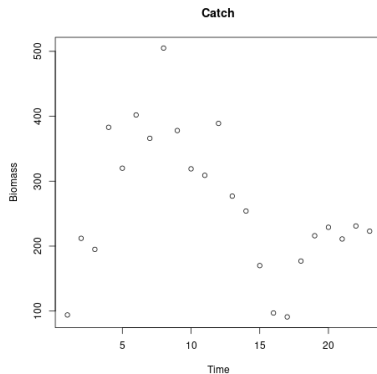
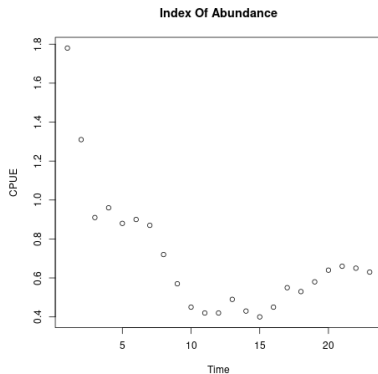


Metamodeling for Bias Estimation of Biological Reference Points Under Two-Parameter SRRs

Nick Grunloh

14 March 2022





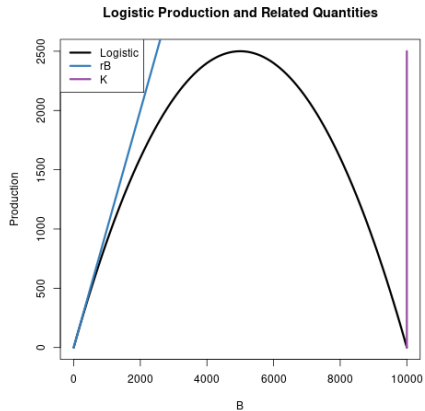
$$I_t = qB_t e^{\epsilon} \quad \epsilon \sim N(0, \sigma^2)$$

$$\frac{dB(t)}{dt} = P(B(t); \theta) - C(t)$$

Schaefer Model

$$P_{\theta}(B) = rB \left(1 - \frac{B}{K}\right)$$

$$\theta = (r, K)$$

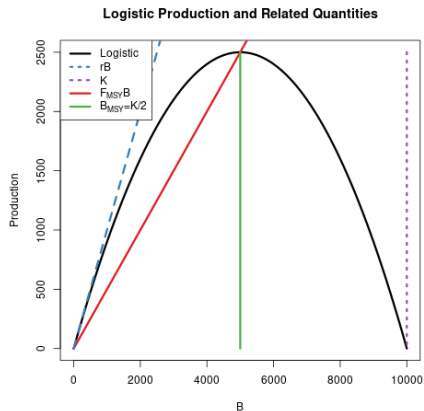


Schaefer Reference Points

$$F^* = \frac{r}{2}$$

$$\frac{B^*}{B_0} = \frac{1}{2}$$

$$MSY = \frac{rK}{4}$$



Conceptually:

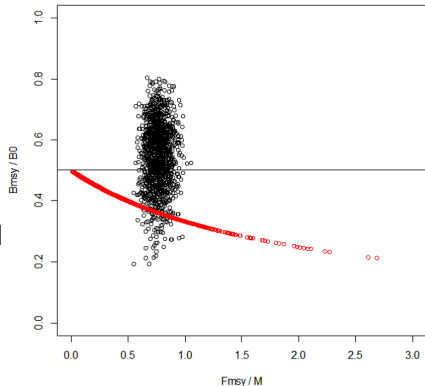
$$F^* \in \mathbb{R}^+ \quad \frac{B^*}{\bar{B}(0)} \in (0, 1)$$

Mangel et al. 2013, CJFAS:

- BH Model:

$$F^* \in \mathbb{R}^+ \quad \frac{B^*}{\bar{B}(0)} = \frac{1}{F^*/M+2}$$

- Similar Constraint for Ricker and other 2 Parameter Curves



Conceptually:

$$F^* \in \mathbb{R}^+ \quad \frac{B^*}{\bar{B}(0)} \in (0, 1)$$

Mangel et al. 2013, CJFAS:

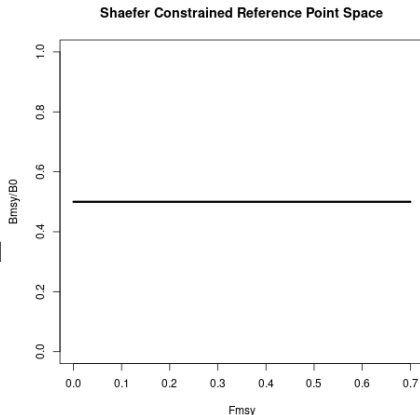
- BH Model:

$$F^* \in \mathbb{R}^+ \quad \frac{B^*}{\bar{B}(0)} = \frac{1}{F^*/M+2}$$

- Similar Constraint for Ricker and other 2 Parameter Curves

Schaefer Model:

$$F^* \in \mathbb{R}^+ \quad \frac{B^*}{\bar{B}(0)} = \frac{1}{2}$$



Pella-Tomlinson Production Model

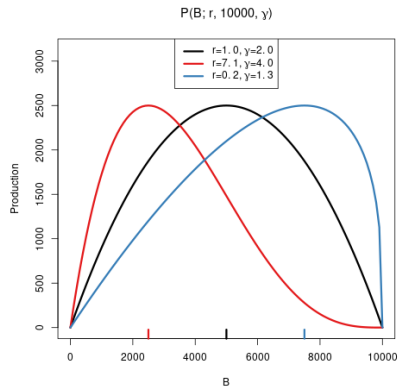
$$I(t) \sim LN(qB(t), \sigma^2)$$

$$\frac{dB(t)}{dt} = P_{\theta}(B(t)) - F(t)B(t)$$

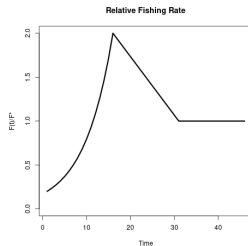
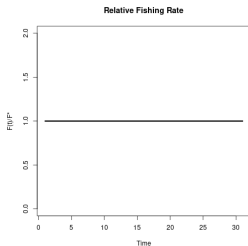
$$P_{\theta}(B) = \frac{rB}{\gamma - 1} \left(1 - \frac{B}{K}\right)^{\gamma-1}$$

$$\theta = (r, K, \gamma)$$

$$\gamma = 2 \Rightarrow \text{Schaefer Model}$$

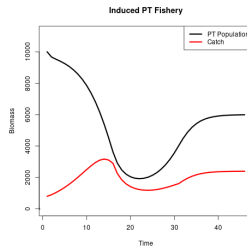
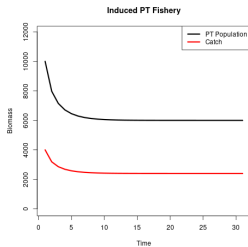
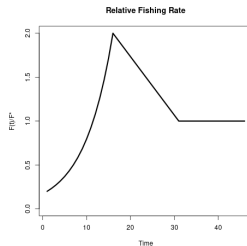
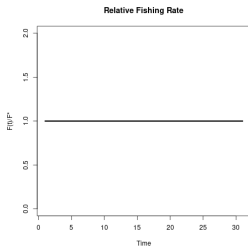


Catch

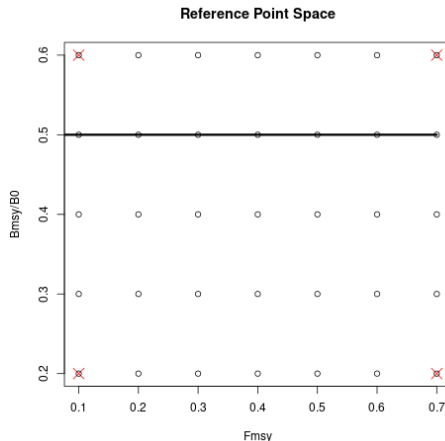


$$\begin{aligned} C(t) &= F(t)B(t) \\ &= F^* \left(\frac{F(t)}{F^*} \right) B(t) \end{aligned}$$

Catch

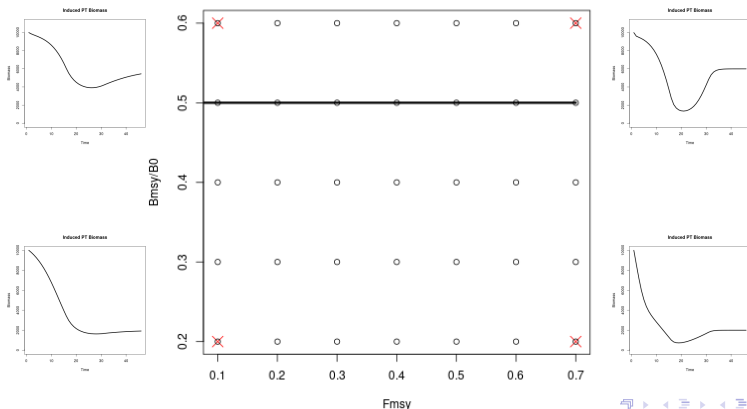


$$\theta = \left[r = F^* \left(\frac{1 - \frac{B^*}{\bar{B}(0)}}{\frac{B^*}{\bar{B}(0)}} \right) \left(1 - \frac{B^*}{\bar{B}(0)} \right)^{\left(\frac{\frac{B^*}{\bar{B}(0)} - 1}{\frac{B^*}{\bar{B}(0)}} \right)}, K = 10000, \gamma = \frac{1}{\frac{B^*}{\bar{B}(0)}} \right]$$



$$\theta = \left[r = F^* \left(\frac{1 - \frac{B^*}{\bar{B}(0)}}{\frac{B^*}{\bar{B}(0)}} \right) \left(1 - \frac{B^*}{\bar{B}(0)} \right)^{\left(\frac{\frac{B^*}{\bar{B}(0)} - 1}{\frac{B^*}{\bar{B}(0)}} \right)}, K = 10000, \gamma = \frac{1}{\frac{B^*}{\bar{B}(0)}} \right]$$

Reference Point Space



Metamodel

Squared Exponential Kernel to extend estimates across RP space

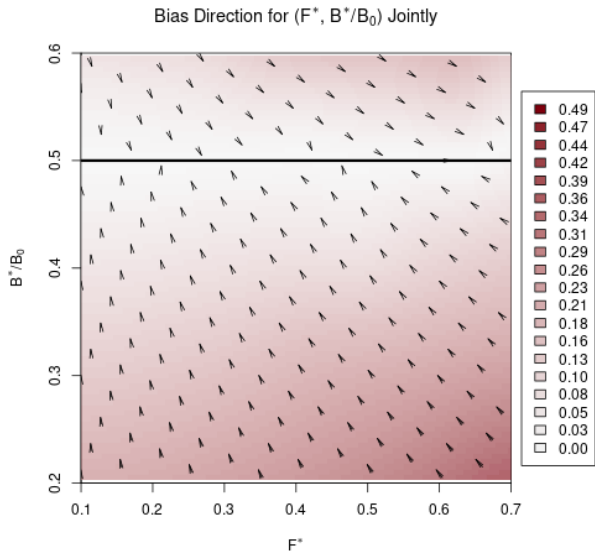
Let $\check{\cdot}$ decorate the extended estimates.

The true RPs are known and the metamodel extends simulation grid estimates across the RP-Space.

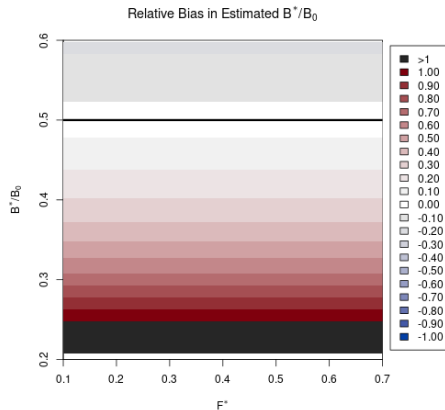
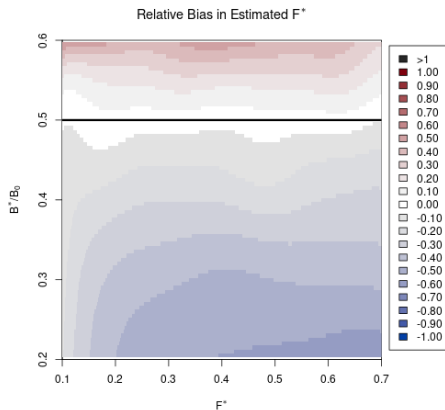
$$\check{B}^* = \frac{\check{K}}{2} \quad \check{F}^* = \frac{\check{r}}{2}.$$

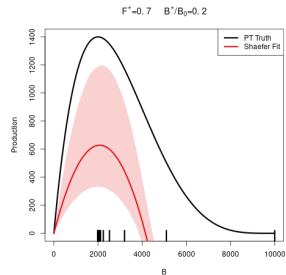
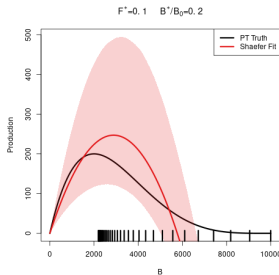
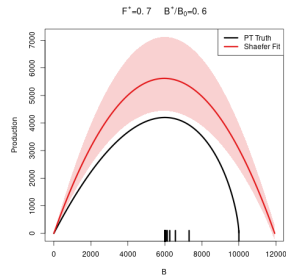
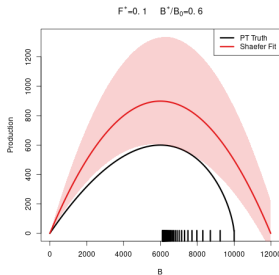
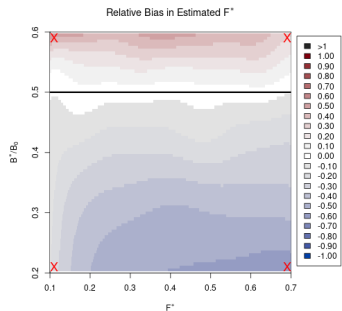
We'll look at bias results in the terms of percent error.

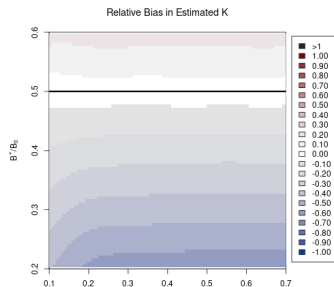
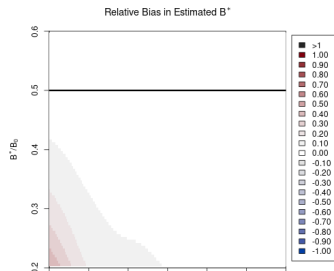
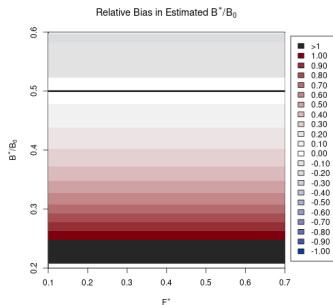
$$\text{Relative Bias} = \frac{\check{RP} - RP}{RP}$$



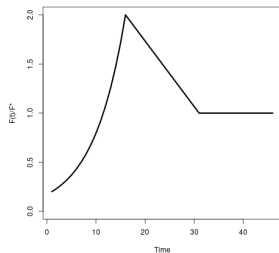
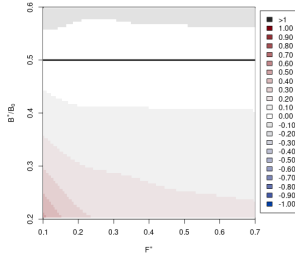
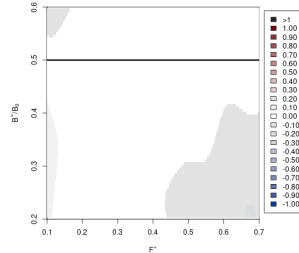
Components of Bias



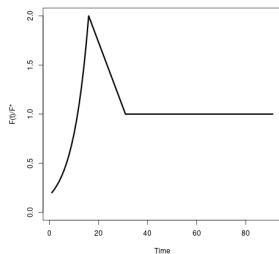
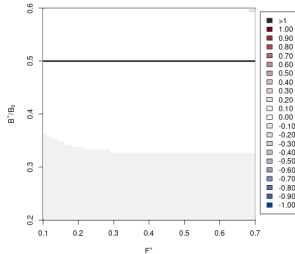
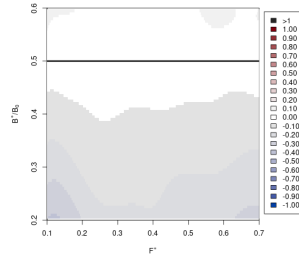




Relative Fishing Rate

Relative Bias in Estimated B^* Relative Bias in Estimated F^* 

Relative Fishing Rate

Relative Bias in Estimated B^* Relative Bias in Estimated F^* 

Summary

F^* , B^* , B_0 are not directly observable, but rather modeled quantities

Model Misspecification, Posterior Uncertainty, Bias

RP bias can be very large when production function assumptions are wrong

As model misspecification increases, RP biases tend to increase

B^* often appears relatively less sensitive to model misspecification than either F^* or B_0

F^* bias is strongly catch dependent

Bias depends on how similar the modeled and true production functions can be at the observed biomasses

A rich simulation-based method for describing global RP bias and a stepping stone for understanding other models

BH and Ricker SRRs

Age-Structured and Delay Difference Models

Productivity Extension

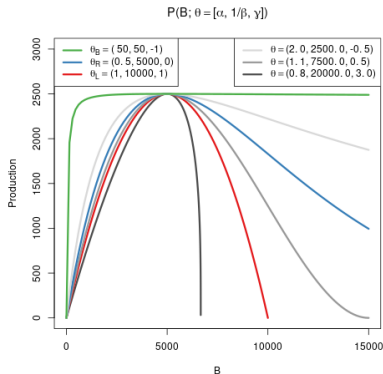
$$\frac{dB}{dt} = P(B; \theta) - (M + F)B$$

$$P(B; [\alpha, \beta, \gamma]) = \alpha B(1 - \beta\gamma B)^{\frac{1}{\gamma}}$$

$\gamma = -1 \Rightarrow$ Beverton-Holt

$\gamma = 1 \Rightarrow$ Logistic

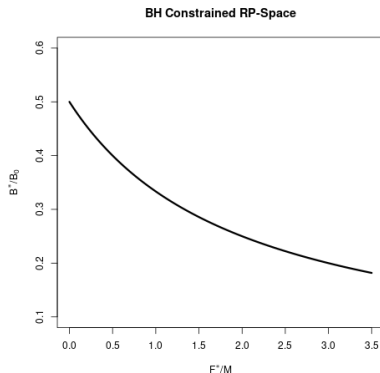
$\gamma \rightarrow 0 \Rightarrow$ Ricker



Productivity Extension

$$P_{\text{BH}}(B; [\alpha, \beta, -1]) = \frac{\alpha B}{(1 + \beta B)}$$

$$\frac{B^*}{\bar{B}(0)} = \frac{1}{\frac{F^*}{M} + 2}$$



Growth Extension

$$\begin{aligned}\frac{dB}{dt} &= \overbrace{w(a_0)R(B; \theta)}^{\text{Recruitment Biomass}} + \overbrace{\kappa [w_\infty N - B]}^{\text{Net Growth}} - \overbrace{(M + F)B}^{\text{Mortality}} \\ \frac{dN}{dt} &= R(B; \theta) - (M + F)N\end{aligned}$$

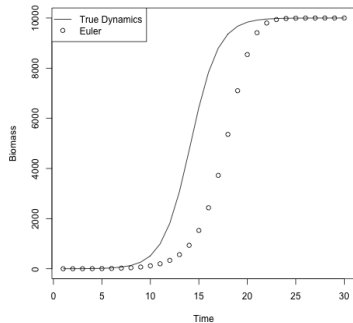
$$\begin{aligned}R(B; [\alpha, \beta, \gamma]) &= \alpha B(t - a_0)(1 - \beta\gamma B(t - a_0))^{\frac{1}{\gamma}} \\ w(a) &= w_\infty(1 - e^{-\kappa a})\end{aligned}$$

bullets of primary points of individual growth and maturity

Common Discretization

$$\frac{dB}{dt} = P_{\theta}(B(t)) - C(t)$$

$$B(\tau + 1) \approx B(\tau) + P_{\theta}(B(\tau)) - C(\tau)$$

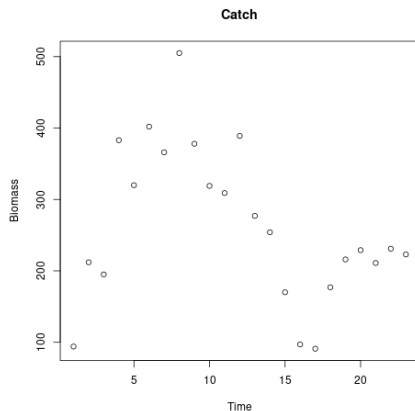


Catch Interpolation

$$t \in \mathbb{R}^+ \quad \tau = \lceil t \rceil - 1$$

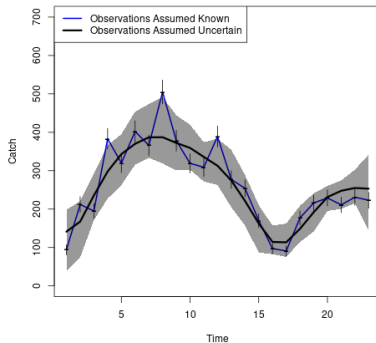
$$\mathbb{E}[y(t)] = \int_{\tau}^t x(t^*) dt^*$$

$$x(t) = \beta_0 + \sum_{j=1}^{T-1} \beta_j (t - \tau_j) \mathbf{1}_{t > \tau_j}$$

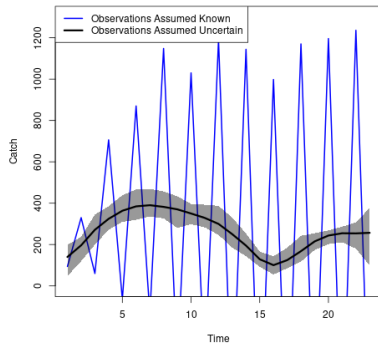


$$y(\tau_i) = \beta_0 + \sum_{j=1}^{i-1} \beta_j \left[\left(\frac{\tau_i^2}{2} - \tau_j \tau_i \right) \mathbb{1}_{\tau_i > \tau_j} - \left(\frac{\tau_{i-1}^2}{2} - \tau_j \tau_{i-1} \right) \mathbb{1}_{\tau_{i-1} > \tau_j} \right] + \epsilon_i$$
$$\beta_j \sim N(0, \phi) \quad \phi \sim \text{Half-Cauchy}(0, 1) \quad \epsilon_i \sim N(0, \sigma_i^2)$$

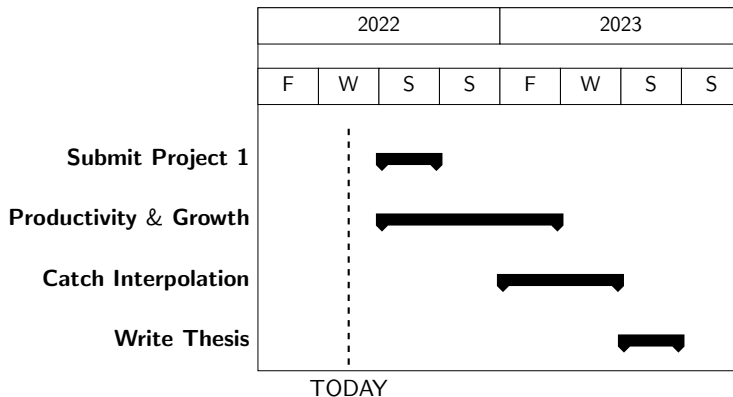
Observed Catch with Predictive Interpolations



Interpolated Instantaneous Catch



Timeline



Thanks and Acknowledgements NOAA, Sea Grant Ecetra

$$\frac{B^*}{\bar{B}(0)} = \frac{\left(\frac{\alpha}{M+F^*}\right)^{\frac{1}{\gamma}} - 1}{\left(\frac{\alpha}{M}\right)^{\frac{1}{\gamma}} - 1}$$
$$\alpha = (M + F^*) \left[1 - \frac{1}{\gamma} \left(\frac{F^*}{M + F^*} \right) \right]^{-\gamma}$$
$$\beta = \frac{1}{\gamma \bar{B}(0)} \left(1 - \left(\frac{\alpha}{M} \right)^{\frac{1}{\gamma}} \right)$$