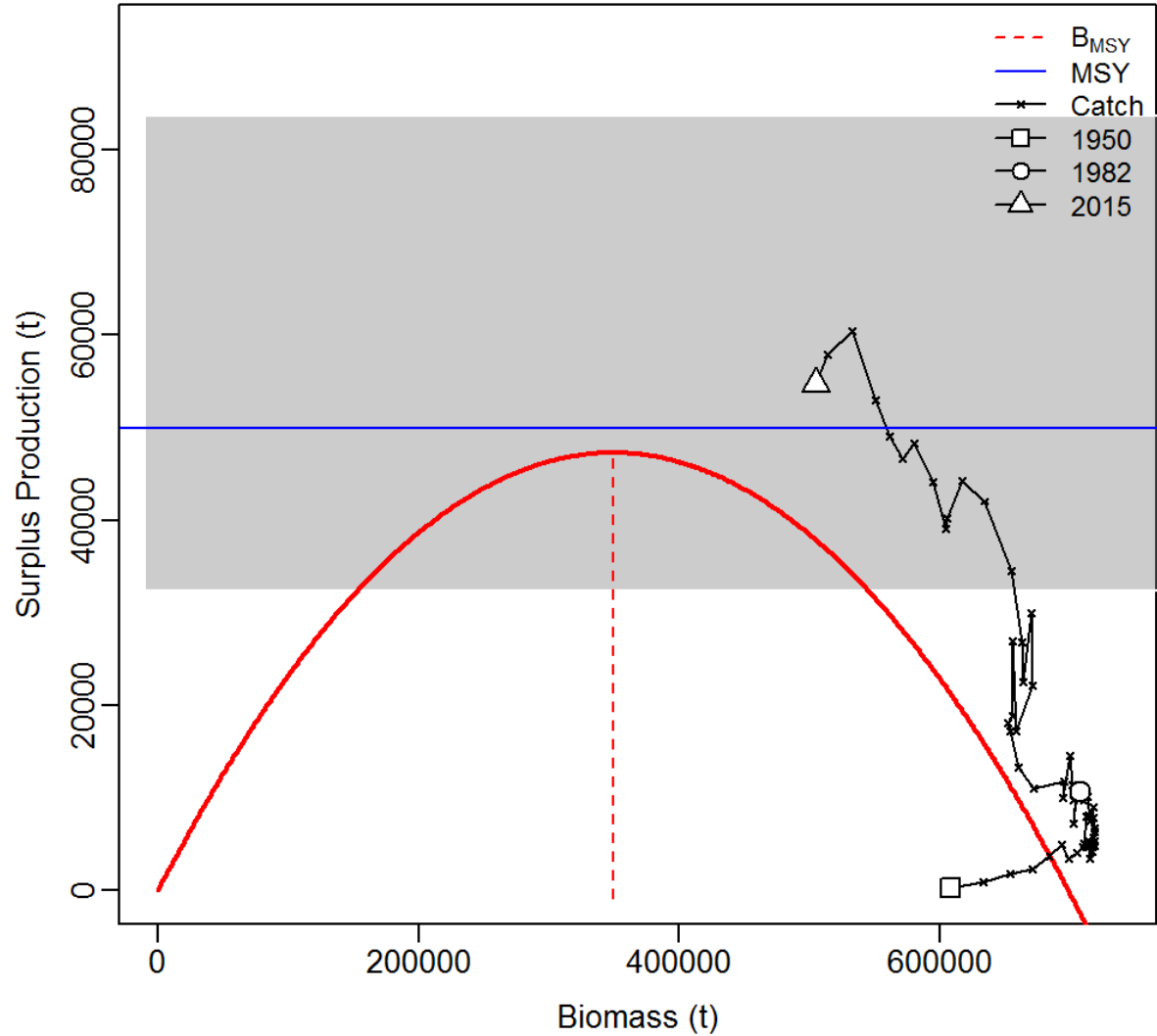
Alternative model

During the 2017 IOTC blue shark assessment, the formulation of alternative scenarios specifically focused on identifying and improving poor fits to CPUE series that may arise from fitting of multiple standardized CPUE time series with conflicting trends.



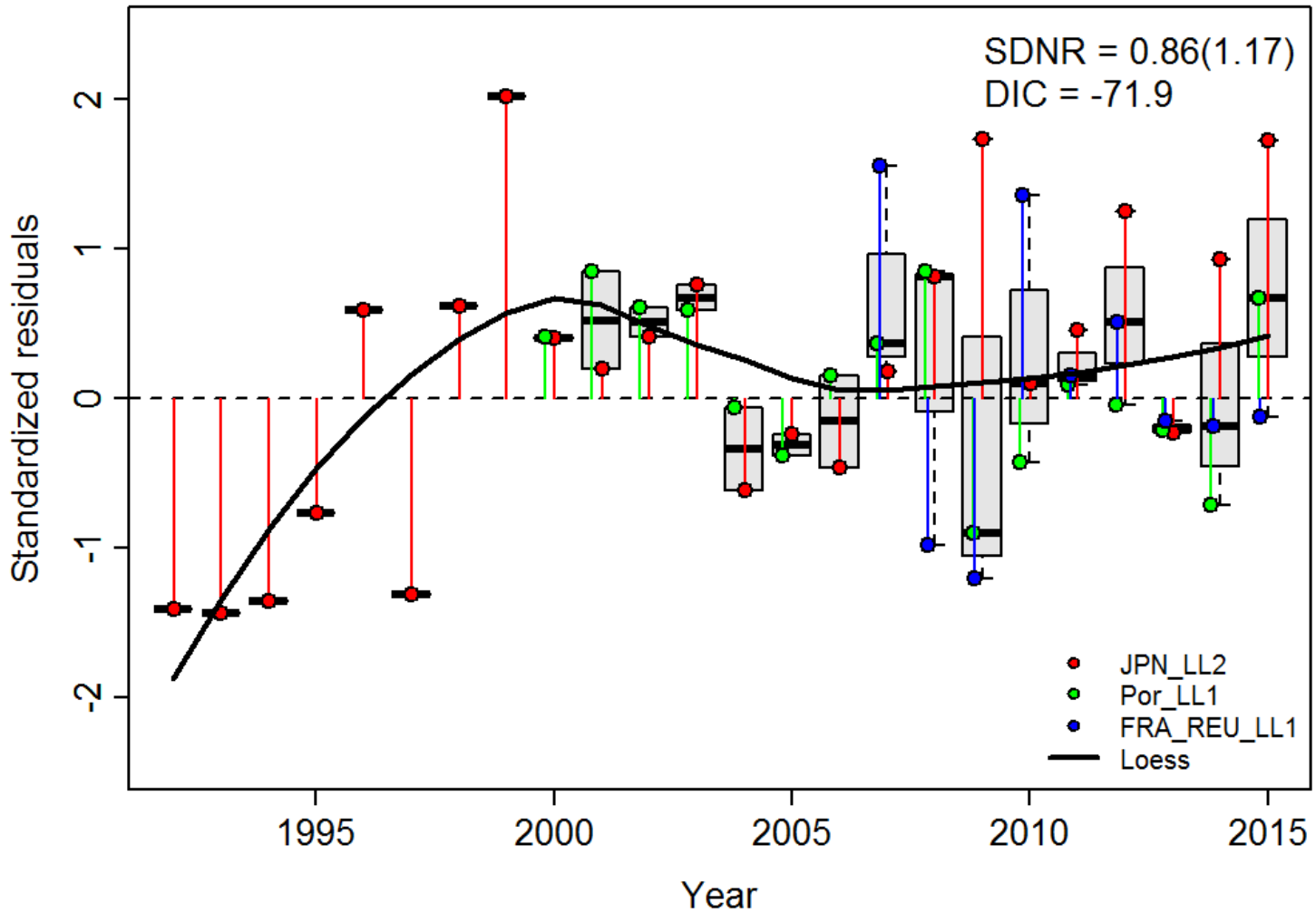
Schaefer Production Function

Base-case



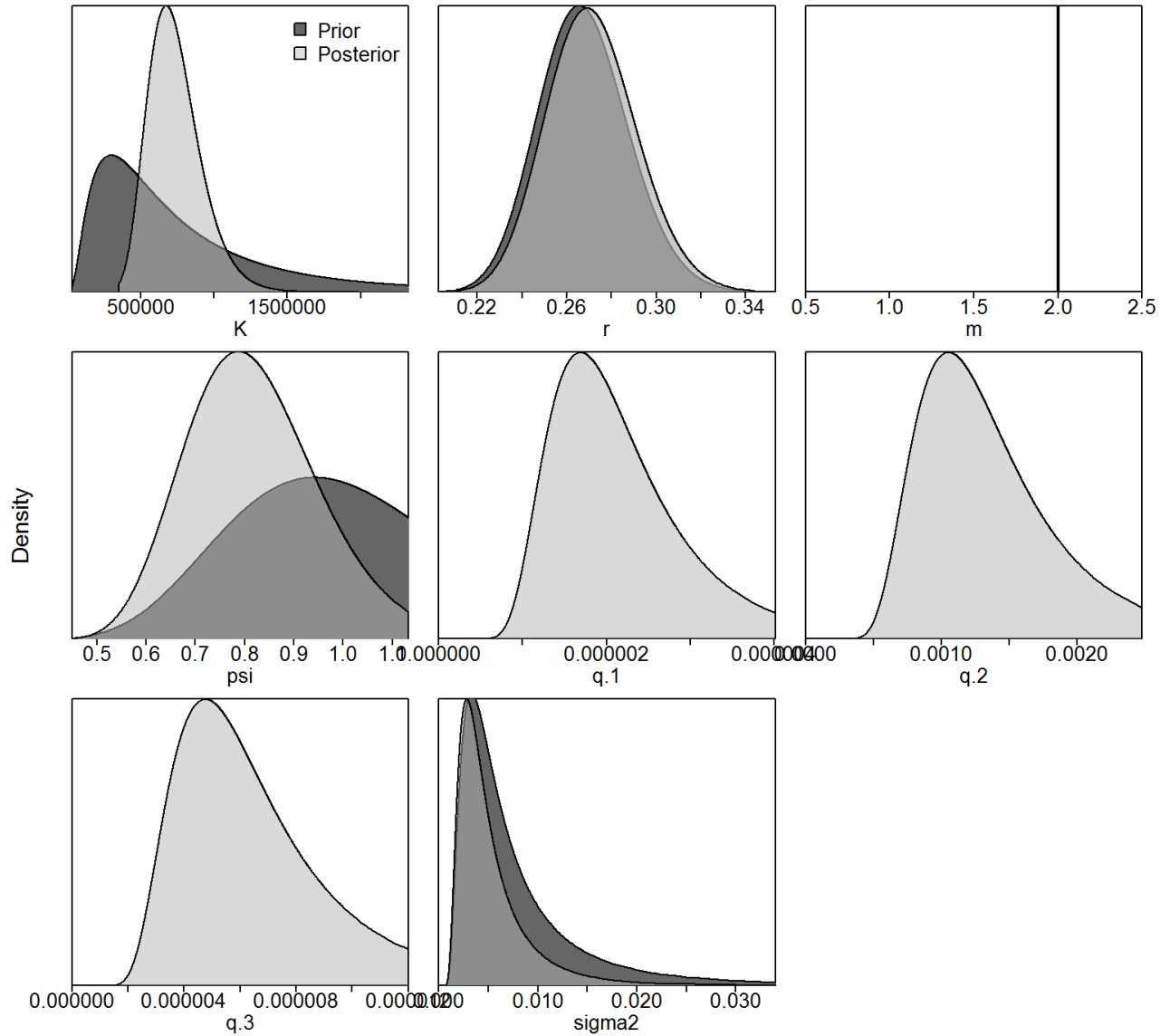
CPUE fits

Base-case



Prior vs. Posterior

Base-case

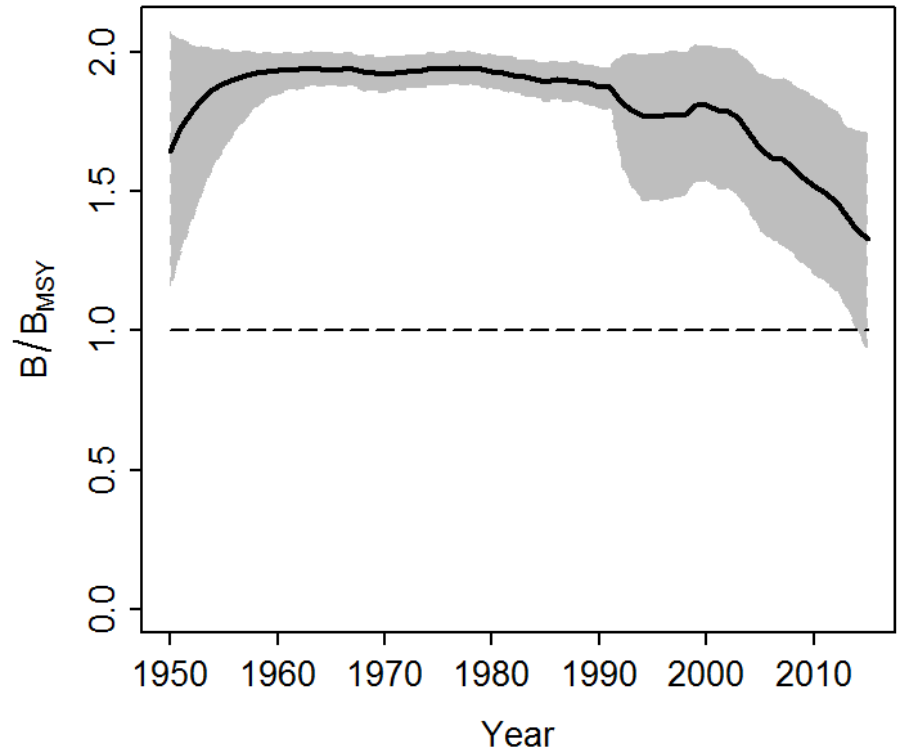
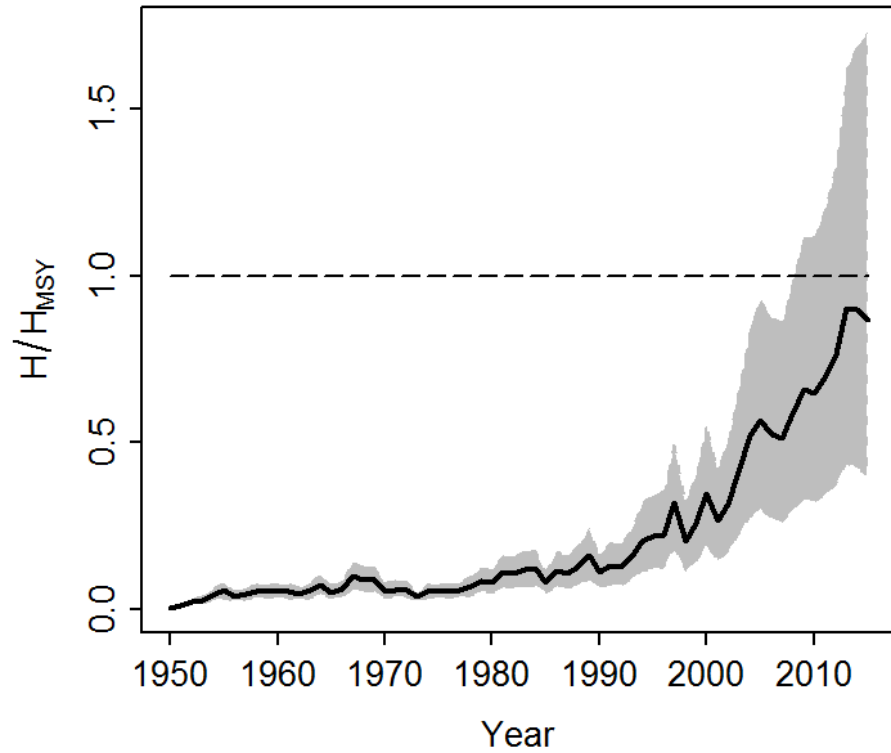


Model Estimates (Base-case)

	Median	2.50%	97.50%
K	698486	476589.9	1233646
r	0.271	0.235	0.313
ψ (psi)	0.824	0.57	1.042
σ	0.045	0.032	0.084
H_{msy}	0.135	0.117	0.156
SB_{msy}	349243	238295	616823
MSY	47355.78	32333.56	83741.82
P₁₉₅₀	0.823	0.57	1.041
P₂₀₁₅	0.666	0.46	0.861
B₂₀₁₅/B_{msy}	1.333	0.92	1.722
H₂₀₁₅/H_{msy}	0.869	0.396	1.738

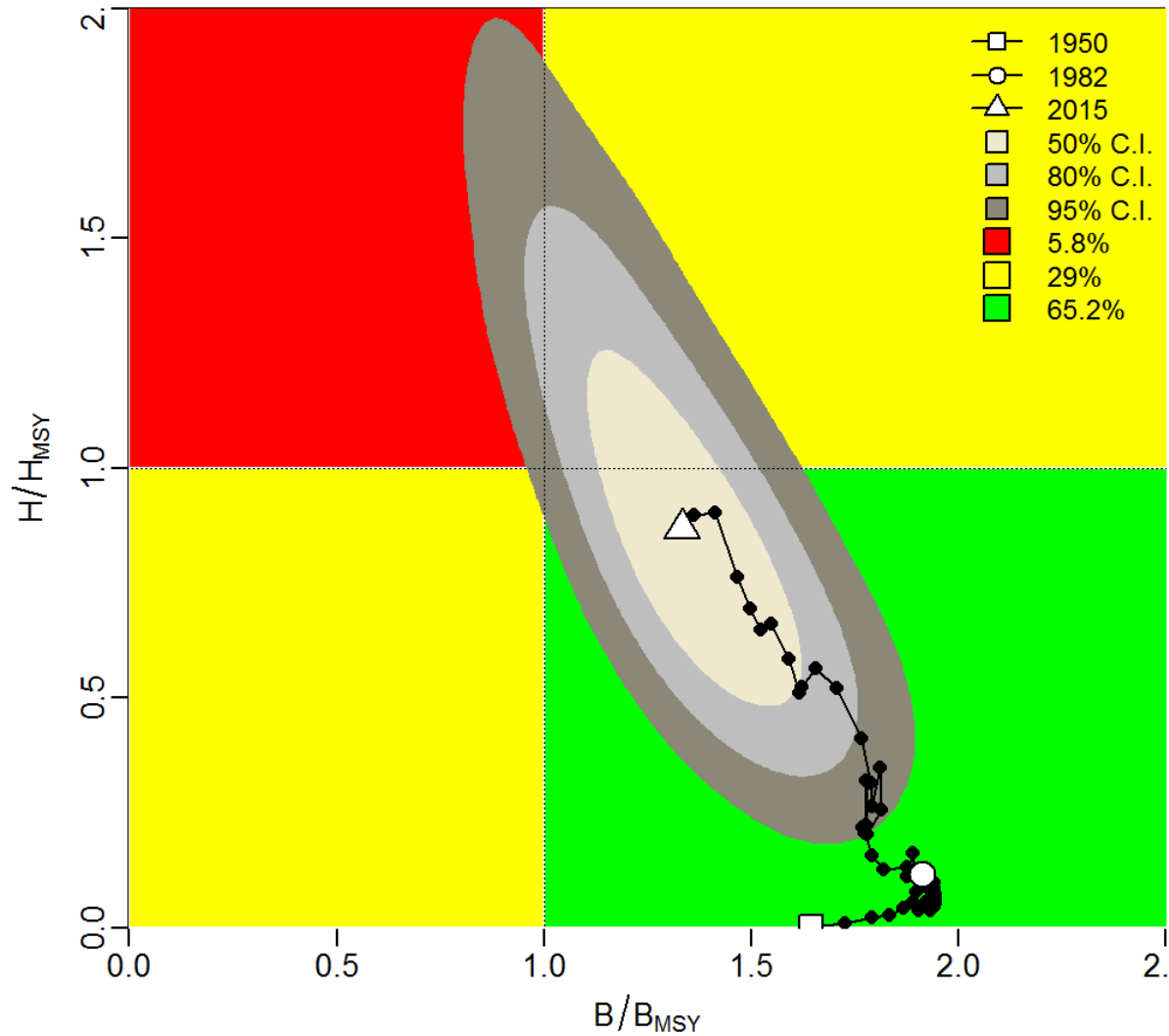
Time series of Reference Points

Base-case

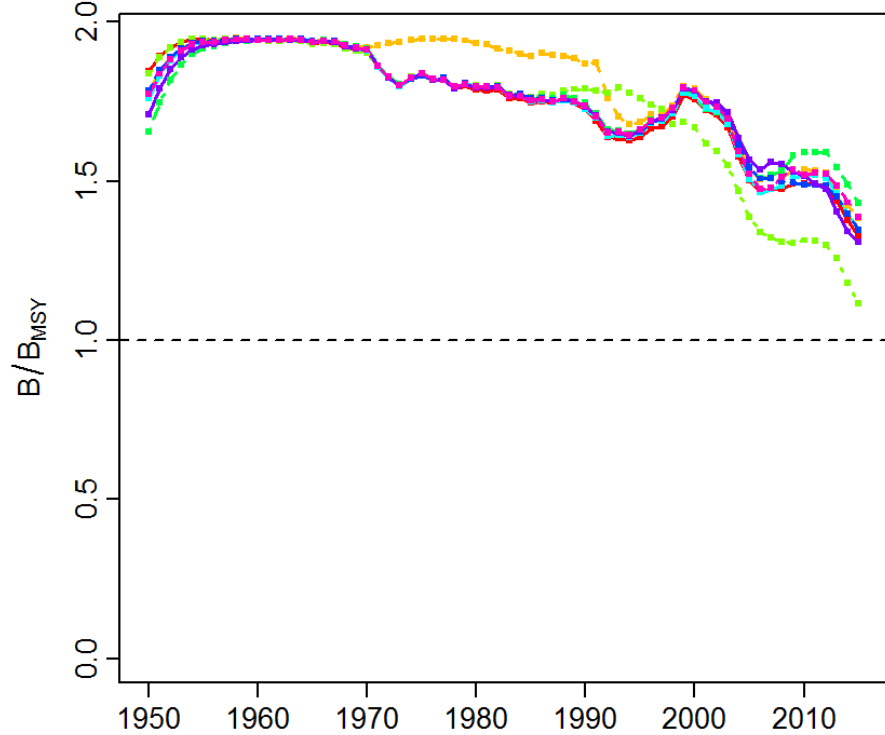
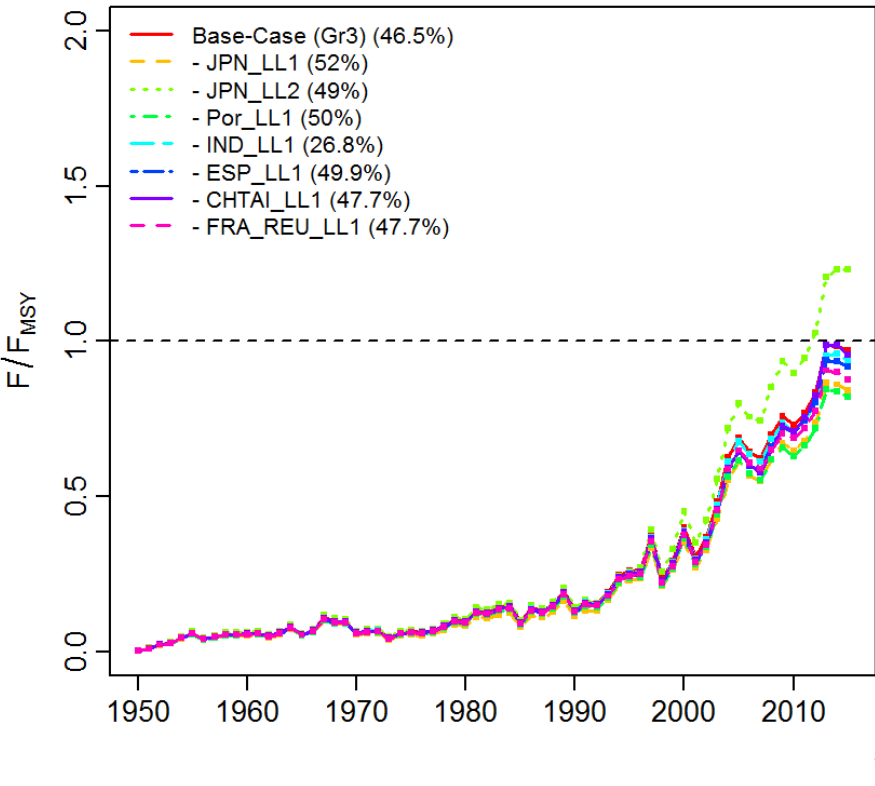


Kobe plot

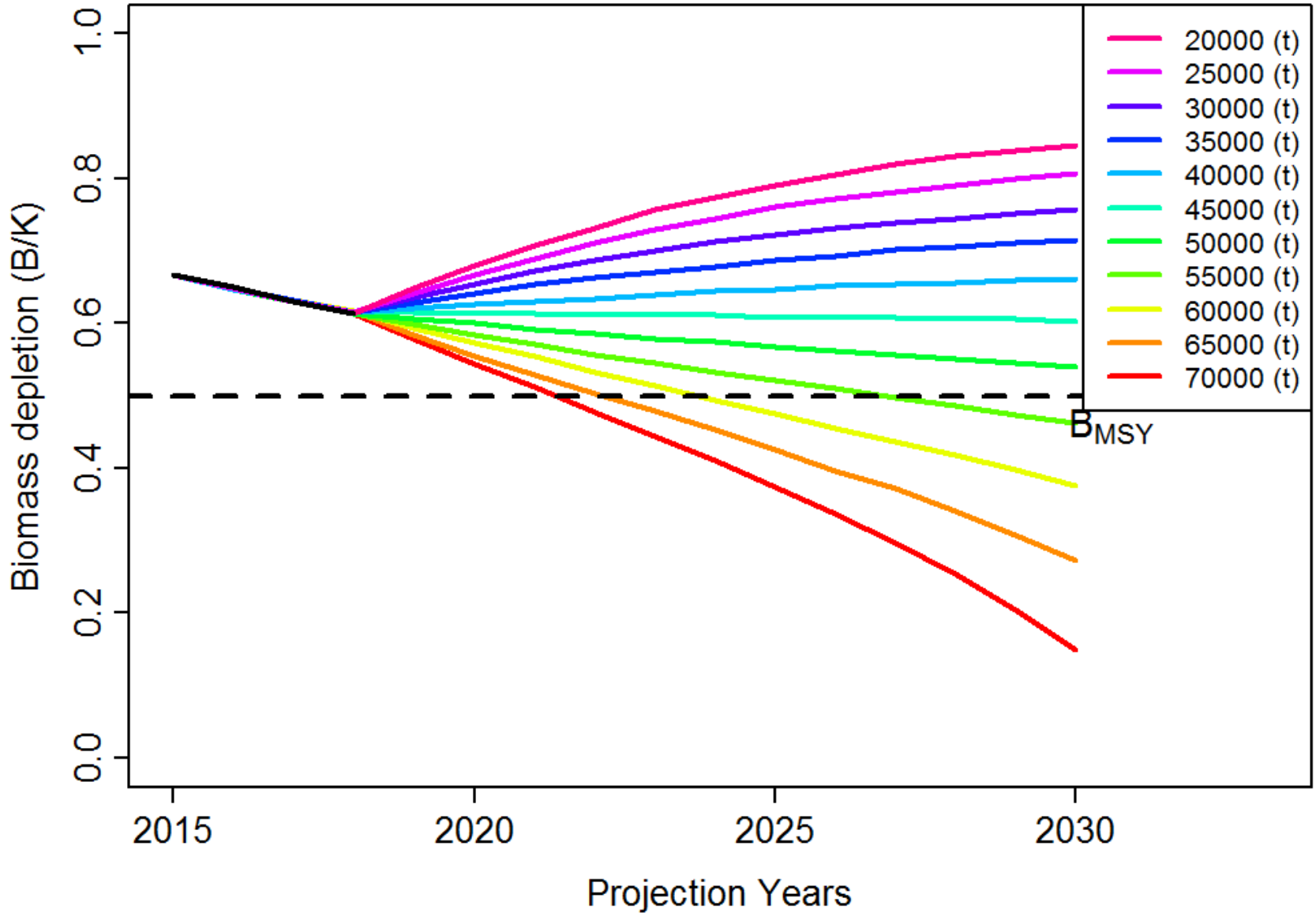
Base-case



Sensitivity runs (drop one from Base-case)



Projections (Base-case)



THANK YOU!