

# Bias Estimation of Biological Reference Points Under Two-Parameter SRRs

Nick Grunloh

In collaboration with:

Dr. E.J. Dick

Dr. H. K.H. Lee

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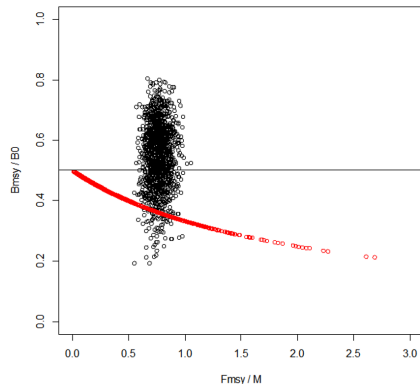
# Mangel et al. 2013, CJFAS

$$\frac{dB(t)}{dt} = \frac{\alpha B(t)}{1 + \beta B(t)} - (M + F(t))B(t)$$

$$h = \frac{\frac{\alpha}{M}}{4 + \frac{\alpha}{M}}$$

$$\frac{F^*}{M} = \sqrt{\frac{4h}{1-h}} - 1$$

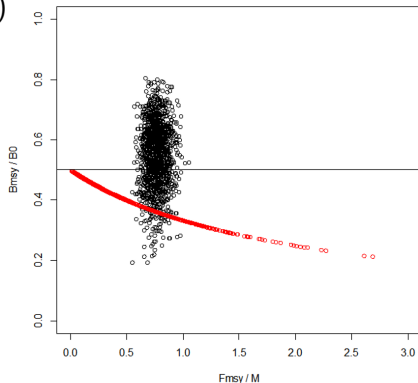
$$\frac{B^*}{B_0} = \frac{\sqrt{\frac{4h}{1-h}} - 1}{\frac{4h}{1-h} - 1}$$



# Mangel et al. 2013, CJFAS

$$\frac{dB(t)}{dt} = \frac{\alpha B(t)}{1 + \beta B(t)^{\frac{1}{\gamma}}} - (M + F(t))B(t)$$

Mangel et al. (2013) suggest exploration of three parameter stock recruit relationships (SRRs) to avoid pre-determined reference points (RP) in assessments



# A'Priori RP Prior Relationships

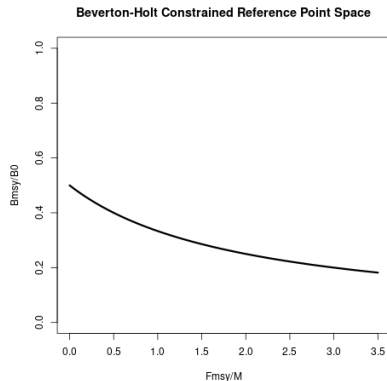
$$\frac{dB(t)}{dt} = \frac{\alpha B(t)}{1 + \beta B(t)} - (M + F(t))B(t)$$

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

$$\log(F^*) \sim N(\mu, \sigma^2)$$



$$2 \frac{B^*}{B_0} \sim \text{logit-N}(\log(2M) - \mu, \sigma^2)$$



# Pella-Tomlinson Production Model

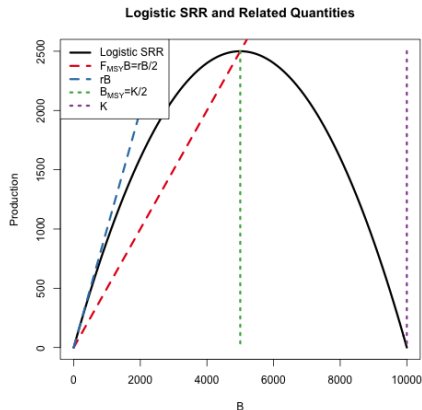
$$I(t) \sim LN(qB(t), \sigma^2)$$

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

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

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

$\gamma = 2 \Rightarrow$  Schaefer Model



# Pella-Tomlinson Production Model

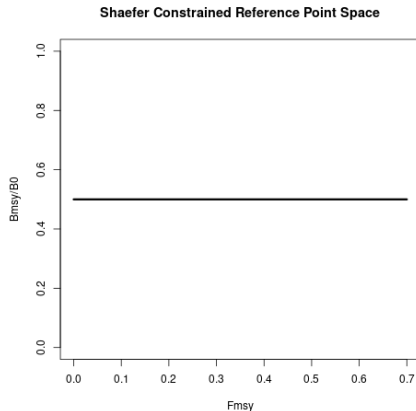
$$I(t) \sim LN(qB(t), \sigma^2)$$

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

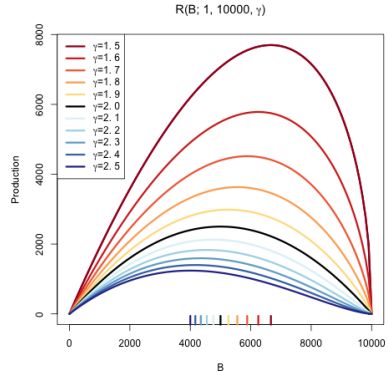
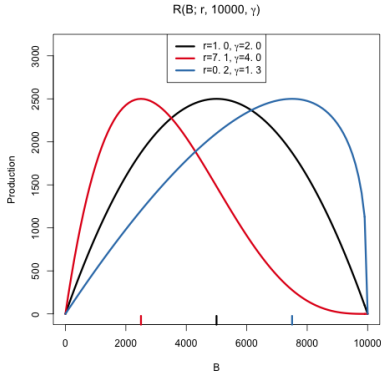
$$R_{\theta}(B) = \frac{rB}{\gamma - 1} \left(1 - \frac{B}{K}\right)^{\gamma-1}$$

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

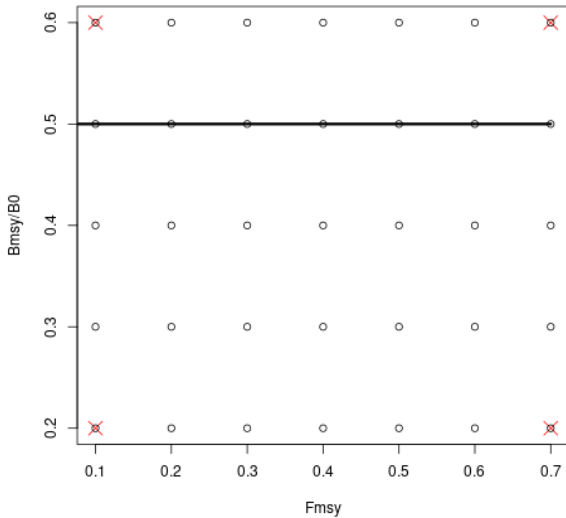
$\gamma = 2 \Rightarrow$  Schaefer Model



# Pella-Tomlinson Family of Curves

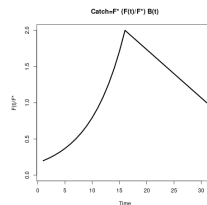
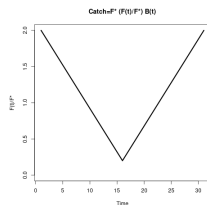
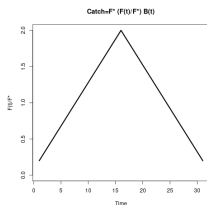
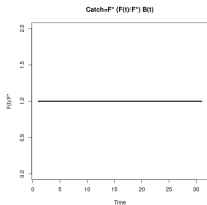


## Reference Point Space

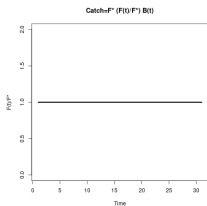




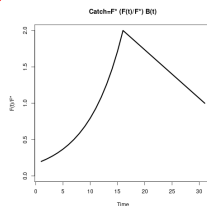
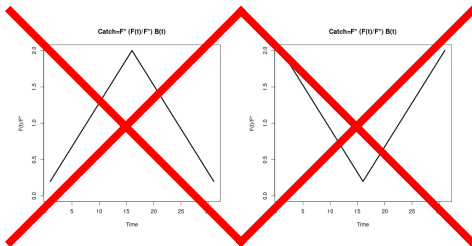
# Catch



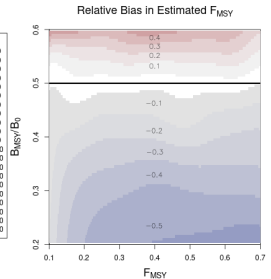
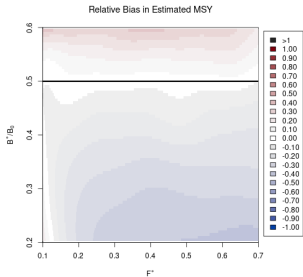
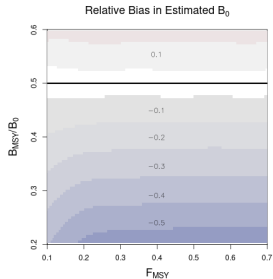
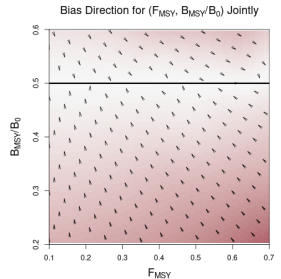
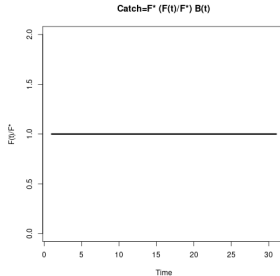
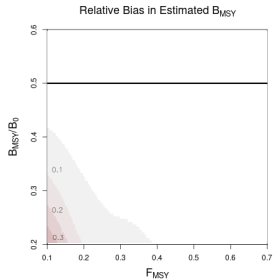
# Catch



Flat

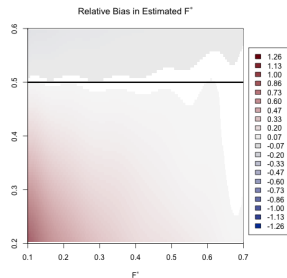
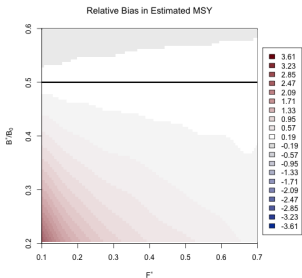
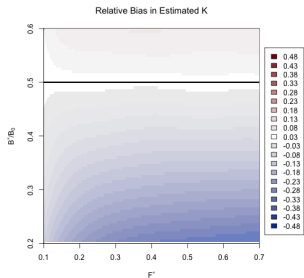
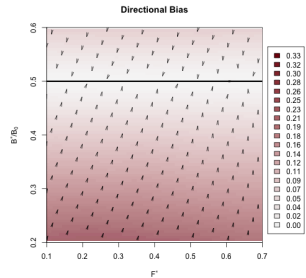
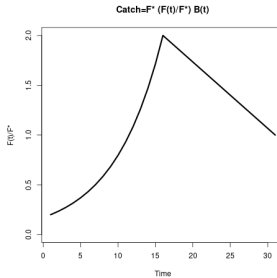
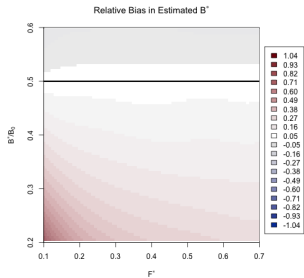


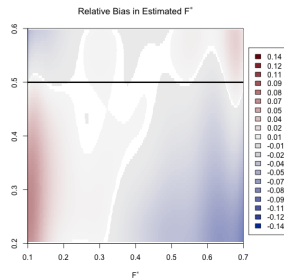
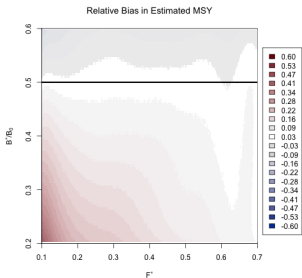
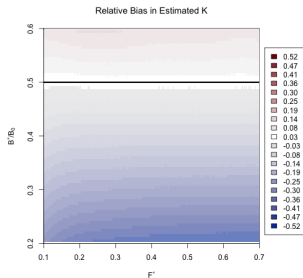
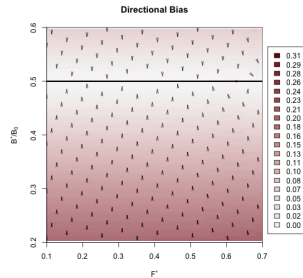
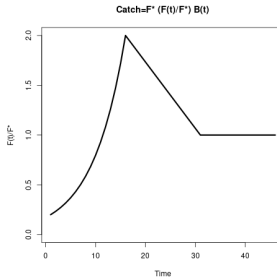
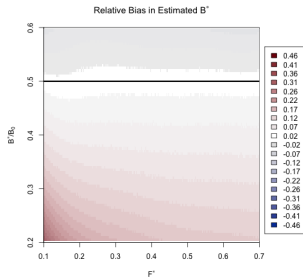
Contrast

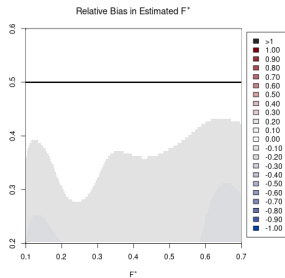
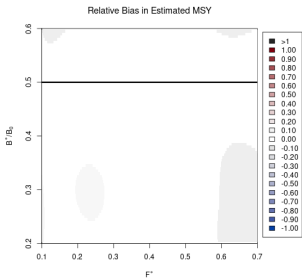
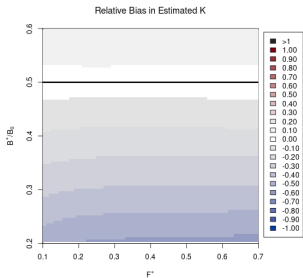
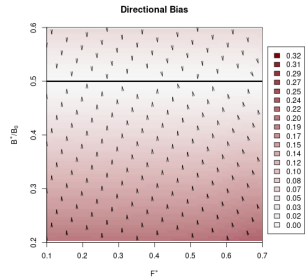
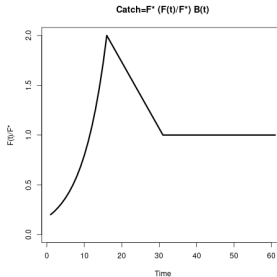
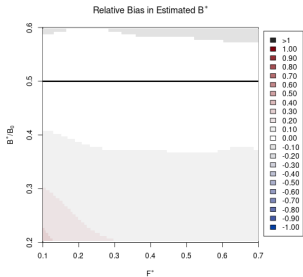


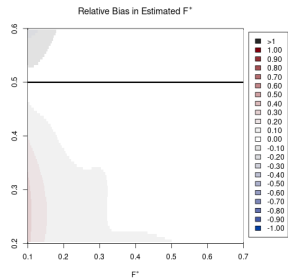
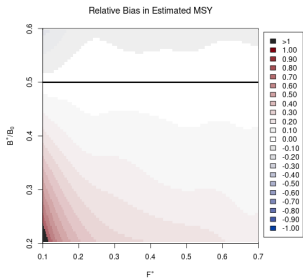
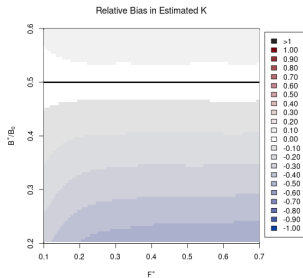
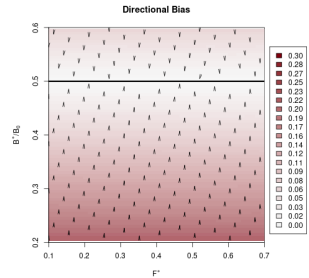
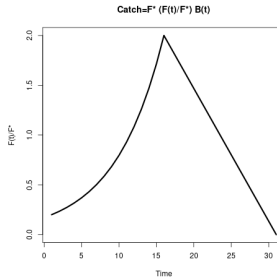
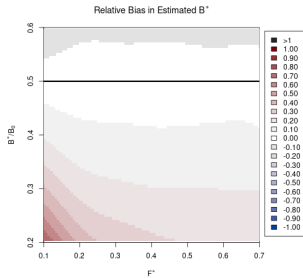
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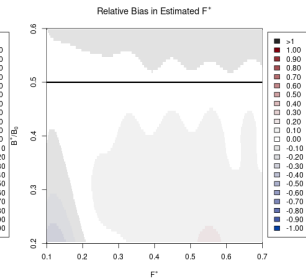
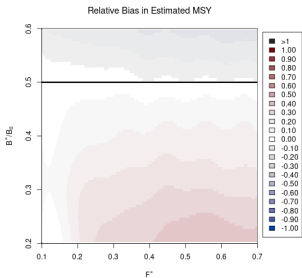
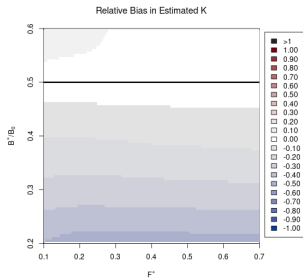
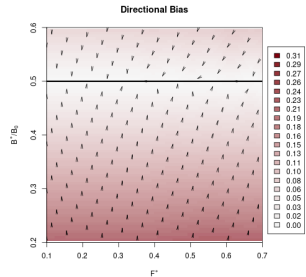
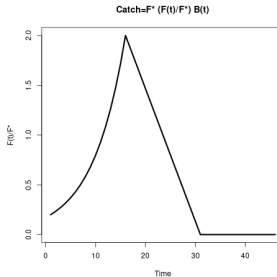
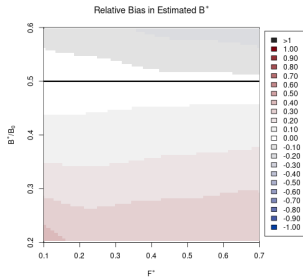
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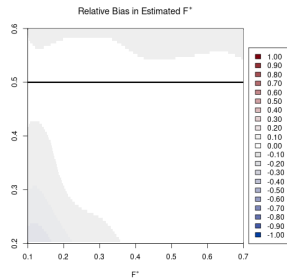
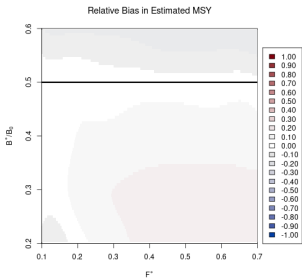
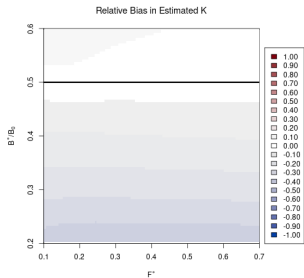
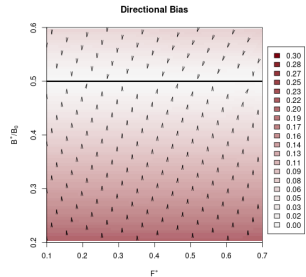
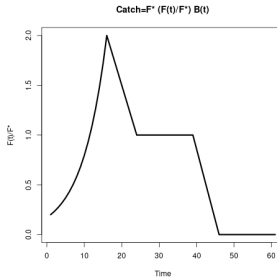
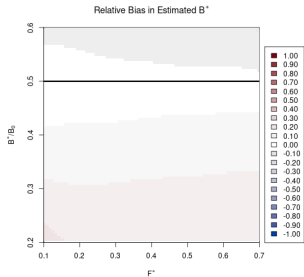


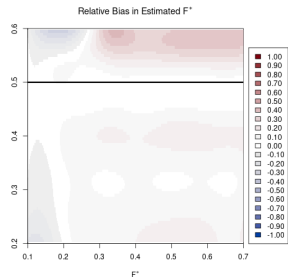
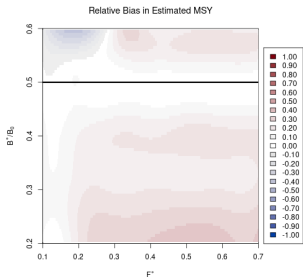
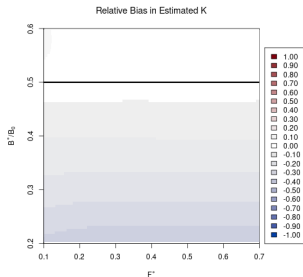
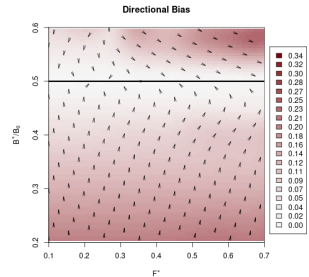
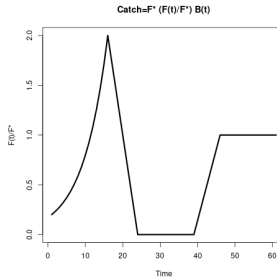
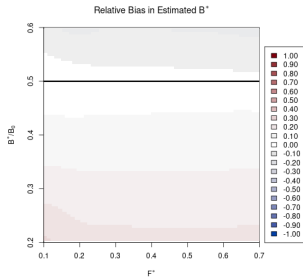








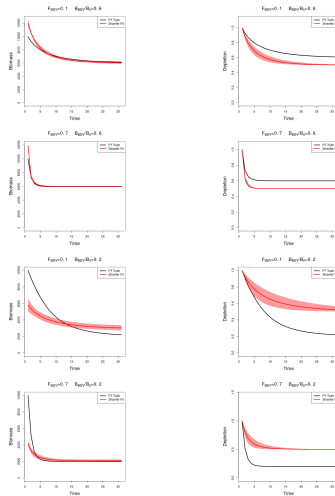
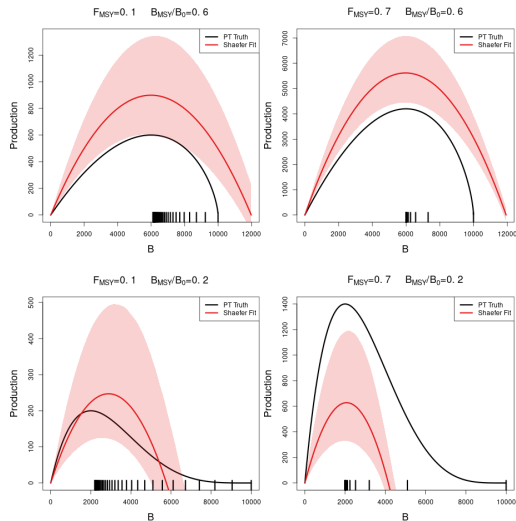




# Flat: Misspecified SRR

## Biomass

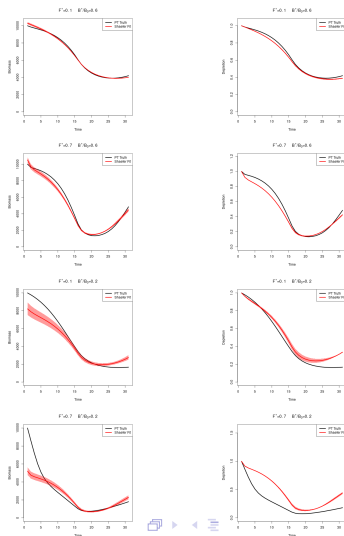
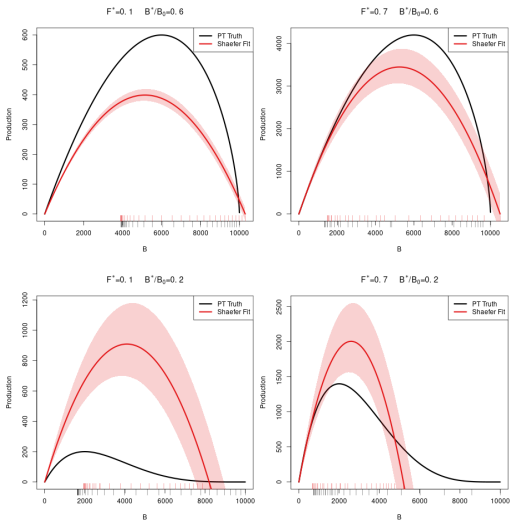
## Depletion



# Contrast: Misspecified SRR

## Biomass

## Depletion



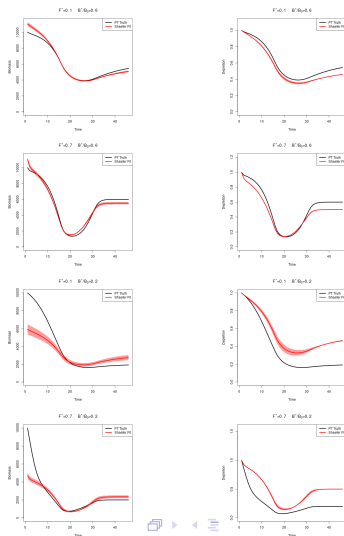
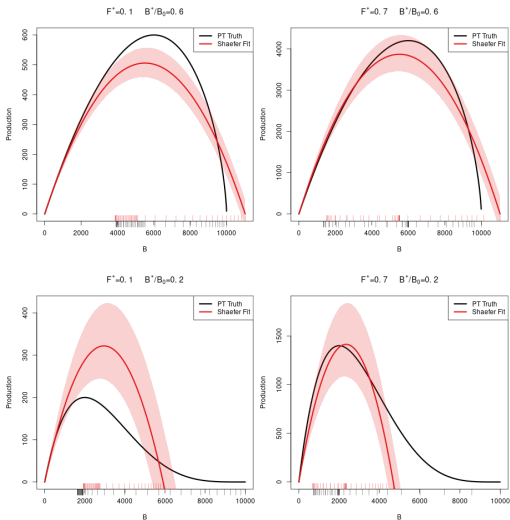
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# ContrastT45: Misspecified SRR

## Biomass

## Depletion



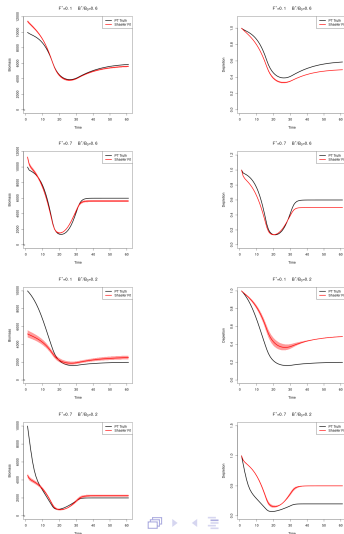
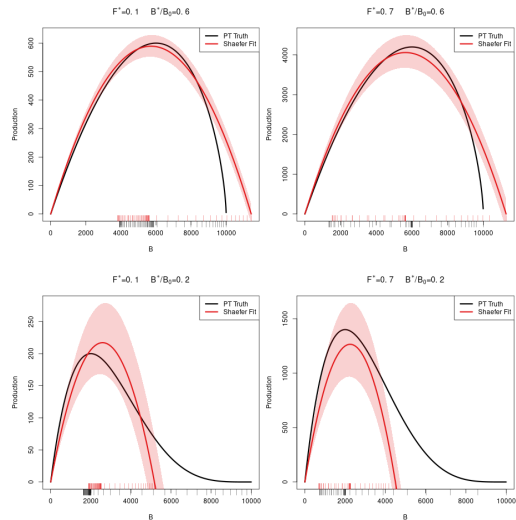
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Bias Estimation of Biological Reference Points Under Two-Parameter SRRs

# ContrastT60: Misspecified SRR

## Biomass

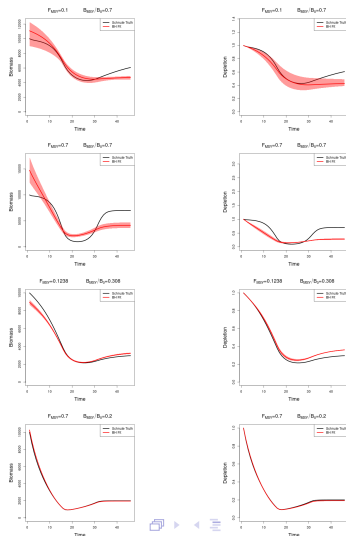
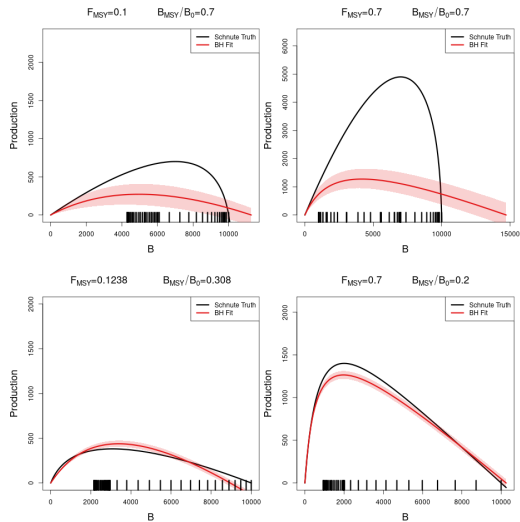
## Depletion



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Bias Estimation of Biological Reference Points Under Two-Parameter SRRs

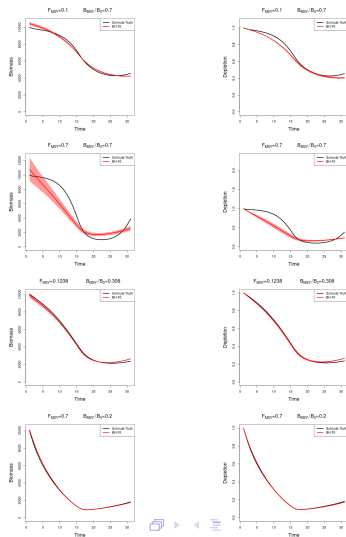
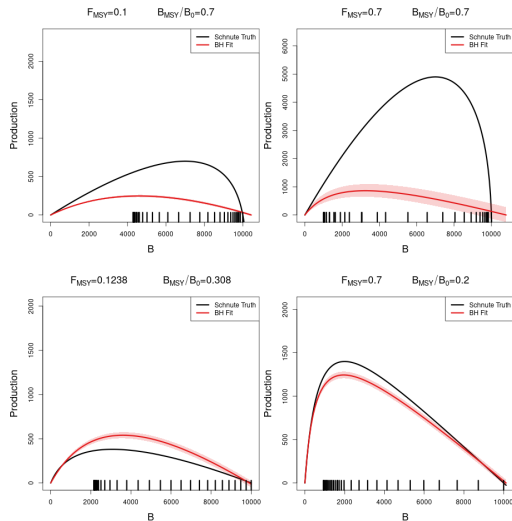
# SchnuteExpT45: Misspecified SRR Biomass Depletion



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Bias Estimation of Biological Reference Points Under Two-Parameter SRRs

# SchnuteExpT30: Misspecified SRR Biomass Depletion

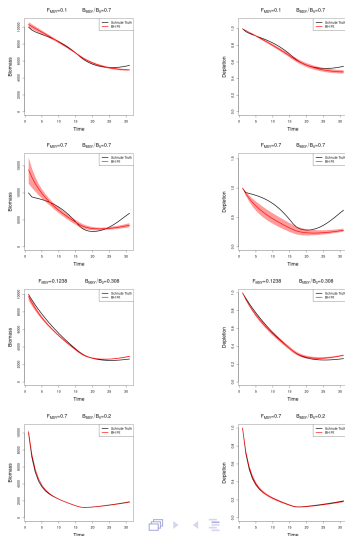
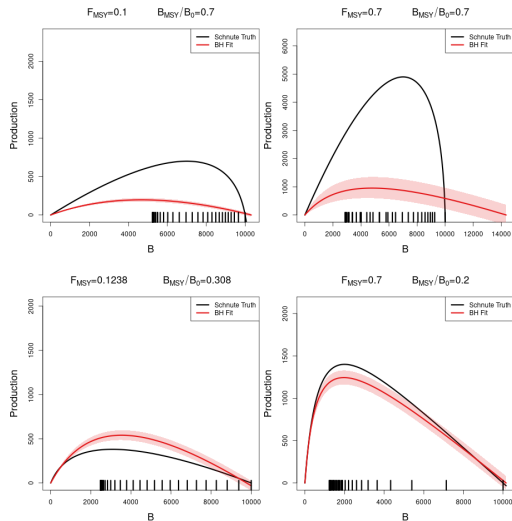




# SchnuteExpT30L2: SRR

Biomass

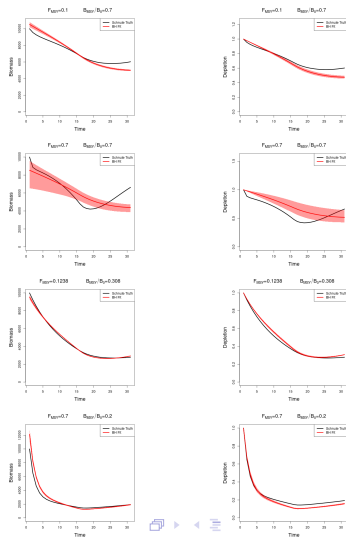
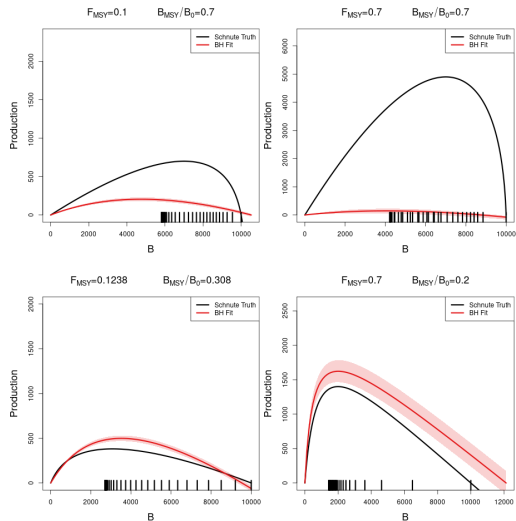
Depletion



# SchnuteExpT30L3: SRR

Biomass

Depletion



# SchnuteFlatT30: SRR

## Biomass

## Depletion

