

Principles of Islands Red King Crab MPT comments May 2016

Continue the work on survey biomass and length frequency weighting issues to improve the model fits to abundance data;

- Addressed in #2 below.

Implement the Francis tuning method to estimate length composition effective sample sizes;

- The Francis effective N calculation was added to the model. In addition, other multipliers on the survey length frequencies were evaluated.

Provide results for a random effects model and three-year weighted average for the September meeting

- The random effects model was fit to the survey biomass data and MMB, OFL, and ABC estimated. The estimates using the three-year weighted average are also included.

Summary of Major Changes:

Management: None.

Input data: Survey (2016) and bycatch (2015) data were incorporated into the assessment.

Assessment methodology: Model output for male only fit is presented with the same model configuration as 2015.

Assessment results: Male biomass estimates from the 3-year running average and a random effects model fit to survey male biomass $\geq 120\text{mm}$ are used to estimate MMB at mating, OFL and ABC.

Crab Plan Team September 2015 comments not addressed

Incorporate a mean-unbiased log normal likelihood for survey numbers

- Next time.

Discuss the poisson vs. negative binomial for survey estimates of abundance and CVs

- Currently all of the data in the model are those that are passed from Bob Foy and the Kodiak lab, but given the over-dispersion in the data, a negative binomial (or something similar) might be more appropriate, particularly for estimates of variance. The CVs sent by Bob are used in the assessment, but bootstrapped variances are much larger.

Consider ADFG pot survey data and retained catch size frequency data

- These data area not yet incorporated, but may be useful in exploring the mechanics of time-varying catchability.

Pribilof red king crab (biomass in tons)

| | MSST | Biomass (MMB mating) | at TAC | Retained Catch | Total Catch | OFL | |
|------|-------|----------------------------|-----------|-------------------|-------------------|-------|---|
| 0/11 | 2,255 | 2,754 ^A | 0 | 0 | 4.2 | 349 | |
| /12 | 2,571 | 2,775 ^{B*} | 0 | 0 | 5.4 | 393 | |
| 2/13 | 2,609 | 4,025 ^{C**} | 0 | 0 | 13.1 | 569 | |
| 3/14 | 2,582 | 4,679 ^{D**} | 0 | 0 | 2.25 | 903 | |
| 4/15 | 2,871 | 8,894 ^{D**} | 0 | 0 | 1.76 | 1,359 | 1 |
| 5/16 | 2,756 | 9,062 ^{**} | 0 | 0 | 0.32 ¹ | 2,119 | 1 |

3 year running average of male biomass ($\geq 120\text{mm}$) at survey time was calculated using the weighted average with weights being the inverse of the variance,

$$BWRA_t = \frac{\sum_{i=1}^3 \frac{MMB_i}{w_i}}{\sum_{i=1}^3 \frac{1}{w_i}} \quad (4)$$

MMB_t

Estimated male biomass ($\geq 120\text{mm}$) from the survey data

w_t

The weight associated with the estimate of MMB in year t

w_t is calculated as the variance of the log(biomass) using the CV estimates of MMB from the survey provided by the Kodiak lab:

$$w_t = \ln((CV_t)^2 + 1)$$

CV_t

Coefficient of variation associated with the estimate of MMB at

Random effects model Likelihood

$$\sum_{i=1}^{n} \left\{ 0.5 \left(\log(2\pi\sigma_i^2) + \left(\frac{(\widehat{B}_i - B_i)^2}{\sigma_i^2} \right) \right) \right\} + \sum_{j=1}^{n} \left\{ 0.5 \left(\log(2\pi\sigma_j^2) + \left(\frac{(\widehat{B}_j - \widehat{B}_{j|j})^2}{\sigma_j^2} \right) \right) \right\}$$

\widehat{B}_i = log of observed biomass in year i

B_i = model estimated log biomass in year i

σ_i^2 = the variance of observed log biomass in year i

σ_j^2 = the variance of the deviations in log survey biomass between years (i.e. process error variance). σ_j^2 was estimated

where α is a parameter estimated in the random effects model.

n = the number of years of survey biomass values

Survey biomass in 2016 declined to 4,150 t from 15,173 t in 2015

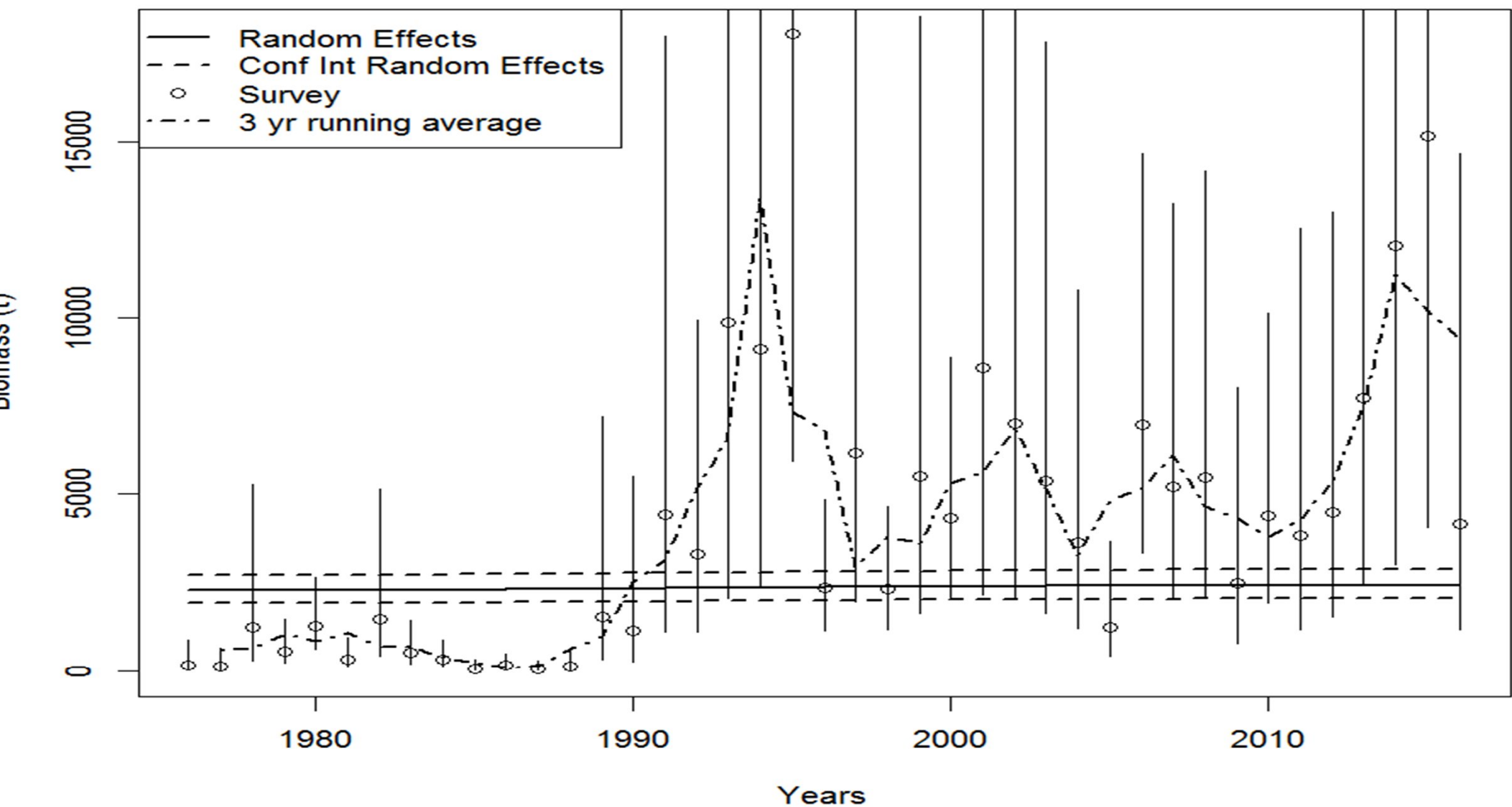


Figure 14. Three-year running average and random effects model fit to male biomass > 120mm at survey.

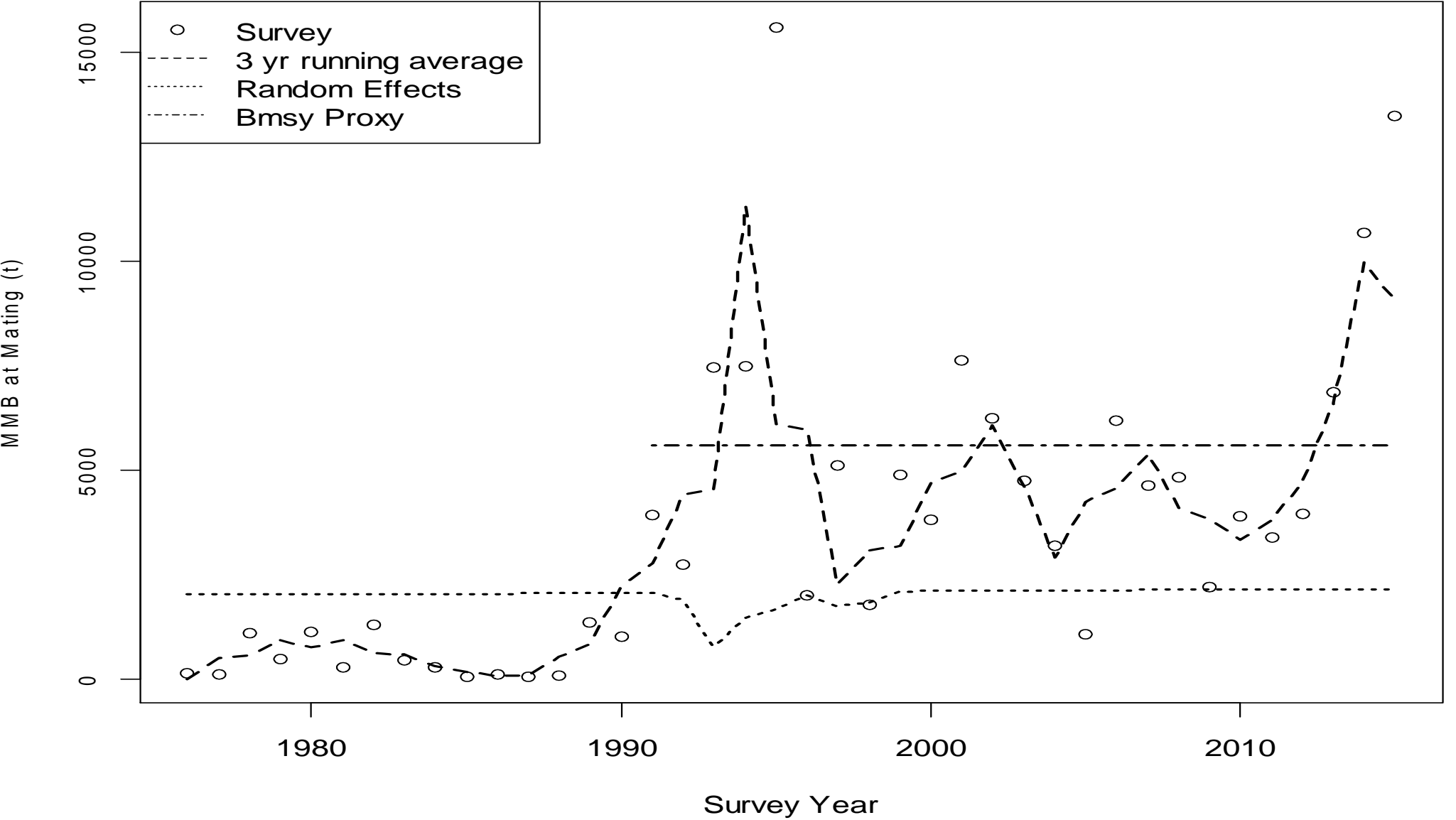


Figure 15. MMB at mating (February 15 of survey year +1) estimated from the survey data, 3 yr running average and the Random effects model. Bmsy proxy is the average of the 1991 to 2015 MMB at mating survey data (February 15 1992 to February 15 2016).

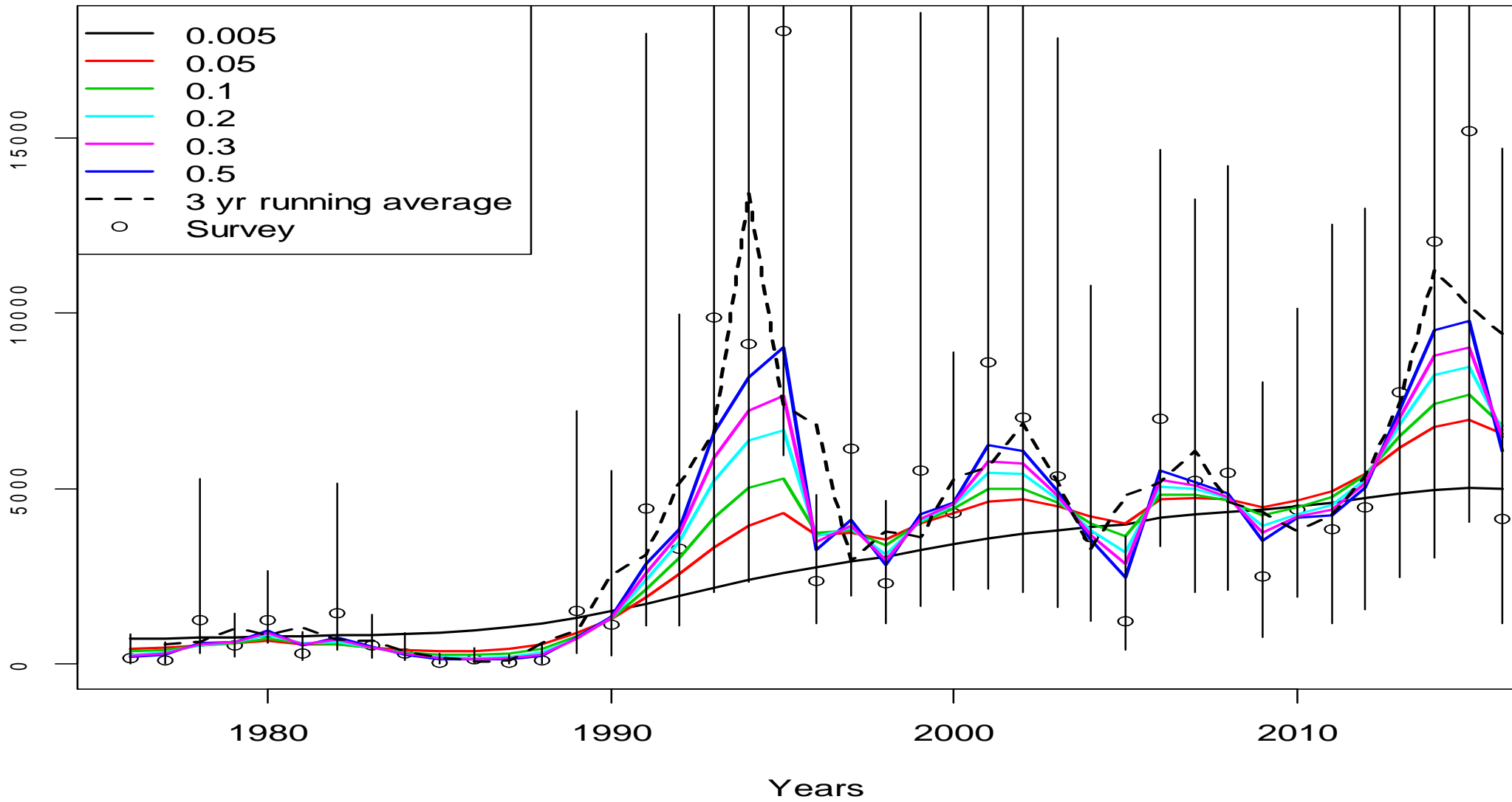
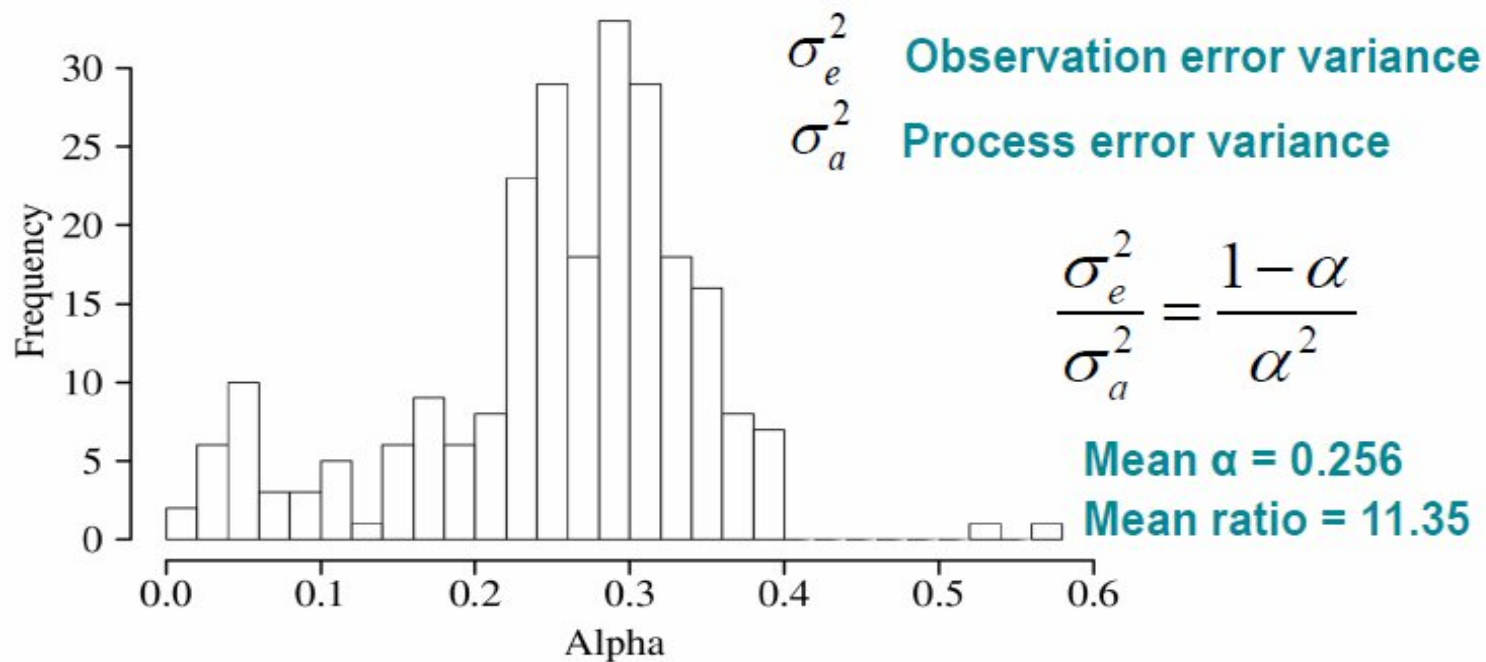


Figure 26. Random effects model estimates of biomass with process error fixed at 0.005, 0.05, 0.1, 0.2, 0.3, 0.5.

From Spencer presentation at Wakefield 2015

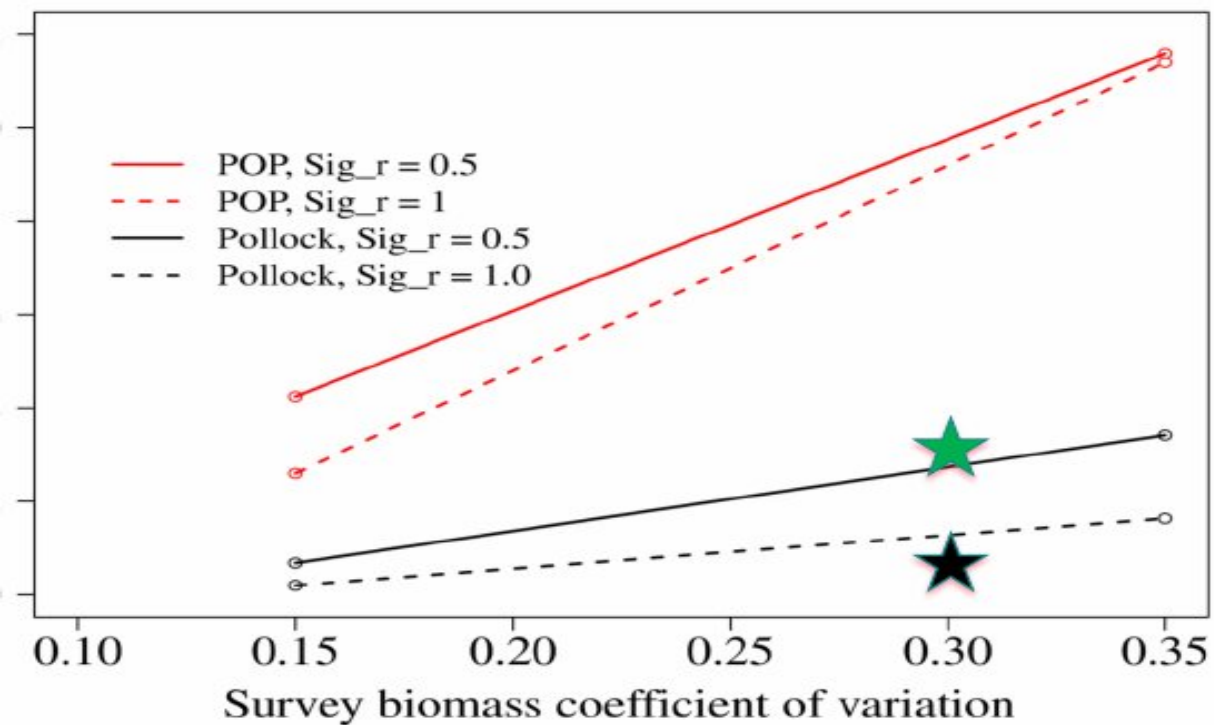
A simple exponential smoothing model can give information on the ratio of variances

$$\hat{z}_t = (\alpha)y_t + (1 - \alpha)[\alpha y_{t-1} + \alpha(1 - \alpha)y_{t-2} + \alpha(1 - \alpha)^2 y_{t-3} + \dots]$$



Spencer Wakefield 2015

The variance ratio is a function of stock longevity, recruitment variability, and survey variability



Used as a prior to constrain the estimate of process error standard deviations

Implied from fit to GOA dogfish

Observation error variance on log scale is $\ln(cv^2+1)$

Mean cv of survey biomass is 0.67

Mean Observation Error variance on log scale = 0.38

Fitting a simple exponential model to Pribilof survey data using HoltWinters function in R gives,

Alpha = 0.396, = variance ratio of 3.75 and process error of $0.38/3.75 = 0.102$.

Pribilof red model variance of first difference in log biomass 0.046,
Bristol Bay red king crab model 0.089.

Integrated assessment model fit to male numbers (male only model)

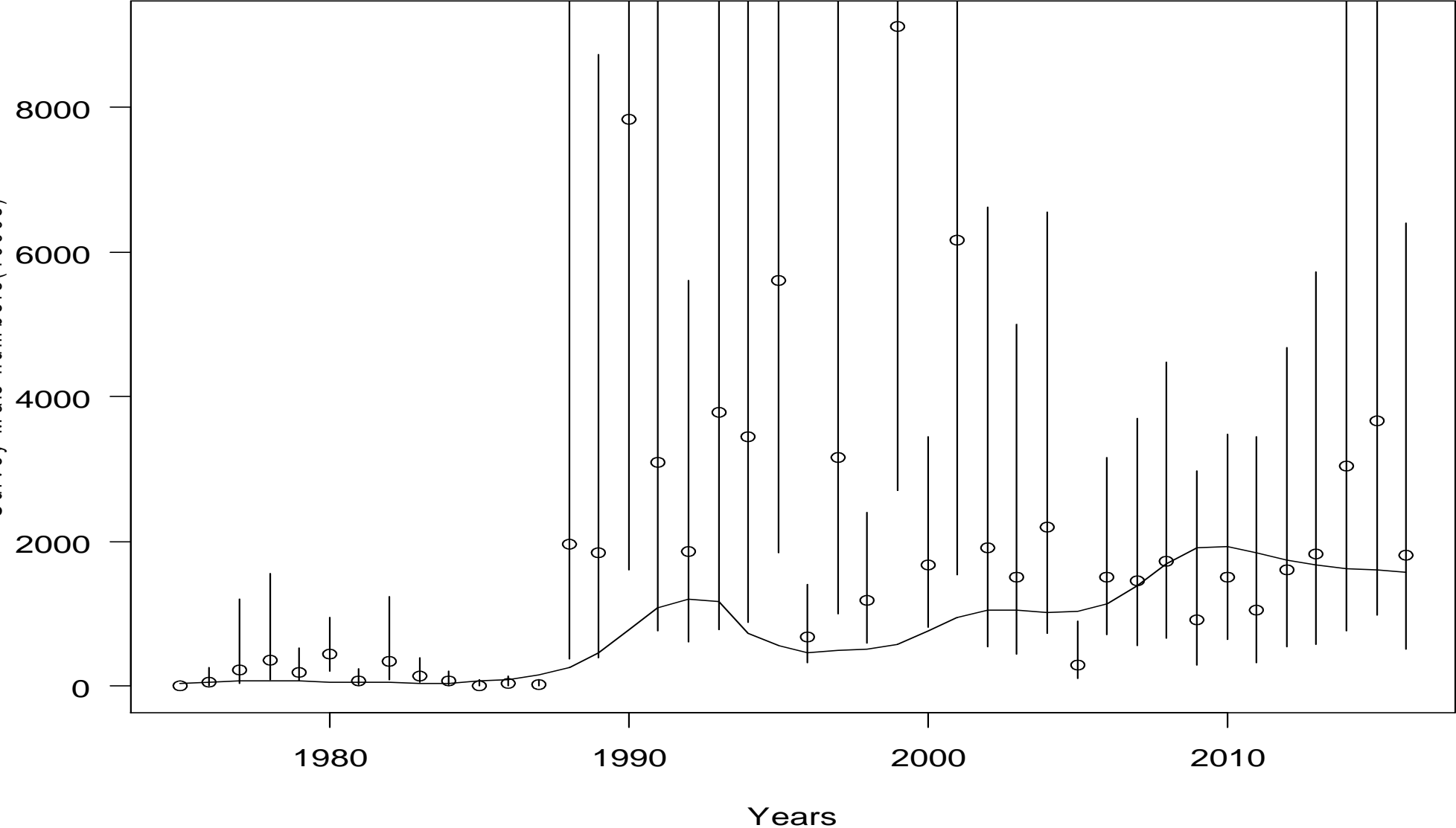
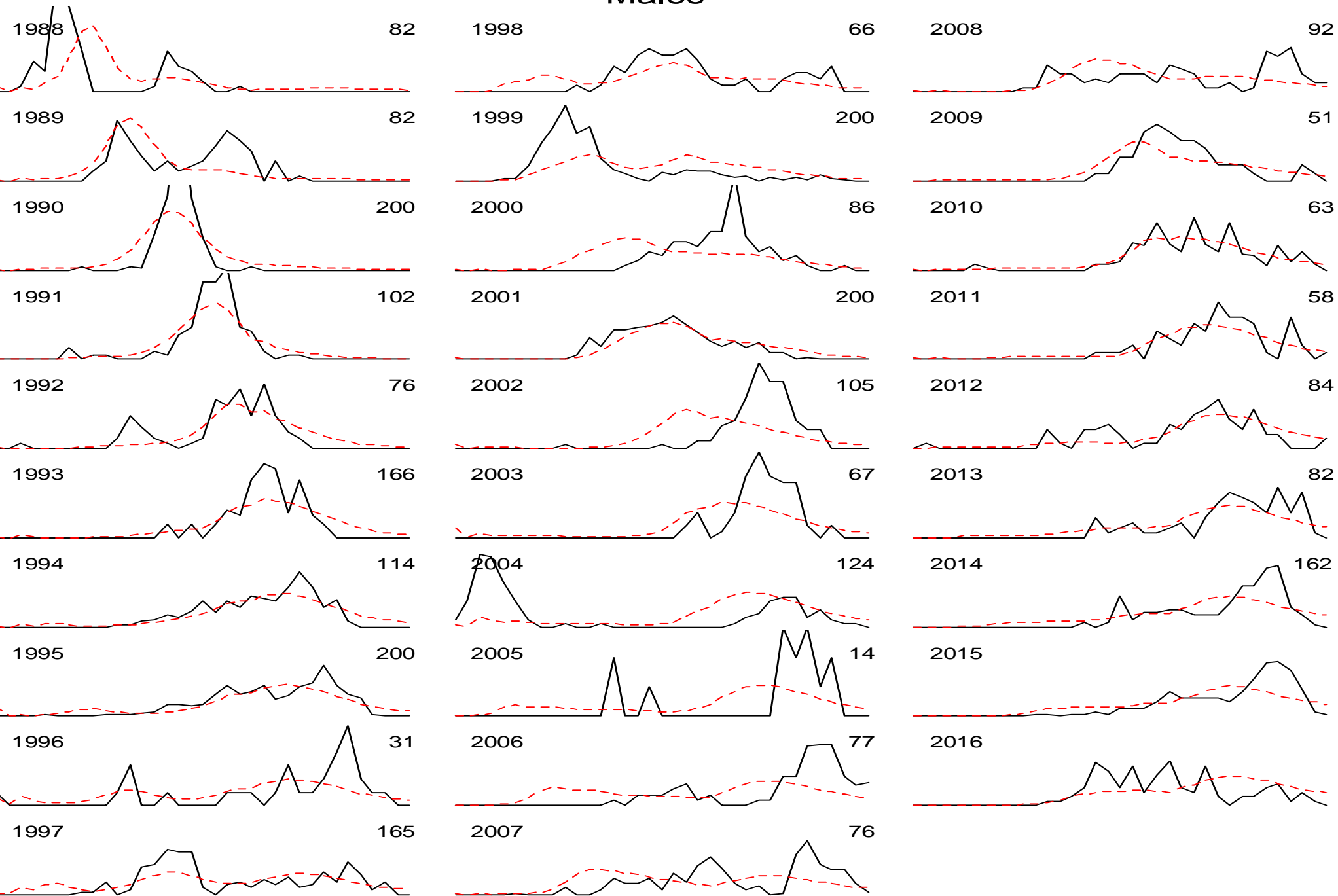


Figure 20. Model fit to survey male numbers.

Males



Survey length sample size reduction – Francis N multiplier 0.05 (model did not converge)

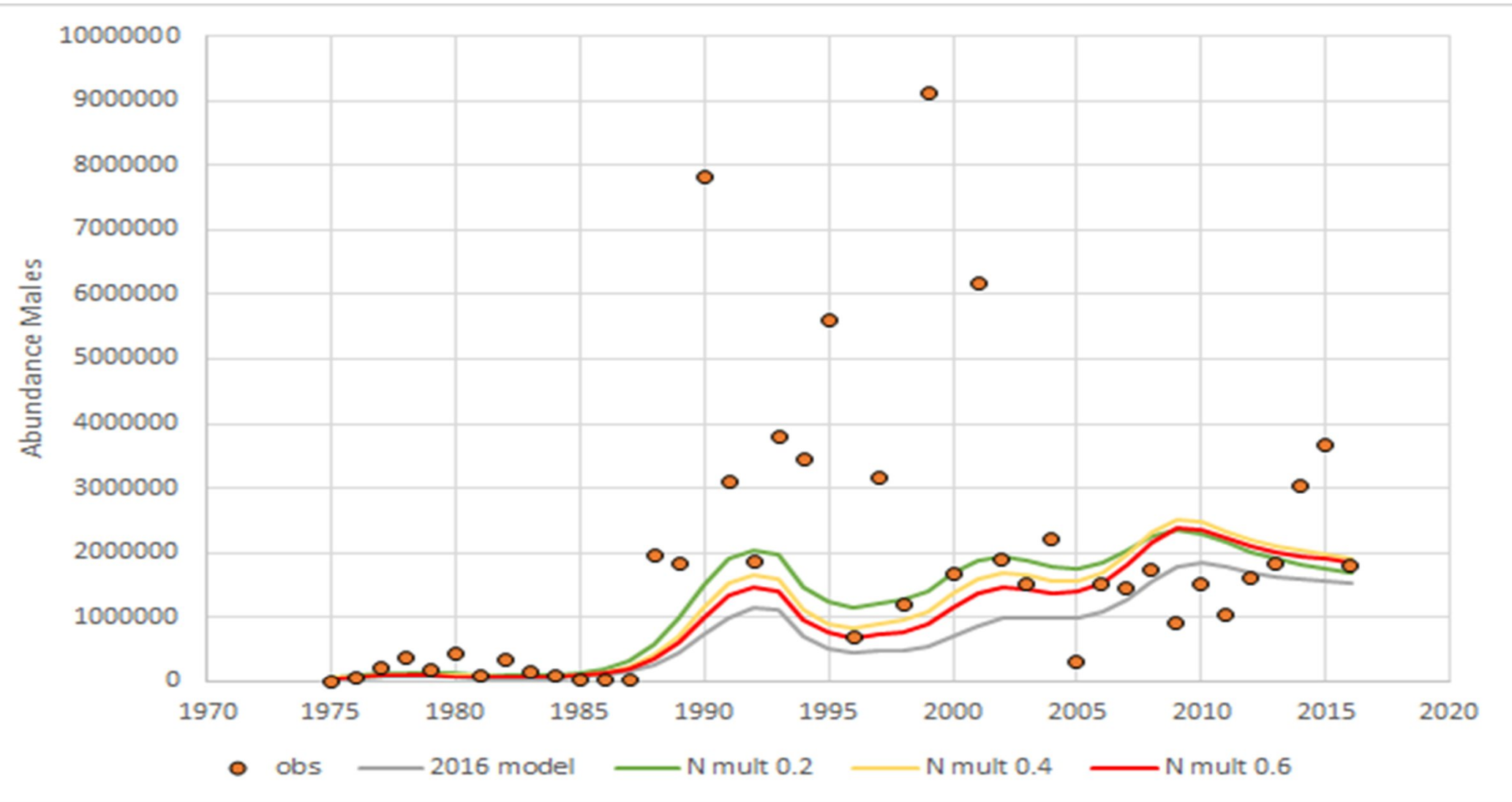


Figure 25. Fit to male abundance for the 2016 base model and model scenarios with multipliers on the survey length sample size of 0.2, 0.4 and 0.6

A4. Likelihood component contribution to the likelihood and associated weights for the assessment model scenarios with multipliers on the survey length sample sizes of 0.2, 0.4, 0.6 and the base model (1.0).

| Likelihood component | Base Model (1.0) | 0.2 | 0.4 | 0.6 | Weighting |
|----------------------------------|------------------|--------|--------|--------|---------------------------------|
| Survey numbers (males) | 45.7 | 29.9 | 32.7 | 36.1 | .36 -1 (CVs) |
| Survey length frequencies (male) | 10,012.3 | 2018.9 | 4024.6 | 6023.7 | 18-200 (Base model sample size) |
| Catch | 0.003 | 0.001 | 0.001 | 0.001 | .005(CV) |
| Trawl | 0.019 | 0.019 | 0.019 | 0.019 | .01 (CV) |
| Smoothness penalties | | | | | |
| Trawl fishing mortality | 38.6 | 38.4 | 38.3 | 38.4 | 1 (CV) |
| Fishing mortality | 4.3 | 4.3 | 4.3 | 4.3 | 1 (CV) |
| Recruitment | 48.9 | 20.4 | 30.1 | 37.5 | 1 (CV) |

Pribilof red king crab alternatives (biomass in tons)

| Assessment method | OFL | B_{MSY} | MMB At mating Feb 15 2017 fishing at OFL | B/B_{MSY} (MMB) | MMB at mating Feb 15 2016 | γ | Years to define B_{MSY} (MMB at mating) | F_{MSY} | ABC ($p^*=0.49$) |
|------------------------------------|-------|-----------|--|-------------------|---------------------------|----------|---|-----------|--------------------|
| Planning average | 1,462 | 5,512 | 6,980 | 1.25 | 9,062 | 1.0 | 1991/92-2015/16 | 0.18 | 1,430 |
| Random Effects Model | 119 | 5,512 | 2,044 | 0.37 | 2,154 | 1 | 1991/92-2015/16 | 0.05 | 119 |
| Observed survey | 370 | 5,512 | 3,332 | 0.60 | 13,457 | 1 | 1991/92-2015/16 | 0.10 | 350 |
| Integrated assessment (males only) | 822 | 3,881 | 5,160 | 1.33 | 6127 | 1 | 1991/92-2015/16 | 0.18 | 822 |
| Integrated assessment (males only) | 1,931 | 1,598 | 4,066 | 2.5 | 6127 | 1 | 1983-present recruitment | 0.49 | 1,931 |