Bias Estimation of Biological Reference Points Under Two-Parameter SRRs

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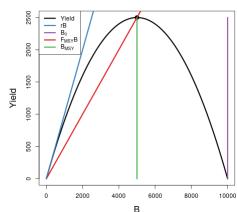


$$I_t = qB_te^{\epsilon} \quad \epsilon \sim N(0, \sigma^2)$$

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

$$RP: MSY, \ \frac{F_{MSY}}{M}, \ \frac{B_{MSY}}{B_0}$$

Yield and Related Quantities





Conceptually:

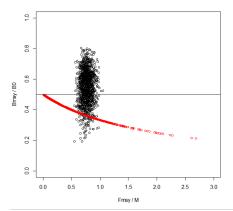
$$\frac{F_{MSY}}{M} \in \mathbb{R}^+ \quad \frac{B_{MSY}}{B_0} \in (0,1)$$

Mangel et al. 2013, CJFAS:

■ BH Model:

$$F_{MSY} \in \mathbb{R}^+$$
 $\frac{B_{MSY}}{\bar{B}(0)} = \frac{1}{F_{MSY}/M+2}$

 Similar Constraints for other Two-Parameter Curves



Conceptually:

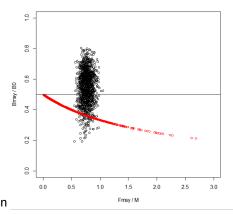
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- Similar Constraints for other Two-Parameter Curves
- Three-Parameter Relationships Allow Independent RP Estimation



- Starting from the rock bottom to build understanding
- And analysis of biases for the Scheaffer model can be seen here
- Attacking this problem from the ground up working towards age structred models is important due to the many computational issues that can arrise in ode modeling.
 - chaos
 - time integration error
 - estimation error
 - etc., etc.

Schaefer Analysis



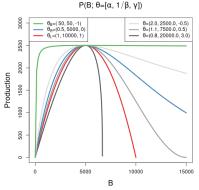
https://ggle.io/5EnI

Schnute 1985, CJFAS

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

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

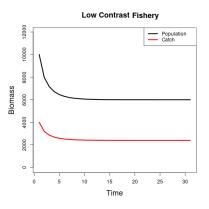
$$\gamma = -1 \Rightarrow$$
 Beverton-Holt $\gamma \to 0 \Rightarrow$ Ricker $\gamma = 1 \Rightarrow$ Logistic

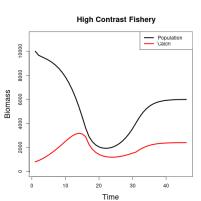


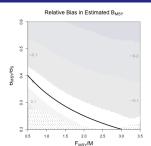
Introish Ideas list

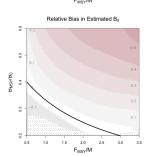
- PT/Schaffer work (link)
- Computational Difficulties
- Schnute Space Filling
- Catch/Contrast

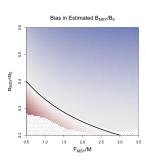
Catch

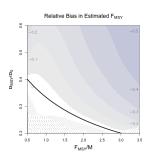


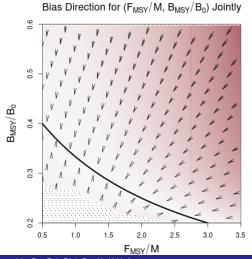




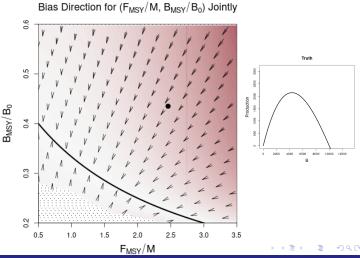


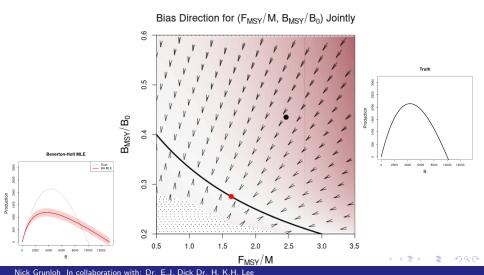


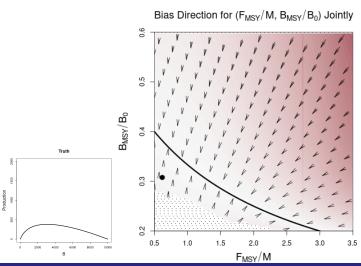




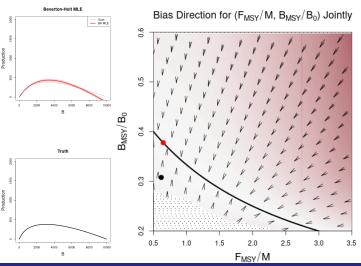




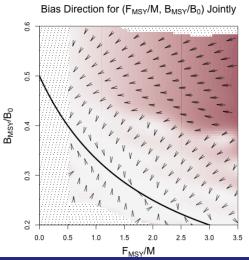




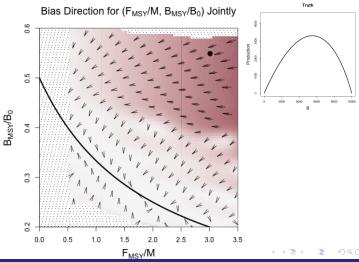


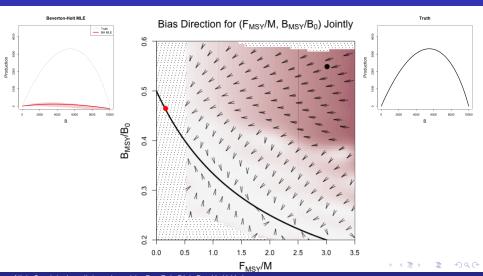


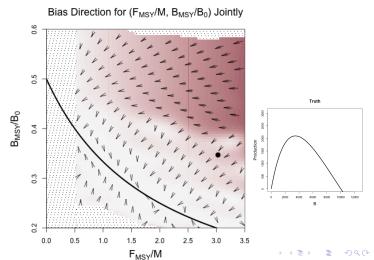


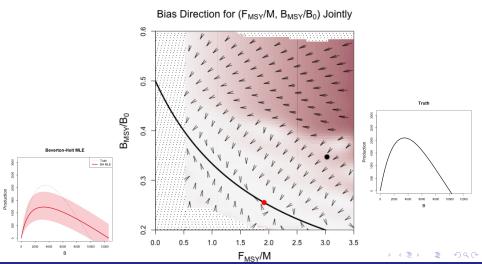












Conclusions

- Contrast story
- Importance of getting the computational details correct for moving to analysis of Delay Difference and age structure