## BSAI Tanner Crab

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## Roadmap for the 2018 SAFE Chapter

- Main text
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## CPT/SSC Comments

## May2018 Crab Plan Team Meeting

The CPT outlined a number of alternative models built on the 2017 assessment model (2017AM) as the base model to be evaluated.
Response: The CPT referred to these models as 2018B0, 2018B1, 2018B2, 2018B3, 2018B4 and 2018B5. These models were all run for this assessment, but renamed as 18A, 18B, 18C0, 18C1, 18D0, and 18D1, where " 18 " refers to the assessment year, $\mathrm{A} / \mathrm{B} / \mathrm{C} / \mathrm{D}$ refers to different datasets included in the likelihood, and $0 / 1$ refers to whether ( 1 ) or not ( 0 ) survey abundance time series were included in the fitting process in addition to survey biomass time series. 2017AM is subsequently referred to herein as 17AM. In addition to the alternative model scenarios requested by the CPT, several additional scenarios were also run: 17AMu, 18C0a, 18Cla, 18C2a, and 18C3a. Scenario 17AMu represents the 2017 assessment model re-run with revised (i.e., "u"pdated) data for the crab fisheries. The "a" in the remaining scenarios refers to ones in which the likelihood component for male maturity ogive data was down-weighted, whereas " 2 " and " 3 " refer to fixing the survey catchability and selectivity parameters to match ones from Somerton and Otto (1999)'s underbag experiment.

## October 2017 SSC Meeting

Comment: "The SSC endorses all of the CPT recommendations with respect to the poor fits to some of the retained catch time series, poor fits to the size composition data for retained catch and survey data, and issues with the total directed fishery selectivity curve for males (in particular the 1996 'outlier')." Response: With respect to the 1996 'outlier', this was a result of the combination of a very small sample size for the 1996 size compositions and the using the mean size-st-50\%-selected for 1991-1996 as the value for the size-at- $50 \%$-selected prior to 1991. Because the sample size for 1996 was small, the 1996 size-at-50\%-selected essentially became a free parameter uninformed by the 1996 data but sensitive to changes in the overall likelihood through changes in the mean value. Regarding the other issues, see the responses to CPT comments below.

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## CPT/SSC Comments

## June 2017 SSC Meeting

The SSC requested an evaluation of all parameters estimated to be at or very near bounds, or substantially limited by priors (unless those priors can be logically defended).
Response: An initial approach to evaluating parameters at or near bounds using ADMB's likelihood profiling capability revealed that errors had apparently been introduced to the profiling algorithm in a recent version (11.2) of the ADMB libraries. These errors have subsequently been resolved, and will be incorporated in the next scheduled version release (11.7). However, likelihood profiling results from the author's version (11.5/11.6) would provide erroneous results.

## May2017 Crab Plan Team Meeting

The CPT noted that the EBS growth data should be used in the assessment if at all possible, that the growth increment function should be adopted, and that the scale parameter should be estimated rather than being set to 0.75.
Response: All three requests have been addressed in the assessment (Model B1 and subsequent models).

The CPT noted that there was a tendency for the model to overpredict the abundance of large crab and recommended that the issue be evaluated by modeling retention with a logistic curve that asymptotes to a value less than one.
Response: The option of fitting a retention curve that asymptotes less than one has been implemented in the model framework. Models B2a, B2b and B3 incorporate this option and address this issue. Results from these models suggest that retention is indeed asymptotically less than one.

## Changes From 2017 Assessment

- Changes to model
- Male maturity data fit in model
- New trawl survey data for 2018
- annual male maturity ogives (1990+)
- size compositions by sex, shell condition, maturity
- male maturity ogives
- New Fishery Data for 2017/18
- Directed Tanner crab fishery
- 2017/18 catch and size compositions
- snow crab pot fishery
- 20117/18 bycatch, size compositions
- BBRKC pot fishery
- 2017/18 bycatch, size compositions
- groundfish fisheries
- 2017/18 bycatch, size compositions for combined gear
- Revised Crab Fishery Data
- Tanner crab pot fishery
- Revised total catch biomass, 1992/93+
- snow crab pot fishery
- Revised total catch biomass, 1992/93+
- BBRKC pot fishery
- Revised total catch biomass, 1992/93+


## Management Regions



## Fishery Trends

## Retained catch




## Bycatch




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## Recent Fishery Trends: Retained Catch



area

- West 166W
- East 166W
- all EBS


## Recent Fishery Trends:

 Total Catch

## Recent Fishery Trends: Directed Fishery Total Catch Size Comps



## Recent Fishery Trends: Snow Crab Total Catch Size Comps




## Recent Fishery Trends: BBRKC Total Catch Size Comps




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## Recent Fishery Trends: Bycatch in the Groundfish Fisheries


gear

- all
- fixed
- trawl


## Bycatch Trends in the Groundfish Fisheries




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## NMFS EBS Trawl Survey Trends



## NMFS EBS Trawl Survey Trends (recent)


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## NMFS EBS Trawl Survey Trends



NMFS EBS Trawl Survey Trends: Preferred Males


$\because$ all EBS

- East 166



## Spatial patterns



## Revisions to Total Catch in the Crab Fisheries

- Previously, Total Catch = Expanded Discard Catch + Retained Catch
- Discard catch: from at-sea observers classifying discards
- Expanded Discard Catch = (pots fished)/(pots obs)*(obs discard catch)
- biomass based on old LW relationships
- included "summary pot" data?
- changes in "target" (total effort, observed pots)?
- Retained catch: from fish ticket data
- Now: TC = Expanded Total Catch = (pots fished)/(pots obs)*(obs total catch)
- observed total catch: from at-sea observers
- ignores observer discard/retained classification
- includes only "detail pot" data
- biomass based on standard LW relationships


## Revisions to Total Catch in the Directed Fishery



- 2016: fishery closed


## Revisions to Total (By)Catch in the Snow Crab Fishery



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## Effects on Assessment Model Results



## Effects on Assessment Model Results




## Effects on Assessment Model Results



## Assessment Model Results: 2017 to 2018




## Tier 3 stage/size-based population dynamics model

- model year runs July 1 to June 30
- sex, shell condition, maturity state, carapace width
- sex/stage-based natural mortality (2 time stanzas)
- trawl survey occurs July 1
- fisheries occur Feb. 15
- directed fishery (retained and bycatch)
- bycatch in snow crab fishery
- bycatch in BBRKC fishery
- bycatch in groundfish fisheries
- sex-specific growth \& maturity (after fisheries)
- pre-molt/post-molt size transition matrix
- size-specific probability of maturing on molt
- terminal molt to maturity
- spawning stock (MMB) assessed at mating, before growth



## New data type: male maturity ogives (based on chela heights)



## Male maturity ogives (based on chela heights)

- size-specific maturity ogives based on classifying new shell male crab by measured chela height-to-carapace width (CH:CW) ratios can be included in model likelihood
- eliminates need to classify male crab by maturity "outside" the model

$$
\begin{aligned}
& \mathrm{L}_{m}=\sum_{y, z} n_{y, z} \cdot\left\{p_{y, z}^{o b s} \cdot \ln \left(p_{y, z}^{\bmod }+\delta\right)+\left(1-p_{y, z}^{o b s}\right) \cdot \ln \left(1-p_{y, z}^{m o d}+\delta\right)\right\} \\
& p_{y, z}^{o b s}=\frac{n_{y, z, n s, m a t}^{o b s}}{n_{y, z, n s, m a t}^{o b s}+n_{y, z, n s, i m m}^{o b s}} \quad p_{y, z}^{m o d}=\frac{n_{y, z, n s, \operatorname{mat}}^{m o d}}{n_{y, z, n s, m a t}^{m o d}+n_{y, z, n s, i m m}^{m o d}}
\end{aligned}
$$

## Model Data Coverage



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## 2017 Assessment Model

| process | time blocks | description |
| :---: | :---: | :---: |
| Population rates and quantities |  |  |
| Population built from annual recruitment |  |  |
| Recruitment | 1949-1974 | In-scale mean + annual devs constrained as AR1 process |
|  | 1975-2017 | In-scale mean + annual devs |
| Growth | 1949-2016 | sex-specific |
|  |  | mean post-molt size: power function of pre-molt size post-molt size: gamma distribution conditioned on pre-molt size |
| Maturity | 1949-2016 | sex-specific |
|  |  | size-specific probability of terminal molt |
|  |  | logit-scale parameterization |
| Natural mortalty | 1949-1979, | estimated sex/maturity state-specific multipliers on base rate |
|  | 1980-1984 | estimated "enhanced mortality" period multipliers |
| Surveys |  |  |
| NMFS EBS trawl survey |  |  |
| male survey q | 1975-1981 | In-scale |
|  | 1982+ | In-scale w/ prior based on Somerton's underbag experiment |
| female survey q | 1975-1981 | In-scale |
|  | 1982+ | In-scale w/ prior based on Somerton's underbag experiment |
| male selectivity | 1975-1981 | ascending logistic |
|  | 1982+ | ascending logistic |
| female selectivity | 1975-1981 | ascending logistic |
|  | 1982+ | ascending logistic |

## 2017 Assessment Model

| Fishery/process | time blocks | description |
| :--- | :--- | :--- |
| TCF | directed Tanner crab fishery |  |
| capture rates | pre-1965 | male nominal rate |
|  | $1965-2016$ | male In-scale mean + annual devs |
|  | $1949-2016$ | In-scale female offset |
| male selectivity | $1949-1990$ | ascending logistic |
|  | $1991-1996$ | annually-varying ascending logistic |
|  | $2005-2016$ | annually-varying ascending logistic |
| female selectivity | $1949-2016$ | ascending logistic |
| male retention | $1949-1990,1991-$ ascending logistic |  |
|  | $1996,2005-2009$, |  |
|  | $2013-2015$ |  |
|  | bycatch in snow crab fishery |  |
| SCF | pre-1978 | nominal rate on males |
| capture rates | $1979-1991$ | extrapolated from effort |
|  | $1992-2016$ | male In-scale mean + annual devs |
|  | $1949-2016$ | In-scale female offset |
| male selectivity | $1949-1996$ | dome-shaped |
|  | $1997-2004$ | dome-shaped |
|  | $2005-2016$ | dome-shaped |
| female selectivity | $1949-1996$ | ascending logistic |
|  | $1997-2004$ | ascending logistic |
|  | $2005-2016$ | ascending logistic |
|  |  |  |

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## 2017 Assessment Model

| Fishery/process | time blocks | description |
| :--- | :--- | :--- |
| RKF | bycatch in BBRKC fishery |  |
| capture rates | pre-1952 | nominal rate on males |
|  | $1953-1991$ | extrapolated from effort |
|  | $1992-2016$ | male In-scale mean + annual devs |
|  | $1949-2016$ | In-scale female offset |
| male selectivity | $1949-1996$ | ascending logistic |
|  | $1997-2004$ | ascending logistic |
|  | $2005-2016$ | ascending logistic |
| female selectivity | $1949-1996$ | ascending logistic |
|  | $1997-2004$ | ascending logistic |
|  | $2005-2016$ | ascending logistic |
|  | bycatch in groundfish fisheries |  |
| GTF | pre-1973 | male In-scale mean from 1973+ |
| capture rates | $1973+$ | male In-scale mean + annual devs |
|  | $1973+$ | In-scale female offset |
| male selectivity | $1949-1986$ | ascending logistic |
|  | $1987-1996$ | ascending logistic |
|  | $1997+$ | ascending logistic |
| female selectivity | $1949-1986$ | ascending logistic |
|  | $1987-1996$ | ascending logistic |
|  | $1997+$ | ascending logistic |

## Alternative Model Scenarios

| model scenario | number of <br> parameters | description |
| :--- | :--- | :--- | :--- |
| 17AM (B2b) | 344 | 2017 assessment model |
| 17AMu | 344 | 17AM with updated crab fishery data |
| 18A (B0) | 357 | 17AMu with 2017/18 fishery data and 2018 NMFS survey data |
| 18B (B1) | 340 | 18A with fits to male maturity ogives. Reduced number of molt-to-maturity parameters <br> (17 fewer) |
| 18C0 (B2) | 340 | Fitting male maturity ogives, survey biomass by sex, size compositions for males by shell <br> condition and by maturity state and shell condition for females |
| 18C0a | 340 | 18C0, but reduced weight (/100) on fitting male maturity ogives |
| 18C1 (B4) | 340 | 18C1, but reduced weight (/100) on fitting male maturity ogives fitting survey abundanceby sex |

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## The "Underbag" Experiment




- Somerton and Otto, Fish. Bull.,1999
- Results used as priors on survey Q's
- 18C2a: fixes Q, selectivity
- mature males, immature females = "mixed sexes"
- mature females = "mature females"
- 18C3a: fixes Q, selectivity
- all crab = "mixed sexes"

Mature females


Author's Preferred Model: Support from BSFRF SBS data for fixing survey catchability/selectivity using the underbag experiment



## Fits to Data: 17AM, 17AMu, 18A

| Name | Component | Type | Distribution | Likelihood |
| :---: | :---: | :---: | :---: | :---: |
| 17AM, 17AMu, 18A | TCF: retained catch | abundance | -- | -- |
|  |  | biomass | norm2 | males only |
|  |  | size comp.s | multinomial | males only |
|  | TCF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | SCF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | RKF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | GTF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | NMFS survey | abundance | -- | -- |
|  |  | biomass | lognormal | by sex, for mature crab only |
|  |  | size comp.s | multinomial | by sex/maturity |
|  |  | chela height data | -- | -- |
|  | growth data | EBS only | gamma | by sex |

## Fits to Data: 18B

| Name | Component | Type | Distribution | Likelihood |
| :---: | :---: | :---: | :---: | :---: |
| 18B | TCF: retained catch | abundance | -- | -- |
|  |  | biomass | norm2 | males only |
|  |  | size comp.s | multinomial | males only |
|  | TCF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | SCF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | RKF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | GTF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | NMFS survey | abundance | -- | -- |
|  |  | biomass | lognormal | by sex, for mature crab only |
|  |  | size comp.s | multinomial | by sex/maturity |
|  |  | chela height data | binomial | for males |
|  | growth data | EBS only | gamma | by sex |

## Fits to Data: 18C0, 18C0a

| Name | Component | Type | Distribution | Likelihood |
| :---: | :---: | :---: | :---: | :---: |
| $\begin{aligned} & \text { 18C0, } \\ & 18 \mathrm{C} 0 \mathrm{a} \end{aligned}$ | TCF: retained catch | abundance | -- | -- |
|  |  | biomass | norm2 | males only |
|  |  | size comp.s | multinomial | males only |
|  | TCF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | SCF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | RKF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | GTF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | NMFS survey | abundance | -- | -- |
|  |  | biomass | lognormal | by sex, all crab |
|  |  | size comp.s | multinomial | males: by shell condition females: by maturity state, shell condition |
|  |  | chela height data | binomial | for males |
|  | growth data | EBS only | gamma | by sex |

## Fits to Data: 18C1, 18C1a, 18C2a, 18C3a

| Name | Component | Type | Distribution | Likelihood |
| :---: | :---: | :---: | :---: | :---: |
| $\begin{gathered} 18 \mathrm{Cl}, \\ 18 \mathrm{C} 1 \mathrm{a}, \\ 18 \mathrm{C} 2 \mathrm{a}, \\ 18 \mathrm{C} 3 \mathrm{a} \end{gathered}$ | TCF: retained catch | abundance | -- | -- |
|  |  | biomass | norm2 | males only |
|  |  | size comp.s | multinomial | males only |
|  | TCF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | SCF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | RKF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | GTF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | NMFS survey | abundance biomass <br> size comp.s | lognormal lognormal <br> multinomial | by sex, all crab by sex, all crab males: by shell condition females: by maturity state, shell condition |
|  |  | chela height data | binomial | for males |
|  | growth data | EBS only | gamma | by sex |

## Fits to Data: 18D0

| Name | Component | Type | Distribution | Likelihood |
| :---: | :---: | :---: | :---: | :---: |
| 18D0 | TCF: retained catch | abundance | -- | -- |
|  |  | biomass | norm2 | males only |
|  |  | size comp.s | multinomial | males only |
|  | TCF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | SCF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | RKF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | GTF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | NMFS survey | abundance | lognormal | -- |
|  |  | biomass <br> size comp.s | lognormal multinomial | by sex, all crab males: summe over shell cond. females: by maturity state |
|  |  | chela height data | binomial | for males |
|  | growth data | EBS only | gamma | by sex |

## Fits to Data: 18D1

| Name | Component | Type | Distribution | Likelihood |
| :---: | :---: | :---: | :---: | :---: |
| 18D1 | TCF: retained catch | abundance | -- | -- |
|  |  | biomass | norm2 | males only |
|  |  | size comp.s | multinomial | males only |
|  | TCF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | SCF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | RKF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | GTF: total catch | abundance | -- | -- |
|  |  | biomass | norm2 | by sex |
|  |  | size comp.s | multinomial | by sex |
|  | NMFS survey | abundance biomass size comp.s | lognormal lognormal multinomial | by sex, all crab by sex, all crab males: summed over shell cond. females: by maturity state |
|  |  | chela height data | binomial | for males |
|  | growth data | EBS only | gamma | by sex |

## Model Scenarios

| Model <br> Scenario | average recruitment millions | Final MMB 1000's t | B0 1000's t | Bmsy <br> 1000's t | Fmsy | $\begin{gathered} \text { MSY } \\ \text { 1000's t } \end{gathered}$ | Fofl | $\begin{gathered} \text { OFL } \\ \text { 1000's t } \end{gathered}$ | $\begin{gathered} \text { projected } \\ \text { MMB } \\ 1000 \text { 's t } \end{gathered}$ | projected MMB <br> / Bmsy |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 17AM (B2b) | 213.96 | 80.58 | 83.34 | 29.17 | 0.75 | 12.26 | 0.75 | 25.42 | 43.32 | 1.49 |
| 17AMu | 371.11 | 136.48 | 111.38 | 38.98 | 1.25 | 18.03 | 1.25 | 50.85 | 63.55 | 1.63 |
| 18A | 391.22 | 114.10 | 120.00 | 42.00 | 1.22 | 19.24 | 1.22 | 42.01 | 53.87 | 1.28 |
| 18B | 464.60 | 124.18 | 130.45 | 45.66 | 2.61 | 22.35 | 2.61 | 55.40 | 48.01 | 1.05 |
| 18C0 | 536.07 | 122.84 | 124.39 | 43.54 | 3.06 | 24.32 | 3.04 | 56.15 | 43.25 | 0.99 |
| 18COa | 366.37 | 99.63 | 100.92 | 35.32 | 1.07 | 18.13 | 1.07 | 35.44 | 46.25 | 1.31 |
| 18C1 | 540.64 | 128.64 | 129.28 | 45.25 | 2.79 | 25.90 | 2.78 | 58.26 | 45.12 | 1.00 |
| 18C1a | 404.67 | 110.14 | 109.74 | 38.41 | 1.14 | 20.41 | 1.14 | 39.87 | 49.67 | 1.29 |
| 18C2a | 199.49 | 50.12 | 63.01 | 22.05 | 0.91 | 11.54 | 0.91 | 16.76 | 24.06 | 1.09 |
| 18С3a | 188.34 | 49.93 | 63.61 | 22.26 | 0.79 | 10.84 | 0.79 | 15.93 | 25.44 | 1.14 |
| 18D0 | 503.62 | 145.40 | 149.02 | 52.16 | 2.64 | 24.09 | 2.64 | 65.30 | 57.35 | 1.10 |

18D1: did not converge

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## 2018 Model Scenario Results: Fits to Male Catch Biomass



## 2018 Model Scenario Results: Fits to "Other" Data

Growth Data


Maturity Ogive Data


## 2018 Model Scenario Results: Fits to NMFS Survey Biomass



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## 2018 Model Scenario Results: Fits to NMFS Survey Biomass



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## 2018 Model Scenario Results: Recruitment



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## 2018 Model Scenario Results: Mature Population Biomass




## 2018 Model Scenario Results: Population Abundance




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## 2018 Model Scenario Results: Survey Characteristics

Fully-selected Catchability
Capture Probabilities


## Average RMSEs for different fishery data components

| catch.type | data.type | fit.type | x | 18A | 18B | 18C0 | 18COa | 18C1 | 18C1a | 18C2a | 18C3a | 18D0 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| $\square$ total catch | Gabundance | ■BY_TOTAL | all sexes | 1.19 | 1.34 | 1.33 | 1.18 | 1.28 | 1.17 | 1.27 | 1.31 | 1.41 |
|  | $\square$ biomass | ■BY_TOTAL | all sexes | 0.06 | 0.06 | 0.06 | 0.07 | 0.06 | 0.07 | 0.09 | 0.10 | 0.06 |
|  | - $\mathrm{n} . \mathrm{at} . \mathrm{z}$ | GBY_XE | female | 374.07 | 370.62 | 392.96 | 401.93 | 386.04 | 394.18 | 390.27 | 378.12 | 364.70 |
|  |  |  | male | 371.14 | 318.81 | 313.14 | 342.07 | 313.58 | 352.02 | 310.12 | 313.03 | 332.45 |
| —total catch | @abundance | ■BY_X | female | 26.16 | 24.43 | 27.50 | 31.38 | 30.15 | 33.22 | 25.63 | 243.70 | 25.37 |
|  |  |  | male | 19.04 | 18.12 | 18.30 | 19.27 | 18.48 | 19.34 | 19.74 | 19.86 | 18.31 |
|  | $\square$ biomass | ■BY_X | female | 0.01 | 0.01 | 0.01 | 0.01 | 0.01 | 0.01 | 0.01 | 0.03 | 0.01 |
|  |  |  | male | 0.17 | 0.17 | 0.17 | 0.17 | 0.17 | 0.17 | 0.19 | 0.19 | 0.17 |
|  | En.at.z | ■BY_X | female | 50.27 | 49.28 | 50.43 | 51.18 | 49.65 | 49.89 | 51.16 | 42.02 | 53.59 |
|  |  |  | male | 67.04 | 67.88 | 67.46 | 64.88 | 68.12 | 66.73 | 65.41 | 64.38 | 69.35 |
| $\square$ total catch | Gabundance | ■BY_X | female | 12.21 | 12.73 | 12.31 | 13.38 | 11.70 | 12.71 | 11.12 | 14.30 | 12.20 |
|  |  |  | male | 2.71 | 2.69 | 2.68 | 2.70 | 2.70 | 2.71 | 3.01 | 2.95 | 2.67 |
|  | $\square$ biomass | ■BY_X | female | 0.09 | 0.09 | 0.09 | 0.08 | 0.09 | 0.08 | 0.05 | 0.07 | 0.09 |
|  |  |  | male | 0.07 | 0.07 | 0.07 | 0.07 | 0.07 | 0.07 | 0.08 | 0.08 | 0.07 |
|  | - n.at.z | ■BY_X | female | 69.38 | 71.22 | 69.80 | 68.30 | 70.41 | 69.35 | 72.79 | 74.39 | 69.30 |
|  |  |  | male | 346.62 | 351.28 | 341.70 | 333.22 | 311.33 | 309.99 | 270.11 | 280.42 | 361.13 |
| $\square$ retained catch | Elabundance | ■BY_X | female | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
|  |  |  | male | 3.99 | 3.98 | 4.04 | 4.06 | 4.06 | 4.03 | 4.46 | 4.50 | 3.94 |
|  | $\square$ biomass | ■BY_X | female | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
|  |  |  | male | 0.19 | 0.19 | 0.17 | 0.19 | 0.16 | 0.17 | 0.24 | 0.26 | 0.18 |
|  | - n.at.z | ■BY_X | male | 527.37 | 403.22 | 407.14 | 537.08 | 412.10 | 548.23 | 463.03 | 460.23 | 416.42 |
| $\square$ total catch | Eabundance | ■BY_X | female | 70.98 | 56.84 | 61.45 | 74.72 | 66.15 | 80.57 | 58.00 | 59.75 | 66.17 |
|  |  |  | male | 1.16 | 1.09 | 1.11 | 1.20 | 1.12 | 1.20 | 1.10 | 1.09 | 1.06 |
|  | Ebiomass | ■BY_X | female | 0.28 | 0.28 | 0.30 | 0.30 | 0.31 | 0.31 | 0.28 | 0.28 | 0.29 |
|  |  |  | male | 0.19 | 0.18 | 0.18 | 0.20 | 0.18 | 0.19 | 0.20 | 0.20 | 0.18 |
|  | En.at.z | ■BY_X | female | 184.36 | 185.71 | 192.13 | 189.51 | 199.10 | 196.16 | 205.96 | 201.38 | 187.44 |
|  |  |  | male | 346.02 | 413.43 | 410.06 | 337.85 | 405.06 | 334.33 | 317.20 | 309.97 | 406.67 |

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## Average RMSEs for different survey data components

| fleet | catch.type | data.type | fit.type | x | 18A | 18B | 18C0 | 18COa | 18C1 | 18C1a | 18C2a | 18C3a | 18D0 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| $\square$ NMFS (all by XM) | $\square$ index catch | Gabundance | ■BY_XM | female | 2.93 | 2.99 | 2.74 | 2.76 | 2.46 | 2.48 | 2.44 | 2.67 | 2.79 |
|  |  |  |  | male | 3.05 | 3.13 | 3.05 | 3.15 | 2.65 | 2.78 | 2.55 | 2.68 | 3.32 |
|  |  | $\square$ biomass | ■BY_X_MATONLY | female | 2.37 | 2.43 | 2.28 | 2.25 | 2.30 | 2.29 | 2.03 | 2.37 | 2.42 |
|  |  |  |  | male | 2.42 | 2.47 | 2.56 | 2.48 | 2.40 | 2.41 | 2.11 | 2.06 | 2.88 |
|  |  | En.at.z | EBY_XME | female | 425.44 | 400.73 | 400.38 | 403.24 | 317.22 | 335.14 | 414.03 | 226.70 | 370.93 |
|  |  |  |  | male | 456.14 | 520.94 | 513.95 | 393.65 | 495.45 | 388.14 | 323.05 | 324.67 | 518.19 |
| $\square$ NMFS (females by XM) | \#index catch | Gabundance | ■BY_X | female | 3.02 | 3.01 | 2.72 | 2.78 | 2.36 | 2.40 | 2.47 | 2.49 | 2.75 |
|  |  |  |  | male | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
|  |  | $\square$ biomass | ■BY_X | female | 2.48 | 2.50 | 2.30 | 2.31 | 2.22 | 2.23 | 2.05 | 2.26 | 2.40 |
|  |  |  |  | male | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
|  |  | \#n.at.z | ■BY_X_ME | female | 172.54 | 170.56 | 220.98 | 229.80 | 148.23 | 160.05 | 191.31 | 119.83 | 191.30 |
| $\square$ NMFS (females by XMS) | $\square$ index catch | Gabundance | ■BY_X | female | 3.02 | 3.01 | 2.72 | 2.78 | 2.36 | 2.40 | 2.47 | 2.49 | 2.75 |
|  |  |  |  | male | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
|  |  | $\square$ biomass | ■BY_X | female | 2.48 | 2.50 | 2.30 | 2.31 | 2.22 | 2.23 | 2.05 | 2.26 | 2.40 |
|  |  |  |  | male | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
|  |  | En.at.z | ■BY_XM_SE | female | 174.26 | 177.28 | 208.97 | 211.05 | 186.43 | 198.27 | 203.21 | 145.63 | 177.33 |
| $\square$ NMFS (males by X) | $\square$ index catch | $\square$ abundance | ■BY_X | female | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
|  |  |  |  | male | 3.48 | 3.48 | 3.38 | 3.39 | 2.76 | 2.82 | 2.77 | 2.86 | 3.51 |
|  |  | $\square$ biomass | ■BY_X | female | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
|  |  |  |  | male | 2.57 | 2.58 | 2.67 | 2.62 | 2.38 | 2.42 | 2.25 | 2.18 | 2.85 |
|  |  | En.at.z | ■BY_X | male | 203.11 | 189.35 | 191.12 | 201.79 | 189.09 | 193.18 | 159.00 | 154.78 | 191.06 |
| $\square \mathrm{NMFS}$ (males by XS) | ■index catch | छabundance | ■BY_X | female | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
|  |  |  |  | male | 3.48 | 3.48 | 3.38 | 3.39 | 2.76 | 2.82 | 2.77 | 2.86 | 3.51 |
|  |  | $\square$ biomass | ■BY_X | female | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
|  |  |  |  | male | 2.57 | 2.58 | 2.67 | 2.62 | 2.38 | 2.42 | 2.25 | 2.18 | 2.85 |
|  |  | En.at.z | ■BY_X_SE | male | 254.38 | 284.67 | 328.50 | 234.20 | 326.16 | 248.80 | 225.49 | 210.51 | 251.80 |

- "fleet" = dataset
- $X=$ sex, $M=$ maturity state, $S=$ shell condition


## Author's Preferred Model: 18C2a—best of the not very good?

- Fixes NMFS trawl survey catchability and selectivity based on Somerton and Otto (1999) "underbag" experiment
- BSFRF SBS data provides some support for this
- Provides reasonable fits to data
- Provides most reasonable characterization of stock productivity
- Fits to most dis-aggregated datasets
- male survey size compositions fit by shell condition
- female survey size compositions fit by maturity state and shell condition


## Author's Preferred Model: Comparison with 17AM

Feb. 15



Author's Preferred Model: Comparison with 17AM


## Author's Preferred Model: Comparison with 17AM

natural mortality
growth
$\operatorname{Pr}($ terminal molt $\mid$ size $)$




Author's Preferred Model: Comparison with 17AM


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## Author's Preferred Model: Comparison with 17AM


predicted observed
_ 17AM $\quad$ 17AM
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## Author's Preferred Model: MCMC Results



male
female

Status Determination \& OFL Calculations

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Status Determination \& OFL Calculation

- $\mathrm{F}_{\mathrm{MSY}}$
$=0.93 \mathrm{yr}^{-1}$
- mean recruitment
- $\mathrm{B}_{\mathrm{MSY}}$
= 199 million
- 2018/19 MMB-at-mating
- B/B $\mathrm{B}_{\mathrm{MS}}$
- Tier
$=21.87$ thousand t
$=23.53$ thousand $t$
$=1.08$
$=3 \mathrm{a}$



## Quad (Kobe) plot



## Management Reference Points

Basis for the OFL
(a) in 1000 's t .

| Year | MSST | Biomass <br> (MMB) | TAC <br> (East + West) | Retained <br> Catch | Total Catch <br> Mortality | OFL | ABC |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| $2014 / 15$ | 13.40 | $71.57^{\mathrm{A}}$ | 6.85 | 6.16 | 9.16 | 31.48 | 25.18 |
| $2015 / 16$ | 12.82 | $73.93^{\mathrm{A}}$ | 8.92 | 8.91 | 11.38 | 27.19 | 21.75 |
| $2016 / 17$ | 14.58 | $77.96^{\mathrm{A}}$ | 0.00 | 0.00 | 1.14 | 25.61 | 20.49 |
| $2017 / 18$ | $10.93^{\mathrm{C}}$ | $43.31^{\mathrm{A}}$ | 1.13 | 1.13 | $2.39^{\mathrm{C}}$ | 25.42 | 20.33 |
| $2018 / 19$ |  | $23.53^{\mathrm{BCC}}$ |  |  |  | $16.76^{\mathrm{C}}$ | $13.41^{\mathrm{C}}$ |

Management Performance
a) in 1000's t.

| Year | Tier $^{\mathbf{A}}$ | $\mathbf{B}_{\mathbf{M S Y}^{\mathbf{A}}}$ | Current <br> $\mathbf{M M B}^{\mathbf{A}}$ | ${\mathbf{B} / \mathbf{B}_{\mathbf{M S Y}} \mathbf{A}^{\mathbf{A}}}$ | $\mathbf{F o r f}^{\mathbf{A}}$ <br> $\left(\mathbf{y r}^{-1}\right)$ | Years to <br> define <br> $\mathbf{B}_{\mathbf{M S Y}}{ }^{\mathbf{A}}$ | Natural <br> $\mathbf{M o r t a l i t y ~}^{\mathbf{A}, \mathbf{B}}$ <br> $\left(\mathbf{y r}^{-\mathbf{1}}\right)$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| $2014 / 15$ | 3 a | 29.82 | 63.80 | 2.14 | 0.61 | $1982-2014$ | 0.23 |
| $2015 / 16$ | 3 a | 26.79 | 53.70 | 2.00 | 0.58 | $1982-2015$ | 0.23 |
| $2016 / 17$ | 3 a | 25.65 | 45.34 | 1.77 | 0.79 | $1982-2016$ | 0.23 |
| $2017 / 18$ | 3 a | 29.17 | 47.04 | 1.49 | 0.75 | $1982-2017$ | 0.23 |
| $2018 / 19$ | 3 a | 21.87 | 23.53 | 1.08 | 0.93 | $1982-2018$ | 0.23 |

## Future Directions

- incorporate BSFRF SBS index survey data
- develop more extensive MCMC diagnostics
- go back to basics (again): build model from ground up
- explore M / growth / maturity covariation
- explore size-dependent M
- determine appropriate likelihood weightings for growth, male maturity data
- start building Gmacs Tanner crab model
- disaggregate East/West directed fisheries in model

