



**Abstract**—The phenomenon of multiple brooding, the production of more than one brood within a reproductive season, is a poorly understood process in rockfishes (*Sebastes* spp.). Most species produce a single brood, and all stock assessment models for *Sebastes* species assume the reproductive output (fecundity) of a single annual brood. We evaluated how well this life history trait could be predicted by both oceanographic (mean latitude, ocean temperature, dissolved oxygen levels, and depth) and demographic (maximum length, maximum age, and natural mortality rate) factors for 24 species of rockfish found on the continental shelf along the West Coast of the United States. Our results indicate that multiple-brooding species are considerably more common in lower latitudes (32–36°N) or in warmer waters (>9°C at depth) and that the other environmental and demographic factors analyzed were not empirically informative. This work provides a more comprehensive understanding of the reproductive ecology of shelf rockfishes, could contribute to assessments of climate change vulnerability, and should eventually lead to improvements in stock assessment models for economically and ecologically important rockfishes.

Manuscript submitted 20 October 2018.  
Manuscript accepted 19 July 2019.  
Fish. Bull. 117:180–188 (2019).  
Online publication date: 31 July 2019.  
doi: [10.7755/FB.117.3.5](https://doi.org/10.7755/FB.117.3.5)

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## An exploration of factors that relate to the occurrence of multiple brooding in rockfishes (*Sebastes* spp.)

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Population dynamics models that inform management decisions require an understanding of the reproductive biology of the fish species that are being managed. Rockfishes (*Sebastes* spp.) of the northeast Pacific Ocean tend to be long-lived species (maximum ages of some species are between 50 and 150 years) that can reach marketable size before maturity (Love et al., 1990). In addition to being long lived, slow growing, and late to mature, they have internal fertilization, are live-bearing, and usually develop a single brood annually (Love et al., 2002). Given these traits, which tend to be associated with high standing stock biomass and low productivity, rockfishes tend to be among the stocks that are more vulnerable to fishing pressure and presumably to future climate change (Parker et al., 2000; Berkeley et al., 2004; Cope et al., 2011).

Despite the report of multiple brooding, the production of more than one brood within a reproductive season, in some rockfishes over 50 years ago (Moser, 1966), multiple brooding continues to be poorly understood. Multiple brooding can be identified through

macroscopic or histological examination of the ovaries. In macroscopically examined ovaries, multiple broods can be identified when ovaries contain residual larvae or fertilized eggs (primary brood) and developing oocytes (upcoming secondary brood) (Beyer et al., 2015; Lefebvre et al., 2018). Lefebvre et al. (2018) identified multiple brooding through histological examination when ovaries contained post-ovulatory follicle complexes, residual larvae or eyed larvae (primary brood) and late-stage developing vitellogenic oocytes (upcoming secondary brood). Although many rockfish species have been identified as having the capacity to produce multiple broods (Moser, 1967; Love et al., 1990; Beyer et al., 2015; Lefebvre et al., 2018), the potential for multiple brooding is unknown for a great many remaining species for which little data on their reproductive ecology exists.

Traditional stock assessment metrics, such as estimates of spawning stock biomass, depend on the reproductive strategies and annual fecundity of species. When data are not available, assumptions are made that

can result in bias and misspecification of stock status and associated parameters. For example, if size-dependent fecundity cannot be quantified in an assessment but such a relationship exists in a real population, assessment models will be biased toward estimating a stock status that is more optimistic than actually exists (He et al., 2015a; Barneche et al., 2018). There has been a growing awareness in recent years that the biological processes that determine a population's productivity are much more complicated than previously understood. For example, some *Sebastes* species exhibit skipped spawning (Rideout et al., 2005; Conrath, 2017), prolonged adolescent periods (Thompson and Hannah, 2010), and multiple brooding (Moser, 1967; Love et al., 1990; Beyer et al., 2015; Lefebvre et al., 2018). Accounting for all of these factors is critical in the understanding of a species' life history and in specifying life history characteristics in stock assessment models, climate vulnerability assessments, and other evaluations.

Previous studies indicate that multiple brooding more commonly occurs in southerly distributed rockfish species (32–36°N), as well as in individuals residing in the southern extent of the range of their species (Love et al., 1990; Beyer et al., 2015). For example, bocaccio (*S. paucispinis*) range from the Southern California Bight to British Columbia, Canada; individuals in the Southern California Bight frequently exhibit multiple brooding and those in Central California occasionally exhibit multiple brooding, but the phenomenon has never been reported in individuals north of Cape Mendocino (He et al., 2015b). Multiple brooding has not been reported in northern stocks (Love et al., 2002; Conrath, 2017), nor in species typically found in deeper water (on the continental slope at depths >400 m), other than the bank rockfish (*S. rufus*) in Southern California. However, beyond these general observations, the phenomenon of multiple brooding has never been empirically evaluated in relation to environmental or life history factors, and we have yet to determine if multiple brooding can be robustly predicted on a species level to allow estimation of the implications to stock assessments and climate vulnerability assessments.

We sought to quantitatively document the relationship between the phenomenon of multiple brooding in West Coast rockfishes with latitude or other demographic (maximum length, maximum age, and natural mortality rate) and environmental (temperature and dissolved oxygen [DO] at depth, and average depth) factors and to determine which of these factors would be the best predictors of multiple brooding. Specifically, we hypothesized that rockfish species of the continental shelf of the West Coast of the United States that are able to produce multiple broods would be statistically more likely to inhabit lower latitudes (32–36°N) or warmer waters (>9°C at depth). The results of this research are intended to provide a more comprehensive understanding of the reproductive biology of rockfish species and to improve the information available for their sustainable management.

## Materials and methods

### Study area and data collection

Researchers at the Fisheries Ecology Division of the NOAA Southwest Fisheries Science Center have been collecting fecundity samples of a range of rockfishes to improve our understanding of size-dependent fecundity and carefully noting the presence or absence of multiple broods (Beyer et al., 2015; Lefebvre et al., 2018). On the basis of these efforts and previous reports (Moser, 1966, 1967; MacGregor, 1970; Wyllie Echeverria, 1987; Love et al., 1990; Ralston and MacFarlane, 2010), we classified 13 rockfish species found on the continental shelf of the West Coast as being multiple brooders and 11 shelf rockfish species as single brooders ( $n=24$ ; Table 1).

The NOAA Northwest Fisheries Science Center has conducted a fishery-independent groundfish bottom-trawl survey along the U.S. West Coast annually since 2003, in an area extending from the U.S.–Mexico border (latitude 32°30'N) to the U.S.–Canada border (latitude 48°10'N) and including depths from 55 m to 1280 m (Keller et al., 2012). We downloaded physical (trawl location, temperature, DO concentration at depth, and trawl depth) and biological (species and sex) data for shelf rockfishes (as defined in PFMC, 2016) for trawl tows conducted from 2004 through 2015 (FRAM Data Warehouse, Northwest Fisheries Science Center, available from [website](#)). It is important to note that the DO concentration at depth was not collected until 2007 and was not collected consistently until 2010. We downloaded commercial landings over the same time period from the Pacific Fisheries Information Network (PacFIN, available from [website](#)) and assigned port code latitudes based on port locations. We acquired the maximum length (i.e., asymptotic average fork length in centimeters; see Francis, 1988), maximum age (in years), and natural mortality rate of the species of interest from published stock assessments by the Pacific Fishery Management Council (available from [website](#)), Dick and MacCall (2010), and Love et al. (2002) (Table 1). Estimates of the natural mortality rate for shortbelly rockfish (*Sebastes jordani*) were used for halfbanded and pygmy rockfishes (*S. semicinctus* and *S. wilsoni*, respectively) because natural mortality had not been estimated for those 2 species and all 3 species are small, rapidly growing dwarf species. Species were excluded from our analyses if fewer than 20 individuals were collected over all the trawl samples; trawl samples were excluded if the tow of the trawl was conducted deeper than the depth range of continental shelf habitat (depths >400 m).

### Data analysis

To provide some context to our analyses, we explored how cumulative survey and commercial catch (measured as weight in kilograms) varied by latitude for each species by plotting the cumulative distribution of the survey and commercial catch data, and the arithmetic mean

**Table 1**

Multiple brooder classification (MB; Y=yes, N=no), mean latitude (in decimal degrees), maximum fork length ( $L_{\infty}$ , cm), observed maximum age ( $A_{MAX}$ , years), and natural mortality rate ( $M$ ) for each species of rockfish (*Sebastes* spp.) collected by the Pacific Fishery Management Council and NOAA Southwest Fishery Science Center between 2004 and 2015 on the continental shelf along the West Coast of the United States. For species for which parameters varied by sex, values for females are given.

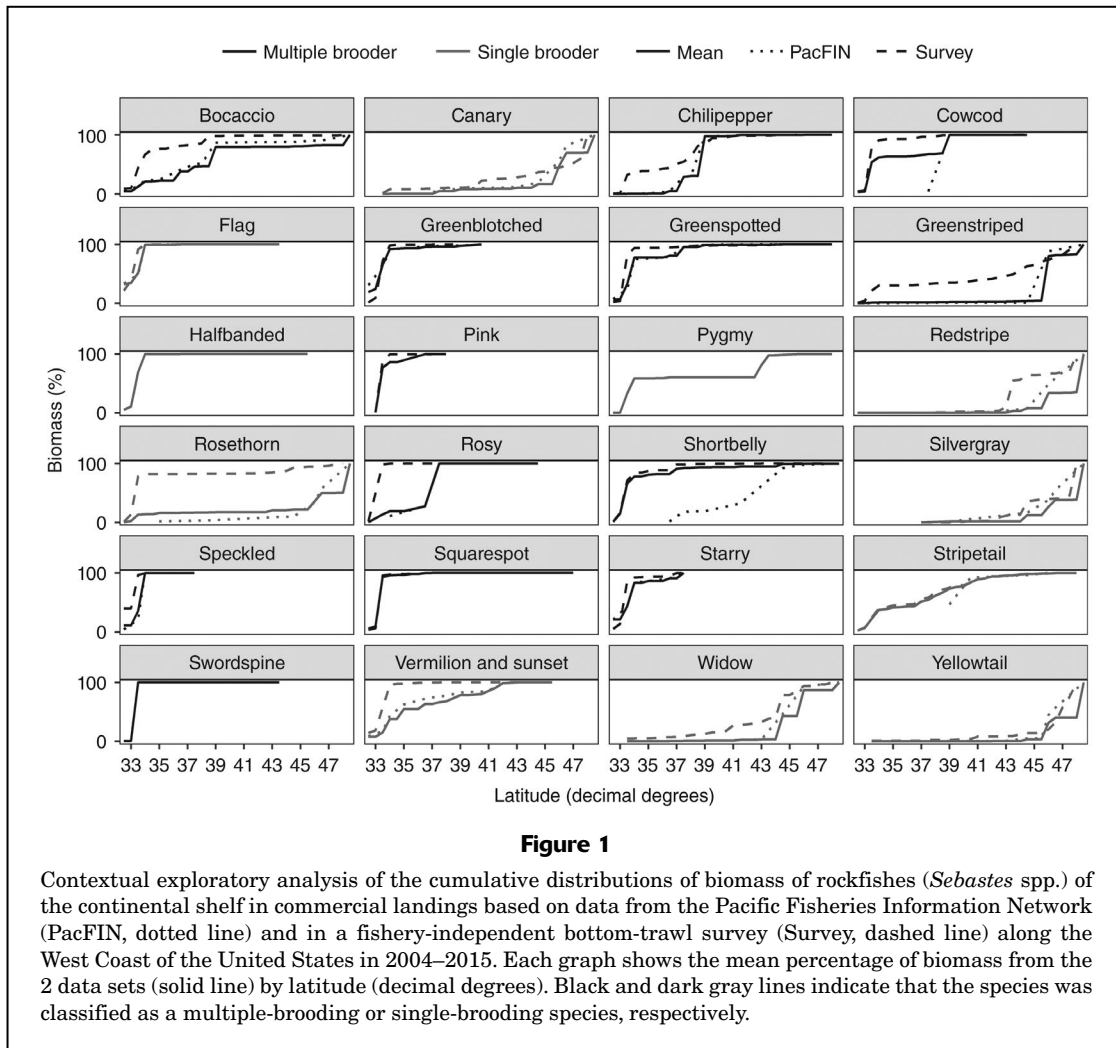
Common name	Scientific name	MB	Mean latitude	$L_{\infty}$	$A_{MAX}$	$M$
Bocaccio	<i>Sebastes paucispinis</i>	Y	35.87	75.90	37	0.15
Canary	<i>Sebastes pinniger</i>	N	44.78	62.00	84	0.05
Chilipepper	<i>Sebastes goodei</i>	Y	37.82	52.00	35	0.16
Cowcod	<i>Sebastes levis</i>	Y	35.50	87.00	55	0.06
Flag	<i>Sebastes rubrivinctus</i>	N	34.24	51.00	38	0.12
Greenblotched	<i>Sebastes rosenblatti</i>	Y	34.06	57.99	50	0.09
Greenspotted	<i>Sebastes chlorostictus</i>	Y	36.53	44.20	51	0.09
Greenstriped	<i>Sebastes elongatus</i>	Y	42.86	37.26	54	0.08
Halfbanded	<i>Sebastes semicinctus</i>	N	34.57	18.14	15	0.26
Pink	<i>Sebastes eos</i>	Y	34.00	56.00	66	0.07
Pygmy	<i>Sebastes wilsoni</i>	N	42.67	23.00	26	0.26
Redstripe	<i>Sebastes proriger</i>	N	45.53	61.00	55	0.08
Rosethorn	<i>Sebastes helvomaculatus</i>	N	43.78	28.66	87	0.05
Rosy	<i>Sebastes rosaceus</i>	Y	33.87	32.90	18	0.27
Shortbelly	<i>Sebastes jordani</i>	Y	36.00	28.50	32	0.26
Silvergray	<i>Sebastes brevispinis</i>	N	45.78	71.00	82	0.05
Speckled	<i>Sebastes ovalis</i>	Y	33.67	49.99	37	0.13
Squarespot	<i>Sebastes hopkinsi</i>	Y	34.82	25.25	19	0.26
Starry	<i>Sebastes constellatus</i>	Y	35.64	45.00	32	0.15
Stripetail	<i>Sebastes saxicola</i>	N	38.30	33.05	38	0.12
Swordspine	<i>Sebastes ensifer</i>	Y	33.76	17.60	43	0.11
Vermilion	<i>Sebastes miniatus</i>	N	34.73	62.40	60	0.07
Widow	<i>Sebastes entomelas</i>	N	42.90	50.34	60	0.08
Yellowtail	<i>Sebastes flavidus</i>	N	46.37	52.20	64	0.15

of both data sources, for each species. We then pooled data from the respective data sets over the study period (2004–2015) and summarized them by calculating the arithmetic mean to provide estimates of mean latitude (in decimal degrees), mean temperature at depth (in degrees Celsius), mean DO concentration at depth (in milliliters per liter), mean survey depth (in meters) for each species in the study ( $n=1$ , for each species). We fit binomial generalized linear models to the probability of each species being identified as a multiple brooder. The capacity to produce multiple broods was the response variable, and the environmental and demographic metrics were predictor variables.

Because temperature and latitude were likely highly correlated, we analyzed the collinearity between all explanatory variables using scatterplots and conducted a variance inflation factor (VIF) analysis. In the VIF analysis, we used an a priori cutoff of 10 (Craney and Surles, 2002) to assess whether collinearity between the variables was problematic. If variables had a VIF greater than 10, they were not used in the same model. Using the results of the VIF analysis, we created 23 candidate models that represented hypothesized potential predictor–response relationships. We compared models using an information-theoretic approach using Akaike information criterion corrected for

small sample sizes ( $AIC_c$ ) (Burnham and Anderson, 2002). Following the general method used by Kowalski et al. (2015), we computed the maximized log-likelihood,  $AIC$  and  $AIC_c$  scores,  $\Delta AIC_c$  values (difference between the  $AIC_c$  of model  $i$  and the smallest  $AIC_c$  among the considered models), and the AIC weight ( $AIC_w$ ) for each model. A priori, we decided to describe the level of empirical support for a model using the terms *substantial*, *considerably less*, and *essentially none* to correspond to  $\Delta AIC_c$  values less than 2, between 4–7, and greater than 10, respectively (as advised by Burnham and Anderson, 2002).

We evaluated the relative importance of influence of each predictor by summing the product of  $AIC_w$  and frequency of occurrence of the variable for all models. We computed the log evidence ratio (LER) in support for each predictor by calculating the  $\log_{10}$  of its evidence ratio (quotient of relative importance and its complement) and used LERs as the basis of inference for predictor comparisons. Following Kass and Raftery (1995), a priori, we decided to use the terms *minimal*, *substantial*, *strong*, and *decisive* to correspond approximately to LERs greater than 0, 0.5, 1, and 2, respectively. All analyses were executed by using R statistical software (vers. 3.4.4; R Core Team, 2018), including the *glm*, *AIC*, and *ggplot* functions of this software.



## Results

The results of exploratory analysis of the cumulative distribution of all rockfishes of the continental shelf that were present in the trawl survey's catch data reveal distinct differences in latitude between single- and multiple-brooding species and similar trends between commercial and survey data (Fig. 1). There appears to have been substantial differences in both latitude and temperature between single and multiple-brooding species and slight differences in DO, depth, and all demographic variables between single-brooding and multiple-brooding species (Table 2).

Unsurprisingly, latitude and temperature were highly correlated (VIFs: 71 and 67, respectively). To address multicollinearity issues, latitude and temperature were included in separate models before retesting all VIF scores, and they remained separate in the rest of the models. Of the 23 models considered, we found substantial support for those models that included latitude and found considerably less support for models that included temperature: (model 1 [M1]:  $\Delta AIC_c=0.0$ ; model 2 [M2]:  $\Delta AIC_c=0.7$ ;

and model 4 [M4]:  $\Delta AIC_c=3.2$ ; Table 3). There was marginal support for models that included some demographic variables (model 3 [M3]:  $\Delta AIC_c=2.7$ ; Table 3). The poorest performing models had essentially no support and were those for which latitude and temperature were separate (as required by the VIF results) but included all other variables (model containing all variables except temperature:  $\Delta AIC_c=11.6$ ; model containing all variables except latitude:  $\Delta AIC_c=14.1$ ; not shown).

Latitude and temperature were the strongest predictors of whether a shelf rockfish species could be a multiple brooder. There was substantial support for M1, which contained latitude and DO, and for M2, which contained only latitude, and there was considerably less support for M4 and M5, both of which include temperature (Table 3). When comparing the coefficients of M1 and M4, which contain DO and latitude (M1) or temperature (M4), the estimate for the latitude coefficient indicates that when DO is fixed and latitude increases by 1°, the odds of a shelf rockfish species being a multiple brooder is 0.6 times that of being a single brooder (for coefficients, see Table 3). Correspondingly,

**Table 2**

Comparison of the arithmetic mean and standard deviation (SD) of environmental and demographic variables between single- and multiple-brooding rockfish species (*Sebastes* spp.): latitude (decimal degrees), temperature (°C), dissolved oxygen (DO) concentration (mL/L), depth (m), maximum fork length ( $L_{\infty}$ , cm), observed maximum age ( $A_{MAX}$ , years), and natural mortality rate ( $M$ ). Data are provided for fish collected along the West Coast of the United States.

Variable	Single brooding		Multiple brooding	
	Mean	SD	Mean	SD
Latitude	41.24	4.83	35.72	2.49
Temperature	8.11	1.29	9.09	0.50
DO	1.94	0.44	1.91	0.46
Depth	157.44	28.18	161.35	34.33
$L_{\infty}$	46.62	17.96	46.89	19.70
$A_{MAX}$	55.36	23.96	40.69	14.19
$M$	0.12	0.08	0.14	0.08

**Table 3**

Top 5 generalized linear models as defined by the Akaike information criterion (AIC) analysis. Models were used to evaluate the probability of each rockfish species (*Sebastes* spp.) being a multiple brooder; all variables were continuous. The number of parameters in the model ( $K$ ), maximum log-likelihood (LL), AIC, AIC corrected for small sample sizes ( $AIC_c$ ), difference between the  $AIC_c$  of model  $i$  and the smallest  $AIC_c$  among the considered models ( $\Delta AIC_c$ ), and AIC weights ( $AIC_w$ ) are provided for each model (models 1–5 [M1–M5]). Variables include those for latitude (Lat), dissolved oxygen (DO) concentration, maximum fork length ( $L_{\infty}$ , cm), and temperature (Temp).

Model name	Model ( $\log(\hat{y}) =$ )	$K$	LL	AIC	$AIC_c$	$\Delta AIC_c$	$AIC_w$
M1	$25.27 - 0.51X_{Lat} - 2.78X_{DO}$	3	-9.95	25.91	27.11	0.00	0.39
M2	$13.32 - 0.35X_{Lat}$	2	-11.61	27.22	27.79	0.68	0.28
M3	$25.14 - 0.52X_{Lat} - 2.85X_{DO} + 0.01X_{L_{\infty}}$	4	-9.87	27.73	29.84	2.73	0.10
M4	$-10.55 + 1.98X_{Temp} - 3.25X_{DO}$	3	-11.54	29.07	30.27	3.16	0.08
M5	$-9.02 + 1.06X_{Temp}$	2	-13.68	31.36	31.93	4.82	0.04

the estimate for the temperature coefficient indicates that when DO is fixed and temperature increases by 1°C, the odds of a shelf rockfish species being a multiple brooder is 7.2 times that of being a single brooder. Finally, the results of evaluation of relative importance of influence of each predictor indicate that there was substantial evidence that species in lower latitudes had a higher probability of being a multiple brooder (LER for latitude=0.63; Table 4). There was minimal evidence indicating that any of the other variables affected the probability of a species being a multiple brooder (all other LERs were <0.5; Table 4).

## Discussion

Our results indicate that species with the capacity for multiple brooding were more likely to occur in lower latitudes (32–36°N) or in warmer waters (>9°C at depth).

This finding agrees with the established knowledge on differing distributions and community assemblages of rockfishes regionally (e.g., by latitude) (Love et al., 2002; Williams and Ralston, 2002; Gunderson and Vetter, 2006). Oceanographic conditions in waters of Southern California and Central California are characterized by warmer conditions and more variable, less seasonal upwelling patterns relative to conditions in the northern part of the California Current (e.g., north of Cape Mendocino), where ocean temperatures tend to be cooler and upwelling is both stronger and more strongly seasonal, conditions that result in higher primary and secondary productivity (Parrish et al., 1981; Ware and Thomson, 2005; Checkley and Barth, 2009). Consequently, rockfish species in waters of Southern California, where primary and secondary productivity and therefore presumably forage availability are both lower and characterized by greater interannual variability, may have an

**Table 4**

Summary of predictor variables considered in generalized linear models used to evaluate the probability of a rockfish species (*Sebastes* spp.) being a multiple brooder. Predictor variables include latitude (decimal degrees), temperature (°C), dissolved oxygen (DO) concentration (mL/L), depth (m), maximum fork length ( $L_{\infty}$ , cm), maximum age ( $A_{MAX}$ , years), and natural mortality rate ( $M$ ). Relative importance (RI), evidence ratio (ER), log evidence ratio (LER), and direction of effect (Direction) on the probability of a species being a multiple brooder for each predictor are shown.

Predictor variable	RI	ER	LER	Direction
Latitude	0.81	4.27	0.63	–
DO	0.64	1.79	0.25	–
$L_{\infty}$	0.18	0.21	–0.67	+
Temperature	0.15	0.17	–0.76	+
$A_{MAX}$	0.07	0.08	–1.10	–
$M$	0.03	0.03	–1.58	+
Depth	0.01	0.01	–2.19	+

evolutionary advantage in producing multiple broods over those species constrained to a single brood. The presumed mechanism for such an advantage would be to increase the probability of post-parturition larvae to encounter appropriate foraging conditions in this more variable, less seasonal region of the California Current. This is consistent with the notion of Cushing's match-mismatch hypothesis in which the effect of highly variable prey abundance for early life history stages of marine fish species may be partially mitigated by spreading spawning effort across a broader time period (Cushing, 1990).

Temperature affects individuals directly by influencing their growth and metabolism (Boehlert and Yoklavich, 1983; Boehlert et al., 1991). Temperature also influences egg and larval development; for example, recent laboratory studies have shown that multiple-brooding rosy rockfish (*S. rosaceus*) produced and released broods more quickly in warm water (relative to cool water) treatments, indicating faster development of eggs and larvae (Beyer<sup>1</sup>), and that faster development could result in the production of multiple broods during potentially narrow environmental windows for spawning. Temperature may tend to covary with other oceanographic variables that drive ocean productivity and consequently the availability of food resources, given that past studies have shown that the condition and reproductive output of some rockfish species can degrade during the warmer ocean conditions associated with El Niño–Southern Oscillation events (VenTresca et al., 1995; Harvey et al., 2011).

The importance of depth in this study is consistent with other studies that demonstrate that depth is a key driver of benthic community structure (Gunderson and Vetter, 2006; Tolimieri and Anderson, 2010). We did not include rockfish

species of the continental slope in this analysis because of our desire to focus on the multiple-brooding phenomenon, which is largely limited to species found on the continental shelf of the West Coast. However, we recognize that depth may more generally drive a broader range of reproductive strategies for rockfishes, such that species in deeper waters (>400 m) may respond to variable conditions by abortive maturity, skipped spawning, or atresia (Nichol and Pickett, 1994; Lefebvre and Field, 2015; Conrath, 2017), rather than the production (or lack thereof) of multiple broods. Although we did not formally evaluate the phylogeny of rockfish species relative to the capacity to produce multiple broods, it would appear that there are examples of both single and multiple brooders across the phylogenetic range of the genus *Sebastes* (Hyde and Vetter, 2007).

The similar predictive ability of temperature and latitude reflects the prominence of the influence of these variables on current probability of shelf rockfish species being multiple brooders. However, ocean conditions are changing. As the effects of climate change become more pronounced, the environmental conditions that are associated with certain regions (i.e., latitudes) will also change. Additionally, we did not find any evidence indicating that demographic variables are substantially associated with the phenomenon of multiple brooding. Although the maximum length and maximum age of species did not serve as a substantial predictor of multiple brooding among species, within species the occurrence of multiple brooding has been shown to increase with both size and age (Beyer et al., 2015; Lefebvre et al., 2018). Our results are also relevant to consideration of how observations of multiple brooding within a stock or species may vary with future warming or with the increasingly variable ocean conditions expected, and potentially already being observed, as a result of climate change (Ryckaczewski et al., 2015; McCabe et al., 2016). For example, Lefebvre et al. (2018) found that during

<sup>1</sup> Beyer, S. 2018. Unpubl. data. Dep. Ecol. Evol. Biol., Univ. Calif. Santa Cruz., 130 McAllister Way, Santa Cruz, CA 95060.

atypically warm years, when upwelling and forage conditions were above average, the frequency of occurrence of multiple brooding in chilipepper (*S. goodei*) was common and nearly equal in both Southern and Central California, whereas previous studies had not detected multiple broods in chilipepper of Central California (Wyllie Echeverria, 1987; Stafford et al., 2014). This finding might indicate that species capable of producing multiple broods do so more frequently under warmer ocean conditions when food resources are sufficient, potentially providing some advantage over single-brooding species in the face of warming trends.

Fisheries managers are beginning to prepare for the effects of climate change on fisheries by incorporating climate variables (such as ocean temperature trends) into harvest policies (Hill et al., 2017) and by conducting climate vulnerability assessments (Morrison et al., 2015; Hare et al., 2016). Climate vulnerability assessments evaluate which species will be at the greatest risk (or most vulnerable) to the effects of climate change, and these assessments consequently are intended to inform and prioritize research and management actions. Spawning frequency is 1 of 12 biological attributes used in the current criteria for evaluating vulnerability to climate change, with species that spawn in one single event per year considered to be more vulnerable than those that spawn several times per year (Hare et al., 2016). Therefore, multiple-brooding species may have an evolutionary advantage relative to single-brooding species in the face of either warming or more variable ocean conditions, necessitating an accurate assessment of the reproductive ecology of all *Sebastes* species in evaluating the vulnerability of different species to climate change.

Accounting for multiple brooding as a maternal effect in population models can substantially affect stock assessments, particularly if the likelihood of producing multiple broods (increasing overall reproductive output) is size dependent, as demonstrated for chilipepper (Lefebvre et al., 2018). Currently, stock assessment models of multiple-brooding species do not account for such factors in estimating reproductive output and stock status, potentially biasing assessment results if the effect is to nominally increase reproductive output of larger, older individuals. Taking multiple brooding into account in general, as well as in the context of climate change, will improve stock assessments and help managers develop appropriate management responses to climate variability and change now and into the future (Barneche et al., 2018). Our results provide robust predictions of this likelihood and provide context to the observed distribution of multiple-brooding species relative to single-brooding shelf rockfish species. However, our insights are not strictly mechanistic and are only the first step toward understanding the mechanisms and evolution of multiple brooding in rockfish species of the continental shelf. Future research should explicitly explore the mechanisms that drive multiple versus single brooding to aid in closing the knowledge gap for multiple-brooding rockfish species.

## Acknowledgments

We would like to thank S. Sogard, S. Beyer, L. Lefebvre, N. Kashef, and D. Stafford at the Fisheries Ecology Division of the Southwest Fisheries Science Center and the University of California's Cooperative Institute for Marine Ecosystems and Climate because their combined efforts were critical to inform and guide this effort. Additionally, we are deeply appreciative of the detailed comments provided by 3 anonymous reviewers and feel that incorporating their recommendations has greatly improved this manuscript.

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