Environment-driven trends in fish larval abundance predict fishery recruitment in two temperate

reef congeners: Mechanisms and implications for fishery recovery under a changing ocean

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Abstract

Environmental and biological processes acting on fish larvae can drive fishery cohort

strength, but predictive ability oftentimes falls short, and larval abundance is generally

considered more useful as a proxy for spawning biomass. Under a changing ocean, studies that

relate environmental covariates, larval abundance, and fishery recruitment are worthy of

continued research, especially in data-limited contexts. We focus on a popular, recreational-only,

multispecies saltwater bass fishery (genus *Paralabrax)* whose population status and recovery

potential are uncertain. We used 54 years of ichthyoplankton data (1963-2016) and a species

distribution model to 1) deconstruct species-specific standardized indices of larval abundance, 2)

test these indices as indicators of adult stock status or predictors of future fishery recruitment,

and 3) evaluate spatiotemporal trends in their population dynamics relative to environmental

variables. Contrary to expectation, species-specific larval abundance predicted future catch, with

recent elevated larval abundance suggesting imminent fishery recovery. Additionally, we

identified strong relationships with environmental variables, thereby providing additional tools

for predicting fishery recruitment and anticipating population change. Our findings paint a path

forward for improving estimates of current and future fishery status under changing natural and

anthropogenic influences and the incorporation of ecosystem considerations into fishery

management.

Keywords

Ecosystem-based management, data limited, species distribution model, fishery recovery,

recruitment processes, fishery cohort strength, environmental indicators

Introduction

An ecosystem approach to fishery management (Dolan et al. 2016) includes consideration

of the many biological and environmental factors that have historically, and are currently,

influencing fished populations. This, of course, also requires having information and resources to

adequately evaluate the status of fish populations. Unfortunately, many small-scale fisheries lack

such a robust approach, partly because data and resources are lacking. Given that the economic

and ecological impacts of some marine recreational fisheries have rivaled those of commercial

fisheries (Cooke & Cowx 2004, Lewin et al. 2019), and that overexploitation dampens the

ecosystem resilience of fish populations (Perry et al. 2010, Ziegler et al. 2023), there is a clear

need for research that leads to more robust population assessments for recreational-only and

small-scale commercial fisheries.

One of the most popular and economically important recreational-only marine fisheries in

California, USA, and perhaps even the world, is the southern California saltwater bass

Paralabrax spp. fishery, which consists of Barred Sand Bass *P. nebulifer* (BSB), Kelp Bass *P.*

clathratus (KB), and Spotted Sand Bass *P. maculatofasciatus* (SSB). This multi-species fishery

has persisted since the early 20th century, but catches have remained at historic lows since 2013

and are thought to primarily reflect a depressed population of BSB (a species targeted during

spawning aggregations; Jarvis et al. 2010, 2014a, Erisman et al. 2011). Dramatic catch declines

for BSB and KB began in 2005, following a period of increasing fishing mortality and

population recruitment failure (Jarvis et al. 2014a). In contrast, SSB is almost entirely catch-and-

release, limited to bays and estuaries, and presumed to have a healthier population. In 2013,

regulations for the saltwater basses became more restrictive (Cal. Code Regs. Tit. 14, § 28.30;

Jarvis et al. 2014a), but the fishery has not recovered, casting further uncertainty into the status

and recovery potential of the BSB and KB populations. Adding to that uncertainty is the

unknown impact that a warming ocean will have on these populations. Thus, research on the

population dynamics of the saltwater basses may help with anticipating change and aid in the

rebuilding and future management.

Differences in the geographic ranges, habitats, and reproductive strategies of the three

saltwater basses suggest different sensitivities to fishing pressure and environmental conditions,

and thus, differences in their resilience to climate change impacts. KB have occurred as far north

as the cool temperate coast of the state of Washington, USA, while BSB and SSB have a

maximum northern range extent that is approximately 1,000 km to the south, off the central

California coast (Love and Passerelli 2020). The southernmost occurrence for KB is southern

Baja California, Mexico, and for BSB and SSB, it is equatorward in the tropical waters of

Acapulco, Mexico; SSB also have populations in the Gulf of California (Heemstra 1995). As a

group, the saltwater basses have responded to decadal shifts in oceanographic conditions

throughout the history of the fishery, being relatively more abundant during warmer ocean

temperature regimes (Moser et al. 2001b, Hsieh et al. 2005, Jarvis et al. 2014a) and at one time,

more common north of southern California (Hubbs 1948). Therefore, it is possible that northern

latitudinal shifts in saltwater bass larvae abundances occur naturally, coinciding with decadal-

scale oceanographic cycles and climate-driven increases in sea surface temperature (SST, Auth

et al. 2018). Yet, at finer temporal scales, the saltwater basses may show differential population

responses associated with seasonal or interannual oceanographic variability, i.e., upwelling, El

Niño. Thus, understanding the relative role environmental variability has had on saltwater bass

populations with respect to their vulnerability to harvest, i.e., catch-and-release versus

aggregation-based, should help elucidate which species may be resilient to climate change

impacts.

Southern California is uniquely data-rich in terms of long-term coastal oceanography data

and thus represents an ideal testbed for the development of ecosystem-assessment methods. The

California Cooperative Oceanic Fisheries Investigations (CalCOFI) is the world's largest and

longest-standing fisheries oceanography survey (Gallo et al. 2019) and since 1951 has conducted

quarterly surveys to collect biological samples (e.g., fish larvae abundance, zooplankton

biomass) and hydrographic data at discrete sampling locations throughout the Southern

California Bight (SCB). Due to its long temporal coverage and high spatial resolution, the survey

has enabled researchers to detect natural and anthropogenic influences on larval fish populations

in southern California (Hsieh et al. 2009, Asch 2015, Thompson et al. 2017, 2022) and to assist

resource managers in fishery assessments (Moser et al. 2001b, McClatchie 2014, Gallo et al.

2019). Combined with other oceanographic and fishery data streams, CalCOFI provides the

unique opportunity to assess historical and recent spatiotemporal trends in saltwater bass

populations. Indeed, interest in parsing the CalCOFI *Paralabrax* spp. larval time series to species

funded research in the 1970s and 1980s to develop morphological and genetic tools for doing so

(Butler et al. 1982, Graves et al. 1990), but successful species discrimination across larval stages

was only recently made possible with the development of a robust, species-specific taxonomic

key (Jarvis Mason et al. 2022).

Larval abundance has long been considered a proxy for spawning stock biomass (Hilborn

& Walters 1992, Cowan & Shaw 2002). Indeed, CalCOFI fish larvae have been incorporated in

west coast fishery stock assessments or used for monitoring status and trends (see Gallo et al.

2019 for examples). On the one hand, if we assume bass larval abundance is a function of the

biomass of reproductive females in the population, then fishery managers are provided with an

easy, reliable means of tracking the status and trends of adult stocks. On the other hand, larval

abundance can be used to predict fishery recruitment in some fish (Murphy et al. 2018, Tripp et

al. 2023); that is, since larval fish dynamics are so highly influenced by environmental

variability, the processes affecting larval growth and mortality are thought to drive future fishery

year class strength (Hjort 1914, Lasker 1984, 1987, Cushing 1990, Houde 2001). However, due

to continued abiotic and biotic pressures on subsequent life stages, the accuracy of recruitment

prediction from larval indices is generally considered to be better for species with lower ages of

recruitment to fisheries (e.g., the juvenile or subadult stage, Bradford 1992, Cowan & Shaw

2002). Jarvis et al. (2014a) reported strong relationships between combined *Paralabrax* spp.

larval abundance and species-specific fishery recruitment strength between 1997 and 2012, but it

is unknown whether this relationship translates using species level larval abundance over a

longer period.

115   Here, we unlock the CalCOFI *Paralabrax* spp. larval archive for the first time. In doing

116   so, we intend to,

117   (I)   generate indices of southern California bass larvae abundance to improve species-

118   specific estimates of population status and trends,

119   (II)   test our indices as indicators of adult stock status or predictors of future fishery

120   recruitment, and

121   (III)   evaluate spatiotemporal trends in bass larval abundance relative to environmental

122   variables and climate forcing (e.g., latitudinal shifts in abundance).

In the face of a changing ocean, the ability to identify potential environmental indicators

of population status or future fishery recruitment, i.e., potential management action triggers, will

be ever more important for guiding sustainable fishery management, especially in data-limited

contexts, and may be a key determinate in the fate of this historical fishery.

Methods

Survey and study region

We focused on *Paralabrax* spp. larvae collected in CalCOFI oceanographic cruises

conducted along the continental shelf off southern California, USA and Baja California, Mexico

during July cruises from 1963 to 2016 (Fig. 1a, Table 1) because: 1) July corresponds to peak

spawning (Walker et al. 1987), 2) *Paralabrax* spp. larvae occur nearshore relative to most other

economically important groundfish in the SCB (McGowen 1993, Moser et al. 2001a), and, 3) at

the time of this study, the CalCOFI ichthyoplankton archive was being updated in reverse

chronological order to current taxonomic standards (1963 and 2016 were the most historical and

recent years available, respectively). Two changes to the CalCOFI zooplankton sampling

occurred during this period, 1) in 1969, the depth from which nets were obliquely towed

increased from 140 m to 210 m (or from within 5 m of the seafloor in shallower waters), and 2)

in 1978, the sampling gear changed from ring nets to paired bongo nets (Thompson et al. 2017).

Survey coverage has varied spatially and temporally, with more consistent temporal coverage in

the SCB (Fig. 1a) and additional, shallower, coastal stations added in 2004 (Gallo et al. 2019).

The study region is part of the California Current Ecosystem, in which the offshore

California Current (CC) brings cool, nutrient-rich waters equatorward from north of Pt.

Conception, CA (34.2°N), before branching shoreward toward the U.S.A./Mexico border

(32.2°N), and then poleward along the southern California coast, forming the counterclockwise

southern California Eddy within the SCB in the summer months (Fig. 1b, Hickey 1993,

McClatchie 2014). The southern branch of the CC continues to flow equatorward before veering

westward offshore of Baja California Sur. Thus, the study region generally consists of cooler,

temperate waters to the north and warmer, subtropical waters to the south (Fig. 1b, McClatchie

2014).

Larval species identifications and data processing

Two taxonomists sorted and processed *Paralabrax* spp. larvae from archived CalCOFI

cruise station vials. We assigned the larval stage of each larva according to notochord

development and then assigned species using pigmentation patterns described in Jarvis Mason et

al. (2022). We excluded yolk sac larvae as these could not be reliably identified, and we

excluded *P. auroguttatus* (Goldspotted Sand Bass) because its population is rare north of Baja

California Sur (Moser et al. 1996). We photographed all larvae and recorded the

presence/absence of stage-specific morphological characters described in Jarvis Mason et al.

(2022). For quality assurance, we selected a random, temporally stratified, subset of samples

(10%) to be identified by both taxonomists. We reviewed and resolved all identifications with

taxonomic disagreement. Four station vials contained hundreds of *Paralabrax* spp. larvae, and

we subsampled at least 10% of the contents of these vials and then extrapolated the raw species

counts based on the proportion of each species identified in the subsample, including any

unidentified or yolk sac larvae. The two taxonomists regularly collaborated to resolve

outstanding, difficult-to-identify larvae.

To characterize the overall sample of visually identified *Paralabrax* spp. larvae, we

summarized the raw numbers of larvae by species, larval stage, and region (southern California,

Baja California) and we plotted their geographic distributions. For subsequent analyses, we

standardized species counts in each sample by the standard haul factor to obtain the number of

larvae under ten m^2 of sea surface (Thompson et al. 2017). We refer to this standardized raw

count as larval abundance. There were 26 samples from 12 cruises in which larvae were so

damaged as to not be identifiable to species or there was a discrepancy in the number of

Paralabrax spp. larvae reported in the database versus the number observed in the sample vial

(e.g., missing larvae). In this case, we first rechecked the original station vials and reference

collection for the missing larvae, and if not found, then we estimated larval abundance (the

standardized counts) for the sample based on the average proportion of each species standardized

count at that station over the course of the entire study period.

Standardized index of larval abundance

We used species distribution models (SDMs) to generate standardized indices of bass

larval abundance in southern California. We did not include larval abundance data from Baja

California because July CalCOFI did not survey the entire Baja California region after 1980. We

used a geostatistical generalized mixed effects model framework that can incorporate spatial and

spatiotemporal random fields (effects) to account for latent variables that may contribute to

observations being correlated in space and time. The spatial random field captures latent spatial

variation that is unchanging through time (e.g., distance to shore), while the spatiotemporal

random field captures latent time varying spatial patterns (e.g., dynamic biological and

oceanographic processes not captured by covariates in the model). This type of SDM can

therefore account for unbalanced sampling effort and thus, is more precise than other SDM

methods at estimating abundance (Thorson et al. 2015, Brodie et al. 2020). We fit SDMs with the

R package `sdmTMB`, which estimates parameters of the Gaussian random fields using the

Integrated Nested Laplace Approximation (INLA) and implements maximum marginal

likelihood estimation with Template Model Builder (TMB, Anderson et al. 2022).

We modeled bass larval abundance as a function of variables that may affect

"catchability" but not abundance (e.g., when and where you survey, gear type). For both species,

we considered year, moon phase, day of year, hour, day/night by net type (to account for gear

changes from ring to bongo nets, Thompson et al. 2017), and distance to the mainland coast. We

did not consider the increase in tow depth because southern California *Paralabrax* spp. larvae

are most abundant inshore of the 36 m depth contour (Lavenberg et al. 1986). We modeled bass

larval abundance *Y* at location *s* and time *t* using a Tweedie observation error family (positive

continuous density values that also contain zeros, Shono 2008) and a log link:

$$\ln(\mu_{s,t}) = \mathbf{X}_{s,t}\boldsymbol{\beta} + \omega_s + \epsilon_{s,t},$$

$$\omega_s \sim \boldsymbol{MVNormal}(\mathbf{0}, \textstyle\sum_{\omega_s}),$$

$$\epsilon_{s,t} \sim \boldsymbol{MVNormal}(\mathbf{0}, \textstyle\sum_{\epsilon_{s,t}}), \text{ where,}$$

$\mu_{s,t}$ represents mean abundance at location *s* and time *t*,

$\mathbf{X}_{s,t}$ is a vector of main effects, time varying effects, or spatially varying effects

corresponding to location *s* and time *t*,

$\boldsymbol{\beta}$ is a vector of estimated parameters, and

ω_s and $\epsilon_{s,t}$, are the spatial and spatiotemporal random effects, respectively, assumed to be

drawn from a multivariate normal distribution with covariance matrices $\sum_{\omega_s}$ and $\sum_{\epsilon_{s,t}}$. A user-

specified triangulated mesh approximates the spatial component of the Gaussian random fields

(Supplement S1, Anderson et al. 2022)

Given data gaps for some years, we could not model year as a fixed effect (factor).

Instead, we modeled year effects and all other covariate effects with penalized smooth functions

(penalized regression "P-splines"; Eilers and Marx 1996), similar to generalized additive models,

GAMs, Wood 2017). We allowed the model to select the basis dimension, i.e., k, or wiggliness

for each variable, except for the hour and continuous numeric moon phase covariates, in which

we specified a cyclic smooth over 24-h and 4-w cycles, respectively. To account for potential

day time net avoidance related to the gear change from ring to bongo nets, we specified

day/night by net type as a factor smooth interaction where a separate smooth is created for each

factor level. In cases of non-convergence, we tried simplifying variables to their linear form.

Using Restricted Maximum Likelihood (REML)-generated Akaike (AIC) values, we

identified the best fit random effects model structure as an independent spatiotemporal random

field across time steps (both species) and a spatial field estimated for BSB but not for KB. Using

these species-specific random effects, we then compared model fit across different fixed effects

(= main effects) with AIC (REML off). To allow for subsequent prediction of larval abundance

in years without data, we specified missing years for model estimation of larval abundance. In

the case of BSB, in which the inclusion of a smooth year effect was not supported, we

incorporated year effects as a time-varying random walk (analogous to a dynamic linear model).

We did not consider models that did not pass all model diagnostic checks (e.g., tests for

convergence, positive definite Hessian matrix, large standard errors associated with model

parameter estimates).

To generate an area-weighted standardized index of larval abundance, we used our model

to predict abundance across a fine-scale grid covering the southern California CalCOFI survey

region, with a grid cell resolution of 2 km x 2 km, focusing only on years when larvae were

observed. This was intended to provide an index with less uncertainty across years. However, for

relating larval abundance to catch data (see Relationship with fishery catch data below), in which

missing years are not allowed, we generated the index of abundance based on the fine-scale

prediction grid with missing years included, i.e., abundance was predicted for missing years.

Spatiotemporal trends in larval abundance

To explore changes in larval abundance through time, we plotted spatiotemporal trends in

larval abundance of both species using predicted annual abundances defined by a 2 km x 2 km

resolution grid (exclusive of land and within the southern California CalCOFI survey area) and

the parameter estimates generated from the index standardization model; missing years not

included (Anderson et al. 2022). These trends included both fixed and random effects. We also

separately plotted trends in the spatiotemporal random effects ($\epsilon_{s,t}$) to visually explore what

patterns in the plots of fixed and random effects were being driven by latent spatiotemporal

processes.

Relationship with fishery catch data

For both species, we tested for a positive monotonic correlation between the index of

larval abundance and spawning stock biomass, in which we used catch records from California

Department of Fish and Wildlife (CDFW) Commercial Passenger Fishing Vessel (CPFV)

logbooks (1975-2016) as a proxy for spawning stock biomass (Fig. S2, see Jarvis et al. 2014a for

a description of catch data). The catch data represent the only index of saltwater bass abundance

that matches the geographic extent of the CalCOFI larval abundance data. Typically, catch-per-

unit-effort (CPUE) is considered an appropriate catch metric to use as an index of abundance

because it accounts for changes in fishing effort over time. However, in this fishery, catch and

CPUE directly correspond (Jarvis et al. 2014a) and species-specific CPUE at the trip level is not

available prior to 1980. CPFV catch records reflect only fish kept (harvested catch) on

commercial sportfishing vessels, the predominant fishing mode for BSB and KB (Jarvis et al.

2014a).

We considered only biologically plausible lags, i.e., catch cannot lead larvae, from zero

to ten years and standardized all data to a mean of zero. We performed cross-correlation analysis

using the R package `funtimes` (Lyubchich et al. 2023) in which a 95% confidence band

provided guidance for interpreting correlations that may be influenced by the presence of

autocorrelation in either or both datasets, i.e., coefficients falling within the band may be an

artifact of non-stationarity.

We also explored the relationship between species-specific larval abundance estimates

and estimates of *total* catch, i.e., estimates of harvest plus catch released alive or dead, by all

fishing modes combined (e.g., CPFVs, private/rental boats, shore fishing). These data are

available from the Marine Fisheries Statistical Survey (MRFSS, 1980-2003) and the California

Recreational Fisheries Survey (CRFS, 2005-2016; PSMFC 2024). Estimates are derived from

angler-intercept surveys of catch on a subset of fishing trips and extrapolated based on a phone

survey of effort. MRFSS and CRFS are calculated using a different sampling design; however,

since we were only concerned with monotonic relationships between larvae and catch, i.e., an

increase in one variable predicts an increase (or decrease) in the other, we combined the time

series and standardized values to a mean of zero (Fig. S2). Given that cross-correlation analysis

does not allow for missing values, we imputed catch for four missing years in the total catch

estimates for both species using the R package `mice` and the Predictive Mean Matching (PMM)

method (van Buuren & Groothuis-Oudshoorn 2011).

We assumed a positive correlation between larvae and catch at a lag of 0 years to indicate

a positive relationship between spawning stock biomass and larval abundance, where larval

abundance is a function of the biomass of reproductive age females in the water, i.e., the higher

the spawning stock biomass, the higher the larval abundance.

Relationship with environmental variables

We also used the R package `sdmTMB` to model the influence of site-specific prey

availability and temperature, as well as larger-scale oceanographic indices on bass larval

abundance. For these models including environmental data, we specified a minimum distance

cutoff of ten km in constructing the mesh.

Not all environmental variables of interest were available as far back as 1963, so we first

modeled a smaller set of variables across the entire time series (1963 and 2016, Model 1) and

then incorporated additional variables over the shorter period (1984 to 2016, Model 2). In

addition, data available for modeling bass larval abundance was limited to those years for which

a July CalCOFI cruise took place. For example, July CalCOFI cruises occurred triennially

between the mid-1960s and mid-1990s, otherwise, July temperature and zooplankton data were

available annually (Fig. 2). The larger scale environmental variables, North Pacific Gyre

Oscillation (NPGO) index and Oceanic Niño Index (ONI), were also available annually, but the

Biologically Effective Upwelled Transport Index (BEUTI), Coastal Upwelling Transport Index

(CUTI), and isothermal layer depth (ILD) for southern California were only available from 1980

onward, while kelp canopy extent was available from 1984 onward (Fig. 2).

For both species in Model 1, we included the station-specific CalCOFI temperature (°C,

averaged across the upper ten meters) and square root transformed zooplankton biomass

measured as zooplankton displacement volume (cm^3/1000 m^3 strained). We also included the

NPGO index for the month of July and the ONI for June/July, and distance to the mainland coast

(km, Fig. 2). The NPGO index is based on sea surface height variability throughout the north

Pacific and tends to track regional nutrient fluctuations (or, mechanisms driving plankton

ecosystem dynamics; Di Lorenzo et al. 2008). The ONI is based on surface temperature

anomalies in the east-central equatorial Pacific and defines El Niño Southern Oscillation (ENSO)

states (NOAA 2023).

For both species in Model 2, we included all variables from Model 1, and two additional

variables: an upwelling index and the ILD in meters. We included upwelling because we wanted

a more local measure of potential nutrient flux (versus the larger-scale NPGO), and we included

ILD because spawning BSB orient to the thermocline during spawning season (McKinzie et al.

2014). In the study region, the ILD roughly corresponds to the mixed layer depth (MLD), which

is the depth where temperature differs by 0.5 °C from the sea surface, i.e., we assume no salinity

barrier layer between the MLD and ILD. We used a mean Bakun upwelling index (m^3/sec) for

June/July at latitude 33°N, where positive values indicate wind-driven offshore transport (Bakun

1973).

For both species, we considered another candidate Model 2 where we replaced the Bakun

index with the BEUTI and CUTI specific to the southern California region, which provide a

measure of nitrate flux through the mixed layer and coastal vertical transport, respectively (Jacox

et al. 2018a, data from Hunsicker et al. 2022). Positive BEUTI values indicate nitrate flux into

the mixed layer and positive CUTI values indicate coastal upwelling. For BSB, we additionally

substituted upwelling in the SCB with upwelling off northern Baja California (Bakun index at

latitude 30°N) because older recruits (~ 2-y) showed a negative correlation with upwelling south

of the SCB at a 2-y lag (Jarvis et al. 2004). For KB, we also considered a model that substituted

upwelling in the SCB with kelp canopy extent because Giant Kelp *Macrocystis pyrifera* is

important adult habitat (White & Caselle 2008). We obtained regional areal kelp canopy data by

year from kelpwatch.org. Additionally, for both bass species, another candidate model

considered (in place of zooplankton biomass) biomass of the calanoid copepod, *Calanus pacifus*,

in spring CalCOFI cruises as a potential measure of larval pre-conditioning (Fennie et al. 2023,

Swalethorp et al. 2023, Walsh 2023).

We used AIC to identify the most parsimonious among the Model 1 and Model 2 candidate

models for each species. We ensured these species-specific candidate models were compared using

the same random effects structure. We visually explored the conditional effects of important

explanatory variables using the R package `visreg` (Breheny & Burchett 2019).

For all models, we standardized non-index variables to a mean of zero. We checked for

multicollinearity among model covariates using the R packages `corrplot` (Wei & Simko 2021)

and `performance` (Lüdecke et al. 2021). Like with other mixed models, our spatiotemporal

model in `sdmTMB` did not permit inclusion of NA values, so prior to analysis we used the R

package `gstat` (Pebesma 2004) to fill in missing within-cruise CalCOFI station data with the

inverse distance-weighted interpolation method..

Latitudinal shifts in larval abundance

To detect potential latitudinal shifts in bass larval abundance over time, we used larval

abundance predictions from the standardized index of abundance models using only years with

observed larvae and a species distribution function estimator within `sdmTMB` to calculate

temporal trends in the center of gravity (COG) for both species (Thorson et al. 2016, Anderson et

al. 2022). For larvae of both species, we plotted changes in the mean COG, grouped by decade

and the mean July SST for each decade. We obtained July SSTs off Pt. Dume, CA, which is

approximately centrally located along the southern California coast (Carter et al. 2022).

Data and Code Availability

Data and code pertaining to the sdmTMB models and cross correlation analysis are

available online in a GitHub repository: https://github.com/ETJarvisMason/xxx *(repository*

publicly available upon acceptance). We performed all analyses in R 4.3.2 (R-Core-Team 2023).

Results

We examined a total of 1,267 "*Paralabrax*" spp. larvae sorted from original station vials

(143 samples), of which 1,118 larvae could be positively identified; most (60.2%) were collected

off Baja California, Mexico (Fig. S3). Larvae we observed in samples from southern California,

USA, were 64.6% KB, 31.3% BSB, and 4.1% SSB. In samples taken off Baja California,

numbers were highest for BSB (76.5%), followed by KB (16.5%) and SSB (7.0%). We

encountered preflexion larvae more frequently across species and region; flexion and postflexion

larvae were least common (Fig. S3). By station, KB were relatively more common offshore,

while BSB and SSB were restricted to nearshore stations, with the highest numbers on average

off Baja California Sur, south of Punta Eugenia (Fig. S4). SSB larvae in the CalCOFI time series

were rare, particularly in southern California, indicating CalCOFI is not a representative survey

for SSB larvae in this region (Fig. S3, S4).

There were four sub-sampled stations with high numbers of larvae that occurred off Baja

California in the 1960s. Extrapolated species counts for these stations increased the overall raw

larval count by an additional 1,716 larvae comprised of 93.0% BSB, 5.6% SSB, and 1.2% KB;

we did not further extrapolate the counts by larval stage. The standardized raw counts of species-

specific larval abundance in southern California indicated KB larvae encountered in CalCOFI

bongo nets was generally higher year-to-year than that of BSB larvae (Fig. S5). Off Baja

California, peaks in the standardized raw counts of bass taken in CalCOFI July samples occurred

in the early 1960s and consisted primarily of BSB, while low counts from 1989 onward were

representative of a single CalCOFI station off the very northern part of Baja California (Fig. S5),

in which line 93.4 was the only Baja California CalCOFI line surveyed in those years. Cruises in

1981 and 1984 did not extend south beyond Punta Eugenia, Baja California (Fig. 1).

Temporal trends in larval abundance

The most parsimonious index standardization models included BSB larval abundance as

a time-varying function of distance to mainland coast (smoothed), incorporated all random

effects, including an independent spatiotemporal field across years ("iid"); the resulting index for

KB included larval abundance as a function of year (smoothed) with an independent

spatiotemporal field across years and no spatial random effect (Table 2). Temporal trends in the

standardized index of larval abundance for both species showed similar sporadic larval pulses in

1981, 1989, 2012, and 2014 (Fig. 3). Compared to BSB, KB larval abundance estimates were

generally higher overall and between strong larval pulses. Larval abundance for both species was

low between the late 1990s and early 2000s. On the natural log scale, we observed BSB larval

abundance fluctuated more dramatically than KB larval abundance and declined steadily from

the mid-1990s through the early 2000s before increasing again through the mid-2010s (Fig. 3a).

In contrast, KB larval abundance fluctuated less than BSB larval abundance and showed a steady

decline beginning in the early 1990s through the early 2000s before increasing again into the

2010s to levels similarly observed in the early 1980s (Fig. 3b).

Spatiotemporal trends in larval abundance

Predicted bass larval abundance off southern California also showed geographic

differences through time. BSB larval abundance was generally minimal throughout much of the

SCB, with the highest densities occurring nearshore and primarily along the central and southern

coast (Fig. 4). In contrast, KB larval abundance was relatively higher throughout the SCB, with

hotspots occurring primarily at the northern Channel Islands (Fig. 5, see '2' in Fig. S1 for

location reference). KB larval abundance was particularly low throughout the SCB between 1995

and 2005. Spatiotemporal random effects plots for BSB and KB also showed different patterns

through time (Fig. S6, S7), in which deviations from the fixed effects predictions varied

depending on species, location, and year; KB tended to show more spatiotemporal structuring

than BSB.

Relationship with fishery catch data

We found no relationship at a lag of zero between bass larval abundance estimates and

fishery catch data (Fig. 6, see Fig. S8 for index estimated for all years). In contrast, for both

species, we found marginal to strong lagged relationships between larval abundance and both

sources of catch data. For example, the highest Spearman correlation coefficients between BSB

larval abundance and total catch estimates occurred at 6- and 7-yr lags, corresponding to the age

of most fishery recruits (Fig. 6a). We found a similar trend with the BSB harvest data, in which

the highest correlations indicated larval abundance led catch by six to ten years. For KB, there

were moderate to strong correlations between larval abundance and total catch estimates from 4-

to 10-yr lags, corresponding to a higher catch-and-release rate across sizes for KB; KB larval

abundance showed marginal correlations with harvest at 4-, to 6-yr lags and was strongly

correlated with harvest at 7- to 10-yr lags (Fig. 6b).

Relationship with environmental variables

Among the 66 combinations of environmental variables, most correlations were weak ($<$

0.20, Fig. S9). Despite three relatively strong correlations; one between the BEUTI and the CUTI

(0.71), one between the BEUTI and the NPGO (0.54) and one between the Bakun index 33°N

and the NPGO (0.46), all model checks for multicollinearity produced "Low Correlation" results,

with most covariate variance inflation factor values close to 1 and none greater than 4.

In southern California, over the entire study period, 1963 to 2016, zooplankton biomass

and CalCOFI station-specific temperature (mean from surface to 10m) were positive predictors

of bass larval abundance for both species (Fig. 7, S10, S11). There was also a strong negative

relationship with distance to mainland coast (higher larval abundance nearshore; Fig. 7, S10,

S11). The influence of temperature and distance to mainland coast was stronger for BSB than

KB (Fig. 7).

Over the shorter study period, 1984 to 2016, representing Model 2, the ILD was strongly

negatively related to bass larval abundance (high bass larval abundance with shallower ILD; Fig.

7, S12, S13). Of the different upwelling indices considered (Bakun, BEUTI, CUTI), models were

essentially equally parsimonious, with only slightly lower AIC values for models with the Bakun

index (Table 3); however, none of the upwelling indices showed a strong relationship to bass

larval abundance. The positive relationship with zooplankton biomass and negative relationship

with distance to mainland coast remained important for both species during the shorter period,

while for BSB, temperature also remained important and the NPGO became more important

(Fig. 7, S12). S13).

Latitudinal shifts in bass larval abundance

Bass larval abundance center of gravity within the SCB remained remarkably stable

between the 1960s and 2010s (Fig. 8). The larval BSB center of gravity remained centrally

located along the mainland coast, fluctuating off Santa Monica Bay to the north and Huntington

Beach to the south (Fig. 8a). In contrast, the larval KB center of gravity was shifted offshore

between the northern and southern Channel Islands (Fig. 8b). We observed no consistent

northward or southward directional movement in the center of gravity for either bass species

either by decade or mean decadal July sea surface temperature.

Discussion

Our reconstructed larval indices of abundance span over half a century and represent the

longest species-specific, fishery-independent time series for the saltwater basses in southern

California, USA. Though larval abundance is generally considered to be a suitable proxy for

adult spawning stock biomass (Hsieh et al. 2005, 2006), our results indicate southern California

bass larvae abundance over the last several decades is a better reflection of future catch in the

fishery, as both BSB and KB larval abundance had moderate to strong predictive power in

forecasting fish catches. In addition, for both species, this result was consistent across different

sources of catch data, i.e., harvest reported in CPFV logbooks, and estimates of total catch

(includes harvest and releases across all fishing modes). Moreover, the utility of this predictive

power for fishery management is bolstered by the strong relationships we identified between

bass larval abundance and environmental variables (e.g., temperature, zooplankton biomass,

MLD), and thus, paints a path forward for an ecosystem approach to managing this fishery.

Additionally, our analysis revealed species-specific temporal and spatiotemporal trends, as well

as differences in the strength of environment-species relationships, suggesting potential

differences in their resilience and recovery potential with respect to intrinsic oceanographic

variability and anthropogenic climate change impacts.

Saltwater bass larvae predict future catch

Once considered the "recruitment problem" in fisheries (Haltuch et al. 2019), variability

in fishery year-class strength is now understood to be driven by a multitude of factors affecting

both pre- and post-recruitment stages, including maternal effects, and is likely to vary by species

and population (Houde 2008). Thus, although the challenge of forecasting fishery recruitment

remains, research that identifies even modest prediction capabilities is recognized as valuable for

informing management. This recognition allows us to more freely accept that prediction may be

easier for some species than others, even if we cannot identify all the mechanisms driving those

relationships. The added challenge, however, is that predictive ability has been shown to break

down for some species due to non-stationary environment-species relationships (White et al.

2019, Litzow et al. 2020), making long-term monitoring and re-evaluation an essential part of

these respective research efforts. This is especially critical when environment-species

relationships are identified based on shorter time series, as change may include both magnitude

and direction. A strength of this study is that our results are based on several multidecadal time

series. Thus, at least part of the variability in the relationships we identified between larvae and

environmental covariates reflects any inherent long-term interannual and decadal variability

influencing the strength of those relationships through time.

In the absence of data on spawning stock biomass, very early life stages like eggs and

preflexion larvae, are often a suitable proxy (Hilborn & Walters 1992, Gunderson 1993), such

that changes in early larval abundance should reflect changes in female spawner biomass and

subsequent recruitment of juveniles. Unfortunately, due to additional, variable mortality after the

larval stage, spawning biomass and larval abundance typically show weak relationships with

subsequent juvenile and fishery recruitment (Cury et al. 2014, Szuwalski et al. 2015). Our

findings with the basses are therefore unusual as we found that young larval abundance was a

better predictor of future fish catches than SSB. A strong relationship between spawning stock

biomass and larval abundance in the same year relies on 1) appreciable spawning occurring

every year, 2) spatiotemporally consistent mortality prior to larvae being surveyed, and 3) that

larvae are locally sourced within the region. Thus, in the case of saltwater basses, the breakdown

in this relationship may be partly driven by variable larval mortality or variable spawning

behavior associated with environmental conditions and harvest impacts. For BSB, there also

appears to be a lack of consistent, appreciable, locally sourced larvae from year to year.

We assume the positive relationship between saltwater bass early larval abundance and

future fish catches is associated with a survival advantage through adulthood, perhaps imparted

by fast initial larval growth and a short preflexion larval stage duration (Fontes et al. 2011,

Robert et al. 2023). Though not commonly observed, there are examples when fishery

recruitment is set in the first year of life, such that changes in early life stages reflect changes in

future fishery year-class strength (Cushing 1990, Stige et al. 2013, White et al. 2019, Schilling et

al. 2022). Our findings may also partially be a function of the geographic location of the study.

Southern California represents the northern extent of the geographic range for these species,

especially for BSB, and populations at their geographic margins tend to experience higher

population recruitment variability or recruitment limitation (Myers 1991, Neill et al. 1994, Levin

et al. 1997), which is typically reflected in subsequent population size (Armsworth 2002).

Schilling et al. (2020), for example, found a positive correlation between predicted Bluefish

(Pomatomus saltatrix) larval "settlement" (transformation to the juvenile pelagic phase) and

fishery catch-per-unit-effort at the southern end of its distribution.

Regardless of the mechanism, the moderate to strong correlations between bass larval

abundance and future catch were observed across multiple catch sets and corresponded to

biologically meaningful lags. Both species grow at similar rates, becoming susceptible to hook-

and-line fishing gear at two to three years of age and reaching size at fishery recruitment between

five and seven years for most of the time series (e.g., in2013, the minimum size limit increased

from 12 to 14 inches, which corresponds with fish ages seven to nine years old; Love et al. 1996,

Jarvis et al. 2014a, Walker et al. 2020b). Interestingly, when catch including releases was

considered, we saw that the species known for a higher catch-and-release rate (KB) showed

significant correlations across many more lags (from 3- to 10-y lags) compared to BSB (only 6-

to 7-y lags). The highest correlations for both species across both catch data sets corresponded to

the age of fishery recruits, while the significant correlations across other lags are likely an

artifact of the catch data comprising many cohorts. Thus, restricting the analysis to catch

comprising fish at fishery recruitment size or just those fish representing the modal length of the

catch may result in even stronger relationships at just one or two lags (Jarvis et al. 2014a,

Schilling et al. 2022). We chose not to focus on specific length frequencies because length data

were only available for one catch data set, and we wanted to have the ability to test the

larva/catch relationship across all available catch data sets.

One potential caveat to using the catch data as a proxy for spawning stock biomass, in the

case of BSB, is that hyperstable catches, i.e., stable, high catches, in aggregation-based fisheries

can occur during a population decline, resulting in a nonlinear relationship between catch and

abundance and an overestimate of abundance at low population sizes (Sadovy de Mitcheson

2016). However, a strong positive linear correlation exists between BSB harvest and fishery-

independent adult densities in southern California (Jarvis Mason et al. 2024); thus, there is low

probability our results were overly influenced by hyperstability. Moreover, our findings are

consistent with other studies showing positive relationships between early life history indices of

bass abundance (larvae, young-of-the-year [YOY] juveniles) and future saltwater bass fishery

recruitment, despite different collection methods and study time frames (Jarvis Mason et al.

2024, Jarvis et al. 2014a, Miller & Erisman 2014).

Species-specific differences in bass larval abundance trends

Trends in the standardized indices of larval abundance for both species suggest that the

populations of KB and BSB in southern California experienced similar sporadic, strong larval

pulses. However, our results also indicate that the KB population has more reliably persisted,

being more abundant overall and having less variable and higher larval abundance between

strong sporadic recruitment pulses than BSB. This pattern for KB is consistent with locally

sourced larval recruitment (Selkoe et al. 2007) and higher densities (Warner 1985). In contrast,

BSB larval trends may be more influenced by occasional seeding from Baja California, Mexico

(Jarvis Mason et al. 2024), a region we found to have relatively higher BSB larval abundance.

Arafeh-Dalmau et al. (2023) demonstrated northward transboundary larval connectivity of up to

500 km between northern Baja California and the SCB during the fall for California Sheephead

(*Semicossyphus pulcher*), a species with a slightly longer pelagic larval duration (PLD; 42 days

in Cowen 1991) than that of BSB (~ one month, Findlay & Allen 2002).

The inferred difference in population persistence between the two saltwater basses is also

consistent with previous anecdotal reports suggesting BSB has fluctuated in its contribution to

the saltwater bass catch, depending on temperature-driven availability (Young 1969, Frey 1971,

Feder et al. 1974). Natural log larval abundance trends also identified species-specific

differences in decadal-scale availability, with BSB larvae being relatively more abundant during

the 1980s and 1990s (a warm period) and KB larvae showing a dramatic, sustained decline

during that same period. Nevertheless, larval abundance for both species showed dramatic

increases at the end of the time series, which was one of the warmest periods on record (Jacox et

al. 2018b). Our results suggest these trends are due to different environmental influences on

saltwater bass larvae and different sensitivities to the same environmental influence.

Temperature, zooplankton, and mixed layer depth predict bass larval abundance

We found strong relationships between saltwater bass larval abundance and temperature,

zooplankton biomass, and MLD (here, MLD equals ILD). In general, our model results indicated

bass larval abundance was higher when temperatures were warmer, zooplankton biomass was

higher, and the MLD was shallower. The relationships between larvae and temperature and

zooplankton biomass are not surprising, as peak spawning for KB and BSB occurs during the

warm summer months (Erisman & Allen 2005, Jarvis et al. 2014b) following the spring

transition in the SCB (McClatchie 2014). Our metric for zooplankton biomass was displacement

volume, which can be a crude index of zooplankton community structure as it is highly

influenced by pelagic tunicates (e.g., salps) and other large gelatinous organisms (Lavaniegos &

Ohman 2007). To better elucidate the potential impact of pelagic invertebrate prey, we explored

the relationship between bass larval abundance and biomass of *C. pacificus* from spring cruises

(this data was not available for July cruises). We reasoned that increased copepod biomass

available to important adult bass prey species (e.g., coastal forage fishes) in the spring might

translate into positive maternal conditioning of larvae in summer (Walsh 2023). Inclusion of *C.*

pacificus, however, did not improve model performance. Further research is necessary to

understand the importance of bass larval prey and maternal effects on bass larval survival.

The MLD had the strongest environmental relationship with bass larval abundance, i.e.,

shallower MLDs predicted higher bass larval abundance for both species. This relationship might

be explained by "seasonal trophic amplification" in nutrient-rich regions like the California

Current Ecosystem (CCE; Xue et al. 2022). In these regions, phytoplankton and zooplankton

become more concentrated within shallower MLDs, thereby increasing prey encounter rates and,

consequently, grazing rates of zooplankton, such that higher feeding efficiency in these regions

is driven more by seasonal changes in the depth of the mixed layer than by the amount of food

present (Xue et al. 2022). In an analysis of environmental covariates on the biological response

of larval fish communities in the CCE, the ILD had the highest ability to predict ecosystem state

within the southern California region (Hunsicker et al. 2022). Despite the saltwater basses

showing relationships with the same environmental variables, relationships with temperature and

zooplankton biomass were relatively stronger for BSB than they were for KB, suggesting BSB is

more sensitive to cooler temperatures and declines in zooplankton biomass. Indeed, BSB was

historically considered a southern, subtropical/tropical species with an impermanent southern

California presence (Young 1963, 1969, Frey 1971, Feder et al. 1974).

If waters off southern California continue to warm, we might expect a shallowing of the

MLD due to increased stratification of warm surface waters and thus, higher, less variable, BSB

larval recruitment. However, the assumption that climate-driven increased stratification will

result in MLD shoaling belies evidence to the contrary (Somavilla et al. 2017), including a recent

global analysis that documented a deepening of the MLD over the last five decades at a rate of 5-

10 m dec^{-1}, despite concomitant increases in stratification (Sallée et al. 2021). Thus, if this trend

continues, and southern California saltwater bass abundance is truly a function of the MLD, then

a deeper mixed layer might counteract surface warming benefits by decreasing the concentration

of prey items. This deepening also has the potential to result in shallow, nearshore depths ("kelp-

forest depths") becoming nutrient-poor (Parnell et al. 2010), which could have negative impacts

on KB larval settlement (White and Caselle 2008).

Additional abiotic processes (beyond those examined here) are likely to influence KB

larval recruitment dynamics. For example, the timing, direction, and strength of ocean currents

impacting larval dispersal and connectivity influences if and where they are transported onshore

to settle, particularly in the physically dynamic SCB (Warner 1985). The primary KB larval

hotspot was the northern Channel Islands, which is influenced by the "transition zone" in the

SCB where many ocean currents converge (Hamilton et al. 2010). White & Caselle (2008)

reported that in southern California, the density of Giant Kelp had a positive effect on KB larval

settlement, but the relationship was conditioned on larval supply, whereby adult densities were

largely a function of whether there was a match between areas of high larval supply and areas of

higher kelp stipe density. In that study, the authors explain how eddies and temperature fronts

may drive a match-mismatch between KB larval supply and kelp spore supply. This may

partially explain why we found that Bight-wide kelp canopy extent was not an important driver

of KB larval abundance.

Larval dispersal and retention are likely equally important for BSB. Nearshore flow in the

SCB is mostly alongshore with a minor cross-shelf component (Winant & Bratkovich 1981, as

cited in Barnett et al. 1984). Whereas we found the predicted geographic distribution of BSB

larvae were almost exclusively distributed nearshore along the mainland coast, KB larvae

through time showed broad distribution within the SCB with hotspots at the northern Channel

Islands, suggesting different processes acting on their distributions (White et al. 2019). Indeed,

we observed differences in the two species' model's spatiotemporal fields over time that indicate

different latent forcing of dynamic biotic and abiotic processes. It is therefore possible that

different environmental conditions cue broad scale synchronized spawning events to take

advantage of tides and circulation patterns that help retain larvae close to shore (BSB; Barnett et

al. 1984) or provide longshore and offshore dispersal to increase the probability of reaching areas

with Giant Kelp, including at offshore islands (KB; see asymmetric directionality of KB larval

transport in Watson et al. 2010). Differences in spawning aggregation formation and migration

may also contribute to spatiotemporal variability in these species' larval distributions, as BSB

form large spawning aggregations at a few predictable locations, while KB spawning

aggregations are smaller and more broadly distributed (Erisman & Allen 2006, Jarvis et al.

2010).

All else equal, warmer temperatures for summer spawners seem to be better for higher

growth and survival of eggs and larvae (Gadomski & Caddell 1996); however, during the warm

regime of the 1980s and 1990s, the SCB shifted to nutrient-poor conditions and experienced a

higher frequency of storms, both of which negatively impacted kelp forests in the region (Parnell

et al. 2010). Thus, the steady decline in KB larval recruitment during that period may be

indirectly associated with decadal-scale declines in Giant Kelp (rather than interannual

variability in regional canopy area), despite more optimal temperatures for larval growth and

survival (Jarvis et al. 2014a). This could be one driver explaining why our models revealed the

influence of temperature on larval KB larval abundance was not as strong as that for BSB.

Although both species showed positive trends with temperature, we did not find evidence

of a northward latitudinal shift in the center of larval distribution through time. Although

climate-driven phenological and distribution shifts have been well-documented around the world

(Pinsky et al. 2020) and historical bass populations extended much farther north than southern

California during the region's tropicalization of the mid-1800s (Hubbs 1948), previous research

on larval distribution/phenology shifts in the CCE showed that species responses to climate

change are not omnipresent (Hsieh et al. 2009, Asch 2015, Auth et al. 2018, Thompson et al.

2022). In addition, northward larval dispersal of nearshore species beyond the well-known

geographic break at Pt. Conception in the SCB, at least during the last sixty years, is thought to

be limited more so by hydrographic features than temperature (Warner 1985).

Implications for fishery recovery and management

Despite the importance of the basses to southern California marine anglers, currently no

formal stock assessment exists for either species. Our standardized indices of bass larval

abundance offer the potential to develop a species-specific stock assessment model in which the

larval abundance index is used as a pre-recruit index or as a means to ground truth an

assessment's predicted recruitment deviations. Even without a stock assessment, our indices

offer predictive ability and insight into the potential for each species' fishery recovery. For

example, despite sustained low fishery catches in the early-to-mid-2010s, we found that larval

abundance for both species peaked during this time. Given our findings that larval abundance

predicts future catches, the increase in larval abundance suggests fishery recovery is imminent.

Indeed, fishery-independent indices of adult BSB and KB densities show an upward trend in

recent years (CDFW 2021, 2023). Based on larvae/catch relationships reported here, the first

peak in recent larval recruitment (2012) should be evident in catches as early as 2020,

corresponding to the average fishery recruitment age of eight years. Indeed, recent harvest

reports show increases in harvest in 2020 (KB), 2021 (BSB), and 2023 (BSB; PSMFC 2024). In

fact, although still relatively low compared to historical harvest, BSB harvest in 2023 was the

highest in over a decade, surpassing KB harvest for the first time during the same period

(PSMFC 2024).

Collectively, our results pave the way for an ecosystem approach to management of this

fishery and bring us closer toward readying the fishery for climate change. Such an approach

should consider (among other factors) species-specific differences in larvae-environment

relationships and factors influencing differences in temporal and spatiotemporal trends in larval

abundance and distributions.

Perhaps one of the most important conclusions of our study is that these two basses have

different population dynamics: 1) larval recruitment patterns indicate differences in the

persistence of their populations in southern California (e.g., background recruitment levels for

KB are higher and less episodic, which is consistent with locally-sourced recruitment (Selkoe et

al. 2007) and higher densities (Warner 1985) and has implications regarding their resilience to

harvest impacts and climate change), 2) relative to KB, BSB larval recruitment is more closely

tied to temperature, which is consistent with historical reports of BSB availability in southern

California (Jarvis Mason et al. 2024) and which also has implications for resilience to harvest

impacts and climate change, and 3) the offshore islands in the northern SCB are a notable

hotspot for KB larval recruitment, which has implications associated with habitat protection

(e.g., Marine Protected Areas, Hopf et al. 2022). In contrast, BSB larval recruitment is limited to

the mainland coast and given its sporadic nature and more southerly larval distribution in the

SCB, may be influenced more so by El Niño-driven sporadic northward advection of larvae from

Baja California (Jarvis Mason et al. 2024) as has been suggested for other fishery species in the

region (Smith & Moser 1988, Allen & Franklin 1992, Ben-Aderet et al. 2020). This too has

implications with respect to harvest impacts and climate change, as future transboundary

connectivity between the SCB and Mexico is predicted to be significantly reduced (assuming

decreased PLDs and lower kelp persistence), thus highlighting the importance of protecting

habitats important for larval connectivity and binational conservation efforts to manage the

collective resource (Arafeh-Dalmau et al. 2023).

Environment-species relationships are typically nonstationary and may not only change

in magnitude, but direction (White et al. 2019). Here, we had the opportunity to test whether

strong environmental relationships we identified using data spanning 54 years (1963-2016) held

up using a shorter part of the time series (1984-2016). Although the influence of zooplankton

biomass was important in both periods for both species, the influence of temperature on KB

larval abundance was weakened/lost in the shorter time series. The shorter time series began just

before the well-documented 1988-89 North Pacific climate shift (Hare & Mantua 2000) after

which, some species responses to temperature changed (Litzow et al. 2019) and previously

established physical (e.g., surface temperature) and ecological relationships with the NPGO (Di

Lorenzo et al. 2008) became weaker (Litzow et al. 2020). Thus, we recommend regular

monitoring and re-evaluation of the relationships identified here. This recommendation points to

the value of maintaining long-term monitoring data streams, like CalCOFI, particularly for other

data-limited state-managed and small-scale fisheries. Future research into the role that maternal

effects and prey type and condition play in bass larval dynamics will likely provide additional,

important ecosystem considerations for managing this culturally and economically important

recreational-only fishery.

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Paralabrax spp. larvae to species. The BSB and KB illustrations included in some of the figures

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1078

1079 Table 1. Temporal and spatial characteristics of California Cooperative Fisheries Investigations cruises
1080 selected for analyzing temporal and spatial trends in species-specific bass *Paralabrax* spp. larval
1081 abundance, 1963-2016.

Scale of inference	Observational unit	Number of observations at this scale	Response variable and units
Species: coastal basses **Temporal:** annual July surveys over 37 years spanning 5 decades **Spatial:** continental shelf of the Southern California Bight and Baja California, Mexico	Oceanographic cruise with stations positioned along transect lines that are 40 km apart; along transect lines, offshore (nearshore) stations are spaced 40km (20km) apart	**Minimum:** 2 stations per cruise **Mean:** 73 stations per cruise **Maximum:** 220 stations per cruise	Bass larval abundance (number of larvae under ten m^2 of sea surface)

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1084 **Table 2.** Results comparison of candidate species distribution models* used to generate a standardized
1085 index of larval abundance for Barred Sand Bass *Paralabrax nebulifer* and Kelp Bass *P. clathratus* in
1086 southern California, USA, 1963-2016.
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Barred Sand Bass					
Formula	Time-varying	Spatial	Spatio-temporal	REML AIC	AIC
~ 0 + dist	**y**	**on**	**iid**	**817**	**818**
~ 0 + dist	y	off	iid	828	--
~ 0 + dist + day	y	on	iid	--	818
~ 0 + dist + s(moon, bs = "cc", k=4)	y	on	iid	--	820
Kelp Bass					
Formula	Time-varying	Spatial	Spatio-temporal	REML AIC	AIC
~ s(year)	**n**	**off**	**iid**	**--**	**1484**
~ s(year) + dist	n	off	iid	--	1527
~ s(year) + moon	n	off	iid	--	1486

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*Main effects are listed in the formula; random effects are spatial and/or spatiotemporal. Time-varying =
main effects vary across years as a random walk, iid = independent and identically distributed across
years, REML = restricted maximum likelihood, AIC = Akaike information criteria, dist = distance to
mainland coast, day = day of year, moon = continuous numerical moon phase, bs = basis dimension, "cc"
= cyclic.

Table 3. Results comparison of candidate species distribution models* used to measure the effects of
environmental influence on larval abundance for Barred Sand Bass *Paralabrax nebulifer* and Kelp Bass
P. clathratus in southern California, USA during two time periods (Model 1, 1963-2016; Model 2, 1984-
2016). Bold represents top model.

Barred Sand Bass					
Formula	Spatial	Spatio-temporal	Passed sanity check?	REML AIC	AIC
Model 1, 1963-2016					
~ dist_km + temp.10m + npgo + oni + zoop	**off**	**iid**	**y**	**708**	**700**
~ dist_km + temp.10m + npgo + oni + zoop	off	off	y	766	--
~ dist_km + temp.10m + npgo + oni + calpac	off	iid	y	--	749
Model 2, 1984-2016					
~ dist_km + temp.10m + npgo + oni + zoop bakun30 + south.ild	**off**	**iid**	**y**	--	**576**
~ dist_km + temp.10m + npgo + oni + zoop + bakun33 + south.ild	off	iid	y	**--**	578
~ dist_km + temp.10m + npgo + oni + zoop + south.beuti + south.cuti + south.ild	off	iid	y	--	578
~ dist_km + temp.10m + npgo + oni + calpac bakun30 + south.ild	off	iid	y	**--**	619
Kelp Bass					
Formula	Spatial	Spatio-temporal	Passed sanity check?	REML AIC	AIC
Model 1, 1963-2016					
~ dist_km + temp.10m + npgo + oni + zoop	**off**	**iid**	**y**	**--**	**1270**
~ dist_km + temp.10m + npgo + calpac + oni	off	iid	y	--	1303
Model 2, 1984-2016					
~ dist_km + temp.10m + npgo + oni + zoop + bakun33 + south.ild	**off**	**iid**	**y**	**--**	**939**
~ dist_km + temp.10m + npgo + oni + zoop + bakun30 + south.ild	off	iid	y	--	940
~ dist_km + temp.10m + npgo + oni + zoop + kelp + south.ild	off	iid	y	--	940
~ dist_km + temp.10m + npgo + oni + zoop + south.beuti + south.cuti + south.ild	off	iid	y	--	940
~ dist_km + temp.10m + npgo + oni + calpac + bakun33 + south.ild	off	iid	y	**--**	973

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* Main effects are listed in the formula; random effects are spatial and/or spatiotemporal. iid =
independent and identically distributed across years, REML = restricted maximum likelihood, AIC =
Akaike information criteria, dist_km = distance to mainland coast, temp.10m = average CalCOFI sea
surface temperature to 10 m depth, npgo = North Pacific Gyre Oscillation index for July, zoop = CalCOFI
zooplankton biomass, oni = Oceanic Niño Index for June/July, calpac = CalCOFI spring biomass of the
calanoid copepod, *Calanus pacificus,* bakun33 = Bakun upwelling index at 33º N, south.ild = isothermal
layer depth, bakun30 = Bakun upwelling index at 30º N, south.beuti = Biologically Enhanced Upwelling
Index, south.cuti = Coastal Upwelling Transport Index, kelp = areal canopy extent of Giant Kelp
Macrocystis pyrifera, CalCOFI = California Cooperative Oceanic Fisheries Investigations.

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Figure 1. Maps of a) CalCOFI station locations surveyed along transect lines during July cruises from
1963 to 2016 and numbers of samples per station from which formalin preserved *Paralabrax* spp. larvae
were identified to species, and b) representative sea surface temperature contours off southern California,
USA and Baja California, Mexico in the month of July (NOAA NCDC OISST version2p1 AVHRR
monthly sea surface temperature data). Blue arrows depict the directional flow of the California Current.
CalCOFI = California Cooperative Oceanic Fisheries Investigations, SCB = Southern California Bight.
Shaded temperature contours range from 14 °C in the north to 26 °C off the southern tip of Baja
California and northward into the Gulf of California, Mexico.

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Figure 2. Temporal trends in regional and local environmental variables selected for modeling southern California, USA, *Paralabrax* spp. larval abundance. Temp.10m = CalCOFI ocean temperature averaged from the sea surface down to 10 m depth, zoop = CalCOFI zooplankton biomass, npgo = North Pacific Gyre Oscillation index for July, oni = Ocean Niño Index for June/July, bakun33 = Bakun upwelling index at 33°N, bakun30 = Bakun upwelling index at 30°N, south.beuti = Biologically Enhanced Upwelling Index, south.cuti = Coastal Upwelling Transport Index, south.ild = isothermal layer depth, calpac = spring biomass of the calanoid copepod, *Calanus pacificus*, area = areal canopy extent of Giant Kelp *Macrocystis pyrifera*. CalCOFI = California Cooperative Oceanic Fisheries Investigations.

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1130 **Figure 3.** Standardized indices of larval abundance in normal space (left panels) and link space ln(abun),
1131 right panels) for a) Barred Sand Bass *Paralabrax nebulifer* and b) and Kelp Bass *P. clathratus* in southern
1132 California, USA, 1963-2016. Shaded ribbons denote 95% confidence intervals.
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1135 **Figure 4.** Predicted Barred Sand Bass *Paralabrax nebulifer* larval abundance in southern
1136 California, USA, by year, 1963-2016. Scale bar denotes abundance on the natural log scale. The
1137 legend color ramp ranging from light to dark denotes higher and lower abundance, respectively.

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Figure 5. Predicted Kelp Bass *Paralabrax clathratus* larval abundance in southern California, USA, by year, 1963-2016. Scale bar denotes abundance on the natural log scale. The legend color ramp ranging from light to dark denotes higher and lower abundance, respectively.

Figure 6. Cross-correlation Spearman coefficients between larval abundance and total catch estimates
(left panels, includes estimates of fish kept and released across all fishing modes) and total numbers of
reported harvest on CPFVs (right panels) across yearly lags for a) Barred Sand Bass *Paralabrax nebulifer*
and b) and Kelp Bass *P. clathratus* in southern California, USA. CPFVs = Commercial Passenger Fishing
Vessels. Negative lags represent larvae leading catch. Shaded ribbon denotes the 95% confidence band.
Gradation in shading of points depicts the strength of the correlation, ranging from zero or weak (light) to
marginal/moderate or strong (darker).

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Figure 7. Model 1 coefficient estimates of the generalized linear mixed effects model depicting the relationships between bass larval abundance and environmental covariates from 1963 to 2016 (Model 1) and from 1984 to 2016 (Model 2) for a) Barred Sand Bass *Paralabrax nebulifer* and b) and Kelp Bass *P. clathratus* in southern California, USA. ild = isothermal layer depth, upwelling = Bakun upwelling index at 33°N, zoop = CalCOFI zooplankton biomass, temp = CalCOFI ocean temperature averaged over upper 10 m, oni = Ocean Niño Index for June/July, npgo = North Pacific Gyre Oscillation index for July, shore_dist = distance from survey station to mainland coast. Model coefficients greater (less) than one reflect positive (negative) relationships. Lines depict 95% confidence intervals. CalCOFI = California Cooperative Oceanic Fisheries Investigations**.**

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1165 **Figure 8.** Decadal trends in the mean center of gravity for larval distributions of a) Barred Sand Bass
1166 *Paralabrax nebulifer* and b) and Kelp Bass *P. clathratus* in southern California, USA, 1963 to 2016, by
1167 decade (symbols) and annual mean sea surface temperature (colors). Lines depict latitudinal and
1168 longitudinal 75% confidence intervals. 1 = northern Channel Islands, 2 = Santa Monica Bay, 3 = southern
1169 Channel Islands, 4 = Huntington Beach
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