## GOA rougheye and blackspotted rockfish (operational full)

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November 2023 Groundfish Plan Team
Nov doc, Sep doc, Sep presentation


## RE/BS Background

1. Cryptic species, no directed fishery, lightly exploited, catch << ABC == TAC
2. 2021 assessment: Tier 3a, no model changes since 2015, large retro bias (Mohn's rho=0.6), large drop in population scale and ABC
3. GPT/SSC/author recommended changes to biological assumptions, catchability, selectivity, data (index calculations, fleet structure, comp data), data weighting
4. In Sep 23, proposed staged approach to model dev, starting with new maturity, M, growth, and ageing error
5. Model did not respond well when new data was added in Oct, more work needed


|  |  |  |  |
| :--- | :---: | :---: | :---: |
| Year | Biomass $^{\mathbf{1}}$ | OFL | ABC |
| 2020 | 40,336 | 1,452 | 1,209 |
| 2021 | 40,432 | 1,456 | 1,212 |
| 2022 | 26,062 | 947 | 788 |
| 2023 | 25,957 | 937 | 781 |

## Risk Table Summary

## Recommend reduction from max ABC

## Author-recommended model was not reviewed in September

| Assessment | Population Dynamics | Ecosystem | Fishery |
| :---: | :---: | :---: | :---: |
| 2 - Major Concern | 2 - Major Concern | 1 - None | 1 - None |
| (Base model) <br> - Severe one-way positive retrospective bias <br> - High uncertainty in stock scale <br> (Recommended model) <br> - Improved stability, but poor fit and unable to account for recent declines in survey indices | - Declines in LLS and BTS indices in recent years <br> - 2023 LLS abundance lowest on record <br> - 2021 BTS lowest on record | - Average environmental conditions <br> - Some evidence of long-term declines in structural epifauna | - Incidental catch only <br> - Catch $\ll \mathrm{ABC}$ <br> - Not currently constraining target fisheries |

LLS = longline survey
BTS = bottom trawl survey

GOA GPT Sep 23 (+SSC): "The Team recommended using the authors approach. Additionally, the Team recommended alternative methods be explored that take skip spawning into account."

As discussed in Sep, we plan to address this recommendation in the next assessment.
FYI: it was noted during internal review that samples should be weighted by species-specific abundance. Currently, we assume sampling was proportional to abundance.


GOA GPT Sep 23 (+SSC): "Alternative methods that relied on a more precise prior were discussed such as computing a distribution based on available ages and applying the ageing error matrix to set the prior. The Team supported the author's investigation into $M$ but recommended the author explore the application of the prior variance used for M."

Current prior: Mean=0.03, CV=0.1
Sep 23: Mean=0.04 (oldest specimen=135 y), CV=0.31 (Hamel and Cope 2022) Analysis in response to recommendation: Using the 5 oldest survey and fishery GOA RE/BS specimens ( $126-135 \mathrm{y}$ ) and the age-specific SD from the new ageing error matrix (Punt et al. 2008), we constructed a distribution of Ms using the Hamel and Cope (2022) max age estimator with Mean $=0.042, \mathrm{CV}=0.058$ Nov 23: Author recommended model fixes $M$ at 0.042

SSC Oct 23 (+GPT): "The SSC supports the author and GOA-GPT recommendation to incorporate new data for the aging error matrix, the size-at-age matrix, and weight-at-age vector. "
In the absence of a strong recommendation to use WAA from a weight-based vonB (status quo) or a length-based vonB converted to weight using the weight-length relationship, we decided to remain with the status quo method in order to maintain consistency with the other Tier 3 GOA rockfish.



SSC Oct 21 (+GPT \& author): "...because the surveys exhibit inconsistent trends and partition biomass differently among areas, it is unclear if these signals reflect a genuine conservation concern or are the byproduct of survey data conflicts. The SSC concurs with the author and the GOA GPT that it would be prudent to estimate survey indices using the same depth strata definitions and to examine weighting CPUE by a variable other than total geographic area that may be more relevant to this complex (e.g., Essential Fish Habitat within a stratum)."
Author recommends no action on this topic at this time for the following reasons:
(1) After digging in more, this is a large task.
(2) EFH is defined using the trawl survey and is therefore not independent

See 2021 assessment for detailed CPUE comparisons by depth and area

SSC Dec 21: "The dome-shaped trawl survey selectivity... the GOA GPT noted it was unclear why 40-year-old fish would be so much less selected than a 30 -year-old fish. Future research could consider alternative parameterizations that would allow for more constrained estimates of selectivity at older ages.

Not addressed in this assessment, and it remains an issue. We welcome additional feedback or suggestions.


## Assessment Data Inputs

| Source | Data | Years |
| :---: | :---: | :---: |
| Fisheries | Catch | 1977-2021, 2022, 2023 |
|  | Age | $\begin{aligned} & 1990,2004,2006,2008,2009,2010,2012,2014 \\ & 2018,2020,2022 \end{aligned}$ |
|  | Length | $\begin{aligned} & \text { 1991-1992, 2002-2003, 2005, 2007, 2011, 2013, } \\ & 2017,2019,2021 \end{aligned}$ |
| AFSC bottom trawl survey | Biomass index | 1984, 1987, 1990, 1993, 1996, 1999, 2003, 2005, $2009,2011,2013,2015,2017,2019,2021,2023$ |
|  | Age | $\begin{aligned} & \text { 1984, 1987, 1990, 1993, 1996, 1999, 2003, 2005, } \\ & 2009,2011,2013,2015,2017,2019,2021 \end{aligned}$ |
| AFSC longline survey | Relative Population Number (RPN) | 1993-2019, 2020, 2021, 2022, 2023 |
|  | Length | 1993-2019, 2020, 2021, 2022, 2023 |

## New data in bold

## Abundance trends

LLS: 2023 lowest on record, $34 \%$ below mean

BTS: increase from 2021, which was the lowest on record, and 28\% below mean


## Demographic changes

## Evidence of declines in mean length and age in recent years

Fishery: at least

$\rightarrow$ BTS lengths
$\rightarrow$ Fishery lengths
$\rightarrow$ LLS lengths partially explained by shift away from hook-and-line gear


- BTS ages
$\rightarrow$ Fishery ages


## Alternative Model Configurations



## Selectivity and Comp Data Assumptions

Fishery marginal ages and length comps

- Age-based slx, nonparametric


## Longline survey

- Age-based slx, nonparametric

Bottom trawl survey

- Age-based slx, dome-shaped

Multinomial likelihood for comp data

- Age input $n$ : sqrt of total $n$
- Length input n : sqrt of total n scaled to a maximum of 100
- No data weighting algorithms used

- M15.4_2019
-- M15.4_2021

| Model | Mohn's rho | Key Results | $\begin{aligned} & 2024 \text { Age- } \\ & 3+ \\ & \text { Biomass* } \end{aligned}$ | $\begin{aligned} & 2024 \\ & \text { SSB }^{\star} \end{aligned}$ | $\begin{gathered} 2024 \\ \text { ABC }^{\star} \end{gathered}$ |
| :---: | :---: | :---: | :---: | :---: | :---: |
| Model 15.4 | 1.05 | - Severe positive retrospective bias in SSB <br> - Strong retro patterns and high parameter correlation among both $q$ 's, $M$, mean recruitment <br> - Unreasonably high estimates of trawl survey $q(>2)$ <br> - High reliance on length composition data | 29,081 | 9,642 | 794 |
| $\begin{gathered} \hline \text { Model } \\ 15.4 \mathrm{a} \end{gathered}$ | 1.16 | - Same as Model 15.4 but with even worse retro behavior | 27,574 | 9,245 | 751 |
| $\begin{gathered} \text { Model } \\ 23.1 \end{gathered}$ | 0.42 | - Bad retrospective bias in spawning biomass and strong retrospective trends in global scaling parameters <br> - Unreasonably high estimates of both q's (>2) <br> - Biomass scales that significantly lower than any model result to date <br> - Slight improvements in the fits to the index data | 16,154 | 3,890 | 432 |
| $\begin{aligned} & \text { Model } \\ & \text { 23.1a } \end{aligned}$ | 0.13 | - Greatly improved retro behavior, except for continued retro trends in $M$ <br> - Biomass scales that are consistent with Model 15.4 results before it started exhibiting retrospective patterns <br> - Degraded fits to index data in recent years <br> - Recent biomass trajectories are inconsistent with survey trends | 45,252 | 11,876 | 1,460 |
| $\begin{aligned} & \text { Model } \\ & \text { 23.1b } \end{aligned}$ | 0.14 | - Same as Model 23.1a but with no retrospective pattern in M | 46,129 | 13,022 | 1,305 |

## Summary of Results

| Model | Mohn's rho | Key Results | $\begin{gathered} 2024 \text { Age- } \\ 3+ \\ \text { Biomass* } \end{gathered}$ | $\begin{aligned} & 2024 \\ & \text { SSB }^{\star} \end{aligned}$ | $\begin{gathered} 2024 \\ \mathrm{ABC}^{\star} \end{gathered}$ |
| :---: | :---: | :---: | :---: | :---: | :---: |
| Model 15.4 | 1.05 | - Severe positive retrospective bias in SSB <br> - Strong retro patterns and high parameter correlation among both $q$ 's, $M$, mean recruitment <br> - Unreasonably high estimates of trawl survey $q(>2)$ <br> - High reliance on length composition data | 29,081 | 9,642 | 794 |
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## M15.4a

- Large declines in estimates of population productivity/scale (recruitment, M)
- High estimates of q (LLS_q=1.54, BTS_q=2.63)

Figure 13-9, Table 13-15

q_longline


## What are "reasonable" estimates of $q$ ?

$q<1.0$<br>Survey underestimates abundance<br>(e.g., untrawlable habitat)


$q>1.0$
Survey overestimates abundance (e.g., herding fish into trawl gear)

- Jones et al. 2021, Somerton et al. 1999, Krieger and Sigler 1996 - no mention of slope rockfish, all highlight the importance of size-selectivity
- No clear mechanism for high q of RE/BS in trawl or longline gear


## High parameter correlation

M15.4a pairwise plots of the MCMC marginal posterior distributions
(Figure 13-10)


## What data is informing scale?

## M15.4a likelihood profiles/Piner plots (Figure 13-14)



| Model | Mohn's <br> rho | Key Results | 2024 Age- <br> $3+$ <br> Biomass $^{\star}$ | 2024 <br> SSB $^{\star}$ | 2024 <br> ABC $^{\star}$ |
| :--- | :--- | :--- | :--- | :--- | :--- |

## What happens when we update the biological assumptions as proposed in Sep?

| $\begin{gathered} \text { Model } \\ 23.1 \end{gathered}$ | 0.42 | - Bad retrospective bias in spawning biomass and strong retrospective trends in global scaling parameters <br> - Unreasonably high estimates of both q's (>2) <br> - Biomass scales that significantly lower than any model result to date <br> - Slight improvements in the fits to the index data | 16,154 | 3,890 | 432 |
| :---: | :---: | :---: | :---: | :---: | :---: |
| $\begin{aligned} & \text { Model } \\ & \text { 23.1a } \end{aligned}$ | 0.13 |  | 45,252 | 11,876 | 1,460 |
| $\begin{gathered} \text { Model } \\ 23.1 \mathrm{~b} \end{gathered}$ | 0.14 | - Same as Mode | 46,129 | 13,022 | 1,305 |

M15.4a (drop 1980s BTS)

q_Iongline

q_trawl


LLS_q=1.54, BTS_q=2.63

M23.1 (just new bio assumptions)


NOAA FISHERIES

| Model | Mohn's rho | Key Results | $2024 \text { Age- }$ 3+ Biomass* |  | $\begin{gathered} 2024 \\ \mathrm{ABC}^{\star} \end{gathered}$ |
| :---: | :---: | :---: | :---: | :---: | :---: |
| Introducing scale into the RE/BS model: Constrained q_BTS and q_LLS Mean=1.0, CV=0.05 (BSAI BS/RE, Spencer et al. 2022) Fixed sigmaR at 1.1 (prior mean) |  |  |  |  |  |
| $\begin{aligned} & \text { Model } \\ & 23.1 \mathrm{a} \end{aligned}$ | 0.13 | - Greatly improved retro behavior, except for continued retro trends in $M$ <br> - Biomass scales that are consistent with Model 15.4 results before it started exhibiting retrospective patterns <br> - Degraded fits to index data in recent years <br> - Recent biomass trajectories are inconsistent with survey trends | 45,252 | 11,876 | 1,460 |
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## M23.1a

- Improved model stability, biomass estimates similar in scale to what has been estimated in the past
- Retrospective shift (increase) in estimates of population productivity (recruitment, M)
Figure 13-9, Table 13-15
log_mean_rec




| Model | Mohn's rho | Key Results | $\begin{aligned} & 2024 \text { Age- } \\ & 3+ \\ & \text { Biomass* } \end{aligned}$ | $\begin{aligned} & 2024 \\ & \text { SSB }^{\star} \end{aligned}$ | $\begin{gathered} 2024 \\ \text { ABC }^{\star} \end{gathered}$ |
| :---: | :---: | :---: | :---: | :---: | :---: |
| Model 15.4 | 1.05 | - Severe positive retrospective bias in SSB <br> - Strong retro patterns and high parameter correlation among both $q$ 's, $M$, mean recruitment <br> - Unreasonably high estimates of trawl survey $q(>2)$ <br> - High reliance on length composition data | 29,081 | 9,642 | 794 |
| $\begin{aligned} & \text { Model } \\ & 15.4 \mathrm{a} \end{aligned}$ | 1.16 | - Same as Model 15.4 but with even worse retro behavior | 27,574 | 9,245 | 751 |
| Model 23.1 | 0.42 | - Bad retrospective bias in spawning biomass and strong retrospective trends in global scaling parameters <br> - Unreasonably high estimates of both q's (>2) <br> - Biomass scales that significantly lower than any model result to date <br> - Slight improvements in the fits to the index data | 16,154 | 3,890 | 432 |
| Model 23.1a | 0.13 | - Greatly improved retro behavior, except for continued retro trends in $M$ <br> - Biomass scales that are consistent with Model 15.4 results before it started exhibiting retrospective patterns <br> - Degraded fits to index data in recent years <br> - Recent biomass trajectories are inconsistent with survey trends | 45,252 | 11,876 | 1,460 |
| $\begin{aligned} & \text { Model } \\ & 23.1 \mathrm{~b} \end{aligned}$ | 0.14 | - Same as Model 23.1a but with no retrospective pattern in M | 46,129 | 13,022 | 1,305 |

NOAA FISHERIES


- M15.4_2023
=- M15.4a_2023
- M23.1_2023
- M23.1a_2023
* . M23.1b_2023

Figure 13-1


| Model | MLE <br> Mean | Cl- <br> Lower | Cl- <br> Upper | MCMC <br> Mean | MCMC <br> Median | BCI- <br> Lower | BCl- <br> Upper |  |
| :--- | :--- | ---: | ---: | ---: | ---: | ---: | ---: | ---: |
| M15.4a_2023 | Log Rbar | 0.156 | -0.394 | 0.712 | -0.073 | -0.114 | -0.619 | 0.709 |
| M23.1_2023 | Log Rbar | 0.142 | -0.177 | 0.464 | -0.285 | -0.283 | -0.676 | 0.106 |
| M23.1a_2023 | Log Rbar | 0.934 | 0.611 | 1.261 | 0.581 | 0.579 | 0.268 | 0.898 |
| M23.1b_2023 | Log Rbar | 0.478 | 0.249 | 0.708 | 0.161 | 0.162 | -0.099 | 0.419 |
| M15.4a_2023 | M | 0.031 | 0.026 | 0.036 | 0.032 | 0.032 | 0.027 | 0.038 |
| M23.1_2023 | M | 0.044 | 0.039 | 0.049 | 0.044 | 0.044 | 0.039 | 0.049 |
| M23.1a_2023 | M | 0.052 | 0.047 | 0.056 | 0.052 | 0.052 | 0.048 | 0.057 |
| M23.1b_2023 | M | - | - | 0.042 | 0.042 | - | - |  |
| M15.4a_2023 | Longline q | 1.54 | 0.41 | 2.67 | 1.64 | 1.59 | 0.73 | 2.84 |
| M23.1_2023 | Longline q | 2.53 | 1.84 | 3.24 | 2.64 | 2.64 | 1.94 | 3.37 |
| M23.1a_2023 | Longline q | 1.00 | 0.90 | 1.10 | 1.01 | 1.01 | 0.92 | 1.11 |
| M23.1b_2023 | Longline q | 1.01 | 0.92 | 1.11 | 1.02 | 1.02 | 0.93 | 1.11 |
| M15.4a_2023 | Trawl q | 2.63 | 1.25 | 4.03 | 2.54 | 2.53 | 1.15 | 3.96 |
| M23.1_2023 | Trawl q | 2.72 | 1.95 | 3.50 | 2.70 | 2.69 | 1.96 | 3.52 |
| M23.1a_2023 | Trawl q | 1.04 | 0.93 | 1.14 | 1.03 | 1.03 | 0.94 | 1.12 |
| M23.1b_2023 | Trawl q | 1.06 | 0.97 | 1.16 | 1.06 | 1.05 | 0.96 | 1.15 |
| M15.4a_2023 | sigmaR | 0.78 | 0.69 | 0.88 | 1.04 | 1.04 | 0.92 | 1.17 |
| M23.1_2023 | sigmaR | 0.81 | 0.71 | 0.91 | 1.08 | 1.08 | 0.96 | 1.22 |
| M23.1a_2023 | sigmaR | 1.10 | - | - | 1.10 | 1.10 | - | - |
| M23.1b_2023 | sigmaR | 1.10 | - | - | 1.10 | 1.10 | - | - |

Table 13-15

Trawl survey age comps













## Top cohorts




Trawl survey length comps

Longline survey length comps



Fishery age comps




|  | 2018 |
| :---: | :---: |
|  | M15.4_2023 |
| - - | M15.4a_2023 |
| - | M23.1_2023 |
| - | M23.1a_2023 |
|  | M23.1b_2023 |





 3
 $\begin{array}{llllll}9 & 15 & 21 & 27 & 33 & 39\end{array}$ Age


Fishery length comps


OSA w/o length bin 1

## All around poor fits to comp data <br> Red = <br> overestimating <br> Blue = <br> underestimating

Figures 13-11-13-14


Pearson
60
|

## RE/BS Assessment Summary

- Base model M15.4 first accepted in 2015
- In Sep 2023, we presented new M, maturity, ageing error, and growth
- When new data were added, the retrospective bias went from bad in 2021 (Mohn's rho=0.61) to worse (Mohn's rho=1.05)
- Bias caused by large retrospective patterns in both survey catchabilities (q_trawl>2), M, and recruitment (all estimated in M15.4)
- Constrained scaling parameters in recommended M23.1b stabilized the model (Mohn's rho=0.14); however, M23.1b has degraded fits to the survey data and biomass trajectories that are inconsistent with recent trends in survey abundance
- More work needed to address model misspecification


## Recommended reduction from max ABC

| Quantity | As estimated or specified last year for: |  | As estimated or recommended this year for: |  |
| :---: | :---: | :---: | :---: | :---: |
|  | 2023 | 2024 | 2024 | 2025 |
| $M$ (natural mortality rate) | 0.034 | 0.034 | 0.042 | 0.042 |
| Tier | 3a | 3a | 3a | 3a |
| Projected total (ages 3+) biomass (t) | 25,837 | 25,755 | 46,029 | 46,109 |
| Projected female spawning biomass (t) | 8,554 | 8,514 | 12,986 | 13,005 |
| $B_{100 \%}$ | 14,776 | 14,776 | 21,878 | 21,878 |
| B40\% | 5,911 | 5,911 | 8,751 | 8,751 |
| B35\% | 5,172 | 5,172 | 7,657 | 7.657 |
| $F_{\text {OFL }}$ | 0.046 | 0.046 | 0.045 | 0.045 |
| maximabc | 0.038 | 0.038 | 0.038 | 0.038 |
| $F_{A B C}$ | 0.038 | 0.038 | 0.030 | 0.030 |
| OFL (t) | 930 | 927 | 1,555 | 1,566 |
| maxABC (t) | 775 | 772 | 1,302 | 1,310 |
| ABC (t) | 775 | 772 | 1,037 | 1,041 |
| Status | As determined last year for: $2021$ <br> 2022 |  | As determined this year for: |  |
|  |  |  | 2022 | 2023 |
| Overfishing | No | n/a | No | n/a |
| Overfished | n/a | No | n/a | No |
| Approaching overfished | n/a | No | n/a | No |

If this was a Tier 5 stock: $\mathrm{ABC}=889 \mathrm{t}, \mathrm{OFL}=1,185 \mathrm{t}$
Method: Split the difference between last year's and this year's projected ABCs

- A large increase from last year, but 20\% less than the 2010-2020 average ABC

| Year | OFL | ABC (=TAC) | Catch |
| :---: | :---: | :---: | :---: |
| 2010 | 1,568 | 1,302 | 426 |
| 2011 | 1,579 | 1,312 | 557 |
| 2012 | 1,472 | 1,223 | 599 |
| 2013 | 1,482 | 1,232 | 580 |
| 2014 | 1,497 | 1,244 | 760 |
| 2015 | 1,345 | 1,122 | 564 |
| 2016 | 1,596 | 1,328 | 697 |
| 2017 | 1,594 | 1,327 | 553 |
| 2018 | 1,735 | 1,444 | 795 |
| 2019 | 1,715 | 1,428 | 790 |
| 2020 | 1,452 | 1,209 | 398 |
| 2021 | 1,456 | 1,212 | 407 |
| 2022 | 947 | 788 | 469 |
| 2023 | 930 | 775 | $487^{*}$ |
| 2024 | 1,555 | 1,037 |  |

## Risk Table Summary

## Recommend reduction from max ABC

## Author-recommended model was not reviewed in September

| Assessment | Population Dynamics | Ecosystem | Fishery |
| :---: | :---: | :---: | :---: |
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| (Base model) <br> - Severe one-way positive retrospective bias <br> - High uncertainty in stock scale <br> (Recommended model) <br> - Improved stability, but unable to account for recent declines in survey indices | - Declines in LLS and BTS indices in recent years <br> - 2023 LLS abundance lowest on record <br> - 2021 BTS lowest on record | - Average environmental conditions <br> - Some evidence of long-term declines in structural epifauna | - Incidental catch only <br> - Catch <<ABC <br> - Not currently constraining target fisheries |

LLS = longline survey
BTS = bottom trawl survey

## RE/BS Apportionment Summary

- Two-survey random effects (REMA) model first accepted in 2019
- LLS scaling parameters fixed at 1.0 to balance LLS/BTS data conflict
- Recommend estimating area-specific scaling parameters, greatly improves model performance
- Recommend using the average area-specific proportions of REMA-predicted biomass and REMA-predicted relative population weights from the LLS in order to more appropriately balance BTS/LLS data

| Stock/ |  | $\mathbf{2 0 2 3}$ |  |  |  | $\mathbf{2 0 2 4}$ |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Sssemblage | Area | OFL | ABC | TAC | Catch $^{2}$ | OFL | ABC |
|  | W |  | 180 | 180 | 101 |  | 197 |
| RE/BS | C |  | 232 | 232 | 135 |  | 315 |
| complex | E |  | 363 | 363 | 149 |  | 525 |
|  | Total | 930 | 775 | 775 | 385 | 1,555 | 1,037 |



Apportionment based on predicted biomass and RPWs by area (PROPOSED)


## Planning for 2025+

1. RE/BS model development team
2. Model: q, selectivity, and recruitment
3. Refinement to maturity, survey indices,
 fisheries data, BTS length comps
4. Evaluation of uncertainty with MCMC
5. Continued organization, documentation, and modernization of code


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Fishery and survey data providers
Coauthors: J. Zahner, M. Siple, B. Ferriss
Former authors: K. Shotwell, D. Hanselman
Age and Growth: C. Gburski, J. Short, B. Matta
Genetic stock structure: W. Larson
Teammates: B. William, P. Hulson, C.
Monnahan, J. lanelli
Review: P. Spencer, C. Lunsford

