

Influence of combined temperature and food availability on Peruvian anchovy (*Engraulis ringens*) early life stages in the northern Humboldt Current system: A modelling approach

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Abstract :

In the northern Humboldt Current system (NHCS), the Peruvian anchovy (*Engraulis ringens*) constitutes the bulk of landings and has a significant socioeconomic contribution. Understanding the impact of environment on the early-life stages of anchovy and further population dynamics remains challenging. Climate variability at a variety of scales modulates currents velocity, temperature and food availability, impacting early-life stages drift, growth and survival. In order to investigate these impacts, we developed Ichthyop-DEB, an individual-based model including larval retention processes and a Dynamic Energy Budget (DEB) bioenergetic module for larval growth. First, we assessed the effect of hydrodynamic simulations horizontal resolution on simulated larval retention patterns using a recruitment age-criterion of 30 days. Then, we evaluated the impact of the following biological processes on simulated larval recruitment patterns: i) a minimum size-criterion (2 cm), as opposed to a minimum age-criterion (30 days), to be considered as recruited, ii) the upper larval thermal limit tolerance of the species, for which lab experiments are lacking, and iii) a constant larval mortality rate. We found that using different resolutions of the hydrodynamic model (10 and 2 km) led to similar simulated larval retention patterns. Retention was highest when spawning occurred in the superficial layer (0 - 15 m) in austral winter and in the deepest considered layer (30 - 45 m) in summer. Coupling with the DEB model produced contrasted growth patterns on the continental shelf with a strong month-latitude interaction. Larval recruitment was strongest from 6° to 10° S in austral summer, largely contributing to the average seasonal pattern. Depending on the temperature correction function tested with the bioenergetic module, simulated larval recruitment

could also be strong in the northernmost zone (2° - 4°S), an area not known for abundant anchovy populations, which suggests a possible thermal growth limitation. Finally, sensitivity tests performed on larval growth limitation by food suggested a deficiency in food supply in the southernmost zone (18° - 20°S).

Highlights

► We developed an individual-based model including larval retention and a Dynamic Energy Budget bioenergetic module. ► Results show that Peruvian anchovy larval growth accelerates with increasing temperature, but the upper threshold is still not properly defined. ► Food availability limits anchovy growth and recruitment in southern Peru, only. ► Spawning depth has a significant effect on Peruvian anchovy recruitment with a seasonal modulation.

Keywords : Ichthyop-DEB model, early life stages survival, Peruvian anchovy, larval drift, larval growth

46 1 Introduction

47 The northern Humboldt Current system (NHCS) currently produces more fish catch
48 per unit area than any other marine ecosystem (Bakun and Weeks 2008; Chavez et al. 2008)
49 despite not having the largest primary productivity (Chavez and Messié 2009; Checkley et al.
50 2017). In the NHCS, the Peruvian anchovy (*Engraulis ringens*) is a highly prolific species (~15
51 000 eggs/batch) that reaches its sexual maturity at the age of one year. The anchovy spawns
52 mainly in the coastal zone close to surface (Gutiérrez et al. 2007, 2008). Early life stages are
53 taking advantage of the exceptional continental shelf nursery area thanks to the high
54 productivity from the upwelling zone, which contribute to make *E. ringens* the most abundant
55 species, supporting the world largest mono-specific fishery (Fréon et al. 2003; Alheit and
56 Niquen 2004; Gutiérrez et al. 2016; Checkley et al. 2017; FAO 2020). *E. ringens* fishery is
57 managed based on scientific monitoring of population indicators (Ayón 2000; Gutiérrez et al.
58 2007), but the link between environmental variability and anchovy recruitment, and thereby
59 biomass of the adult population, is still unclear.

60 Modeling studies have been conducted to understand the hydrodynamics of the
61 NHCS (Penven et al. 2005; Colas et al. 2012), its interannual variability (Colas et al. 2008;
62 Espinoza-Morriberón et al. 2017), its potential changes under future climate scenarios (Oerder
63 et al. 2015; Echevin et al. 2020), and the seasonal cycle and intraseasonal variability of surface
64 chlorophyll (Echevin et al. 2008, 2014). These works provided the physical and biogeochemical
65 basis for ecological studies in the NHCS, from testing Bakun's triad hypothesis (Bakun 1998)
66 for small pelagic fish recruitment and early life stages survival (Lett et al. 2007) to simulating
67 Peruvian anchovy recruitment depending on environmental conditions (Brochier et al. 2008,
68 2009, 2011, 2013; Xu et al. 2013, 2015).

69 The first anchovy larval drift modeling study conducted in the NHCS found
70 similarities between simulated anchovy larval near surface retention over the continental shelf
71 and observed egg distribution (Brochier et al. 2008). This results suggests a reproductive
72 strategy of the Peruvian anchovy adapted to maximize reproduction success, a pattern also
73 found in other Eastern Boundary Upwelling Systems (Brochier et al. 2009). Later, Brochier et
74 al. (2013) used a physical-biogeochemical model to evaluate the effect of currents and
75 productivity on nursery areas reduction due to climate change, and found a negative effect on
76 Peruvian anchovy early life stages survival. However, the effects of temperature and food
77 availability on larval growth and survival were not directly taken into account and coastal

78 retention was evaluated using a constant planktonic life duration (PLD) of 30 days (recruitment
79 age-criterion). Xu et al. (2013) applied a 3D full life cycle model to the Peruvian anchovy over
80 the period 1991 - 2007 using a bioenergetic growth model, with a 5-cm limit for the successful
81 recruitment of individuals (recruitment size-criterion). They obtained an increased age at
82 recruitment in 1998 (El Niño conditions) as well as a notable decrease in individuals' survival.
83 Then, Xu et al. (2015) underlined the importance of spatial variability in environmental
84 conditions of the NHCS and thereby on simulated recruitment of Peruvian anchovy.

85 Despite these previous works, the question of the relative contributions of the two
86 main *E. ringens* spawning seasons, the most intense being in September and the second in
87 February-March (Seatersdal and Valdivia 1964; Perea et al. 2011), to the over-all recruitment
88 remains an open debate (Walsh et al. 1980; Perea et al. 2011). In addition, lags in spawning
89 periods due to changes in environmental conditions have been reported (Perea and Buitrón Díaz
90 1999). Here we tested the hypothesis that the higher food availability combined with warmer
91 condition in summer could contribute to better growth conditions in this season, and
92 compensate for the lower retention pattern previously predicted in the surface layer. Such
93 hypothesis would imply the summer spawning to be the main contribution to recruitment, which
94 could have consequences for *E. ringens* fisheries management in Peru. In order test this
95 hypothesis, we developed Ichthyop-DEB, an individual-based model including larval retention
96 processes (Lett et al. 2008) and a Dynamic Energy Budget (Kooijman 2010) bioenergetic
97 module for larval growth. Using this tool, we assessed the effect of hydrodynamic simulations
98 horizontal resolution on simulated larval retention patterns using a recruitment age-criterion of
99 30 days. Then, we evaluated the impact of the following biological processes on simulated
100 larval recruitment: i) a minimum size-criterion (2 cm), as opposed to a minimum age-criterion
101 (30 days), to be considered as recruited, ii) the upper larval thermal limit tolerance of the
102 species, for which lab experiments are lacking, and iii) a constant larval mortality rate.

103 2 Methods

104

105 An individual-based model (IBM) simulates populations and communities by
106 following individuals and their properties (DeAngelis and Grimm 2014). Our IBM description
107 follows the standard protocol proposed for describing IBMs (Grimm et al. 2006, 2010). Our
108 IBM was developed from the modeling tool Ichthyop v.3.2 (Lett et al. 2008,
109 <http://www.ichthyop.org/>) and since our main development is coupling with a bioenergetic
110 growth model based on Dynamic Energy Budget (DEB) theory, hereafter we will refer to our
111 model as Ichthyop-DEB. This new tool is able to simulate the effect of ocean currents on the
112 drift of ichthyoplankton as well as the combined effects of temperature and food on their
113 growth.

114

115 2.1 Purpose

116

117 The general purpose of Ichthyop-DEB is to evaluate the impact of environmental
118 factors experienced by fish (herein anchovy) eggs and larvae on recruitment. Considered
119 environmental factors are described in section 2.2.

120

121 2.2 Entities and state variables

122

123 The model included two types of entities: the environment and the individuals (eggs
124 and larvae). The environment was represented by stored hydrodynamic simulations from the
125 Coastal and Regional Ocean COMMUNITY model (CROCO, <https://www.croco-ocean.org/>, Hilt
126 et al., 2020; Shchepetkin and McWilliams, 2005) coupled to a biogeochemical model (PISCES,
127 Aumont et al. 2015), providing the following forcing state variables: ocean current velocities
128 (m s^{-1}), temperature (in $^{\circ}\text{C}$, which will then be transformed into Kelvin degrees as a requirement
129 of the growth model) and meso-zooplankton concentrations ($\mu\text{mol C l}^{-1}$) over the NHCS. The
130 meso-zooplankton field was chosen as a proxy for food as it constitutes the main energy source

131 for anchovy larvae off Peru (Espinoza and Bertrand 2008, 2014; van der Lingen et al. 2009). In
132 PISCES, the mesozoopankton growth equation includes a quadratic dependency to
133 mesozooplankton in order to depict grazing by the higher, non-resolved trophic levels (Aumont
134 et al., 2015). Individuals were characterized by the following state variables: age (d), location
135 in 3D (longitude, latitude and depth), amount of energy reserve (E , J) and structure (V , cm³).
136 Structure (V) was converted to standard length using a shape coefficient (δ_M).

137 We used three different CROCO-PISCES configurations with contrasted grid size
138 and bathymetry in order to evaluate the model sensitivity to spatial resolution (Table 1). The
139 first configuration (D01) extends from 22 °S to 5 °N in latitude and from 96 °W to 70 °W in
140 longitude, with a horizontal resolution of ~ 10 km and 64 vertical levels. The bathymetry comes
141 from the STRM30 dataset (Becker et al. 2009). It was interpolated on the model grid and
142 smoothed in order to reduce errors in the horizontal pressure gradient. The second configuration
143 (D02) extends from 20 °S to 5 °S with a horizontal resolution of ~ 2 km and 42 vertical levels.
144 The D02 domain is embedded into the D01 domain through an offline 1-way nesting procedure
145 (“roms2roms”; Mason et al. 2010). We used two different bathymetries for the D02 domain,
146 one interpolated from the D01 bathymetry (i.e., similar to the D01 bathymetry) and one
147 interpolated directly from the SRTM30 dataset. Note that consequently the former is smoother
148 than the latter, so in the following we call the two configurations D02s and D02r, respectively
149 and contain no biogeochemical outputs. Horizontal advection of temperature, salinity and
150 momentum is done with a third-order scheme (UP3), and horizontal advection of
151 biogeochemical tracers is with a fifth-order WENO5 scheme (to avoid negative values). The
152 vertical grid is discretized in terrain-following coordinates. The vertical advection of tracers
153 uses a fourth-order Akima scheme and a fourth-order compact scheme, Splines, is used for the
154 momentum vertical advection (Shchepetkin 2015). Vertical mixing is parameterized using the
155 KPP formulation (Large et al. 1994).

156 The three configurations were used to obtain quasi-equilibrium solutions forced by
157 monthly climatologies (over the period 2008-2015) at their surface and lateral boundaries. They
158 all used the same atmospheric forcing fields. The wind stress was computed from a monthly
159 climatology of the Advanced Scatterometer (ASCAT, 1/4° gridded product). Other atmospheric
160 fluxes (shortwave heat fluxes and freshwater fluxes) come from the COADS monthly
161 climatology (da Silva et al. 1994). Model sea surface temperature (SST) was restored to
162 observed climatological monthly SST derived from the merged multi-sensor OSTIA product

163 (Donlon et al. 2012) following the methodology of (Barnier et al. 1995). Open boundary
164 conditions for the D01 domain were taken from a monthly climatology of the GLORYS2V4
165 reanalysis (1/4° horizontal resolution; Ferry et al. 2012) for temperature, salinity, zonal and
166 meridional current velocity components and sea-level height. Biogeochemical conditions were
167 taken from the CARS2009 climatology (Ridgway et al. 2002) for oxygen and nutrients (nitrate,
168 phosphate and silicate), and from the World Ocean Atlas climatology (WOA2005, Conkright et al.,
169 2002) for dissolved organic carbon, dissolved inorganic carbon and total alkalinity. Iron is obtained
170 from a NEMO-PISCES global simulation climatology (Aumont and Bopp 2006). Climatological
171 simulations were run for 10 years, the first 4 years being considered as a spin-up. In the present
172 study, the last three years were used to force Ichthyop-DEB. Note that the same D01 simulation
173 has been used in a recent study (Echevin et al. 2021; NSH simulation therein) in which it has
174 been shown to reproduce the general characteristics of the upwelling dynamics
175 (and biogeochemical tracers' distribution) in the region through a validation against in-situ
176 observations.

177

178 2.3 Process overview and scheduling

179

180 Virtual individuals (eggs/larvae) were spawned in the environment according to a
181 determined spatial (area, depth and bathymetry) and temporal (month and frequency) spawning
182 strategy that constituted the initial conditions (section 2.5). For each time step (2 hours) each
183 egg or larva was passively transported by the 3D current fields, grew according to the 3D food
184 and temperature fields and was then tested for recruitment (section 2.6).

185

186 2.4 Design concepts

187

188 *Stochasticity.* Individuals were initially randomly distributed over the Peruvian
189 continental shelf. We chose the number of individuals released large enough (5 000) such that
190 the variability of simulated recruitment between three replicates of the same simulation was
191 negligible.

192 *Observation.* A spatio-temporal recruitment index was computed for each
 193 simulation and compared with egg presence observational data. Three recruitment criterions
 194 were tested, either based on 30-day retention over the continental shelf (criterion 1), retention
 195 until the larval length reach 2 cm (criterion 2) and mortality-weighted larval worth (see section
 196 2.6 *mortality*) until the larval length reach 2 cm (criterion 3). Standard length of a larva used
 197 for criterion 2 and 3 relates to its structural volume (section 2.6) as follows:

$$198 \quad L_w = \frac{V^{1/3}}{\delta_M} \quad \text{Eqn 1}$$

199 where L_w is the standard length (cm), V is the structural volume (cm^3) and δ_M is a shape
 200 coefficient.

201

202 2.5 Initialization

203

204 In each simulation, individuals were released within the coastal spawning area (Fig.
 205 1) each month at days 1, 10 and 20, during the three climatological years used. The coastal
 206 spawning area was defined as the volume of water between latitudes 2° S and 20° S, depth
 207 range [0 – 45 m] and from the coast to isobath 2000 m. Individuals were released randomly
 208 within that defined volume, leading to a uniform distribution both horizontally and vertically.

209 The initial values of the bioenergetic variables for each individual were set as initial
 210 reserve $E_0 = 1J$ and initial structure $V_0 = 0.000001\text{cm}^3$. Formally, an egg is only composed of
 211 reserve but in practice a very small value for V_0 value was needed in order to avoid division by
 212 zero in the mobilization equation (Equation of the \dot{p}_C flux Eqn 3, see Appendix and Kooijman
 213 2010). We checked that a value of V_0 larger by one order of magnitude did not change the
 214 results.

215

216 2.6 Sub-models

217

218 *Transport.* Virtual eggs and larvae were advected using a trilinear interpolation
 219 scheme of the velocity fields derived from CROCO-PISCES, in space and time, and using a
 220 forward Euler numerical scheme with horizontal diffusion following Peliz et al. (2007). The
 221 transport was assumed to be purely Lagrangian with no egg buoyancy nor larval vertical
 222 migration.

223 *Growth.* Dynamic Energy Budget (DEB) theory (Kooijman 2010, Sousa et al. 2010)
 224 was used to simulate the growth of embryos and larvae. It describes the acquisition and
 225 utilization of energy for metabolic processes during the complete life cycle of an organism
 226 depending on temperature (T) and food conditions (X). An individual is represented by two
 227 compartments: Reserve (E, in J) and Structure (V, in cm³). Energy assimilated from food in
 228 the environment contributes to reserve once the organisms starts feeding. A fraction κ of the
 229 energy mobilized from reserve is first allocated to somatic maintenance (E), and the excess
 230 energy is used to increase the structure (V), i.e., standard length (see Eqn 1). The remaining
 231 fraction of mobilized energy (1- κ) is allocated to development and maturity maintenance. The
 232 equations of the DEB model as implemented in the Lagrangian tool routines can be found in
 233 Supplementary material together with the schemes of the energy fluxes and the state variables
 234 of a DEB model for an embryo and a feeding larva. We here only present the system of two
 235 ordinary differential equations that describe the growth of an individual and how it is impacted
 236 by food (X) and temperature (T) conditions using a Holling type II scaled functional response
 237 (f) and a temperature correction function that describe how physiological rates are impacted
 238 within and outside the optimum temperature range:

$$239 \quad \begin{cases} \frac{dE}{dt} = f c_T \{ \dot{p}_{Am} \} V^{2/3} - \dot{p}_C \\ \frac{dV}{dt} = \frac{\kappa \dot{p}_C - c_T [\dot{p}_M] V}{[E_G]} \end{cases} \quad \text{Eqn 2}$$

$$240 \quad \dot{p}_C = \frac{E \left([E_G] \frac{c_T \{ \dot{p}_{Am} \}}{[E_m]} V^{-\frac{1}{3}} + [\dot{p}_M] \right)}{\kappa \left(\frac{E}{V} \right) + [E_G]} \quad \text{Eqn 3}$$

$$241 \quad f = \frac{X}{(X + K)} \quad \text{Eqn 4}$$

$$242 \quad c_T = \exp \left(\frac{T_A}{T_1} - \frac{T_A}{T} \right) \left(\frac{1 + \exp \left(\frac{T_{AL}}{T_1} - \frac{T_{AL}}{T_L} \right) + \exp \left(\frac{T_{AH}}{T_H} - \frac{T_{AH}}{T_1} \right)}{1 + \exp \left(\frac{T_{AL}}{T} - \frac{T_{AL}}{T_L} \right) + \exp \left(\frac{T_{AH}}{T_H} - \frac{T_{AH}}{T} \right)} \right) \quad \text{Eqn 5}$$

243

244 With $[E_m]$ is the maximum reserve density and f the Holling type II scaled scaled functional
245 response, X the local food concentration (here meso-zooplankton fields coming from CROCO-
246 PISCES), K the half-saturation constant and C_T a non-monotonic temperature correction
247 function, T the water temperature surrounding an individual (coming from CROCO-PISCES),
248 T_1 is the reference temperature (for which flux parameters were estimated), T_A is the Arrhenius
249 temperature (Kooijman 2010) and T_{AL} , T_{AH} , T_L , T_H are constants used to define a curved shape
250 of the temperature correction according to temperature.

251 In the absence of observations for the Peruvian anchovy (*E. ringens*), we used here parameters
252 estimated for the European anchovy (*Engraulis encrasicolus*, Pethybridge et al. 2013, Table 2),
253 a taxonomically close species that is also distributed in upwelling zones. We validated that these
254 parameters were able to describe larval growth in field and lab conditions (Figure S1).

255 *Recruitment.* We considered two criteria for larval recruitment, hereafter referred
256 to as the age- and size-criterion, respectively. For the age-criterion, an individual was
257 considered as recruited if it was within the coastal zone (offshore limit 2000 m isobath) at age
258 30 days, like in the previous modeling study (Brochier et al. 2008). For the size-criterion, an
259 individual was considered as recruited when it was within the coastal zone at a size larger than
260 2 cm. The 2 cm threshold was chosen because Peruvian anchovy larvae reached an average size
261 of 2 cm at 30 days (Castro and Hernandez 2000; Moreno et al. 2011; Rioual et al. 2021).

262 *Mortality.* We used the concept of super-individual (Scheffer et al. 1995; Parry and
263 Evans 2008) by assigning an initial worth of 1 to each individual, then applying a constant daily
264 mortality rate until the age at recruitment. The daily mortality rate was set to 0.1 as proposed
265 for anchovy (Bailey and Houde 1989; Houde 2008).

266

267 2.7 Simulations and sensitivity analysis

268

269 Eight simulations were performed in order to explore the model sensitivity to
270 different environmental forcing fields and larval growth parameters (Table 1). A first set of four
271 simulations was carried out without the growth model, the first three (Sim 1, Sim 2 and Sim 3)
272 in order to test the effect of the spatial resolution of the current velocity fields on simulated
273 retention patterns using the three CROCO-PISCES configurations described in section 2.2 and

274 in order to fit the spatial extent of the 2 km grid, individuals release was constrained in the
275 coastal area between 6° S and 14° S (Fig. 1, dotted box) for all three simulations. The fourth
276 simulation (Sim 4) was similar to Sim 1, but the spawning zone was extended between 2° to
277 20° S aiming to compare to Brochier et al. (2008)'s results. Larval retention was calculated
278 using an age-criterion of 30 days in all four simulations.

279 A second set of four simulations was carried out with the growth model, using the
280 D01 grid and the larval size threshold (20 mm) as a criterion for recruitment (size-criterion).
281 The DEB parameters values are given in Table 2, corresponding to *E. encrasicolus* (Pethybridge
282 et al. 2013) but fitting *E. ringens* larval growth well (supplementary Fig. S1). In order to
283 disentangle the effects of food and temperature on growth, and ultimately on recruitment,
284 simulations were repeated using a half saturation parameter either null, i.e., $f = 1$ (no food
285 limitation) or calculated such that $f = 0.5$ for the average meso-zooplankton concentration over
286 the continental shelf off Peru with a half saturation constant of 1.6. To contrast the effect of
287 temperature on growth, we used two different shapes for the curve of the energy fluxes
288 temperature correction (C_T ; Fig. 2). In both cases, C_T dropped to very low values for temperature
289 higher than 25°C but in the first case the maximum value of C_T was at ~ 19 °C and then it
290 dropped slowly (hereafter referred to as “case 1”) whereas in the second case the maximum was
291 at ~ 23 °C and then it dropped quickly (hereafter referred to as “case 2”). These temperature
292 thresholds were chosen to fit *E. ringens* distribution in Peru (Castillo et al. 2022). All
293 simulations lasted 90 days, a value found from preliminary simulations as long enough for the
294 slowest growing individuals to reach the recruitment size. Larval retention at 30 days (age
295 criterion, Sim 4) was also compared with the size-criterion in all four simulations (Sim 5, Sim
296 6, Sim 7 and Sim 8). In order to quantify the variation of results between simulations with (Sim
297 5, Sim 6) and without (Sim 7, Sim 8) food limitation, we calculated the percentage of variation,
298 e.g. $\frac{Sim\ 7 - Sim\ 5}{Sim\ 5} * 100$.

299 **3 Results**

300

301 *Physical configurations*

302 Simulated retention patterns obtained with the three tested configurations of the
303 hydrodynamic model were very similar (Sim 1, Sim 2 and Sim3, Supplementary material, Fig.
304 S2). The main differences concerned the D02s configuration (Sim 2) that exhibits slightly
305 higher retention values for austral summer months (Supplementary material, Fig. S2a) and for
306 the coastal spawning zone (0-100m isobath, Supplementary material, Fig. S2d). The latitudinal
307 range between 10° - 12° S was the most favorable for larval retention (Supplementary material,
308 Fig. S2b) and a direct relationship was observed between spawning depth and larval retention,
309 being lower near the surface and higher in deeper layers (Supplementary material, Fig. S2c).

310 Globally, in Sim 4 we obtained similar retention patterns as Brochier et al. (2008),
311 who used different physical forcing fields. The interaction of spawning depth and month
312 displayed the same characteristic pattern, with highest retention in austral winter for the
313 superficial spawning depth level (0 - 15 m) and in summer for the deepest spawning levels (30
314 - 45 m; bars in Fig. 3). We also found the same seasonal trends when the spawning area was
315 split into inner shelf (0 - 100 m isobaths) and offshore shelf (100 - 500 m and 500 - 2000 m
316 isobaths; lines in Fig. 3). The results differed most notably in highest values obtained between
317 8°S and 12°S from June to September in Brochier et al. (2008) as opposed to between 6°S and
318 8°S in January-February here (Supplementary material, Fig. S3).

319 *Growth and bioenergetics*

320 When we included larval growth (Sim 5 and Sim 6) and changed the criterion used
321 for retention from age (30 days, Sim 4, grey bars in Fig. 4 and Fig. 5) to size (> 2 cm, Sim 5
322 and Sim 6, black line in Fig. 4 and Fig. 5) we obtained nearly identical results in temperature
323 correction's case 2 (Sim 6, Fig. 5), i.e., when the bio-energetic fluxes decayed abruptly at high
324 temperature.. In case 1 (Sim 5, Fig. 4), when the bio-energetic fluxes decayed smoothly at high
325 temperature, the patterns remained similar, with highest recruitment values obtained in summer
326 (Fig. 4a). at depth (Fig. 4c) and close to the coast (Fig. 4d). When mortality was included (red
327 lines in Fig. 4 and Fig. 5), in case 1 the patterns did not change notably but the trends dampened
328 (Fig. 4), whereas in case 2 (Fig. 5) we obtained a stronger seasonal variability highlighting the

329 difference between summer and winter (Fig. 5a) and highest values for the northern part of the
330 domain (2° - 4° S) instead of the central (10° - 14° S) and southern (18° - 20° S) zones without
331 mortality (Fig. 5b). Most notably, recruitment was highest for the intermediate spawning depth
332 level (15 - 30 m) as opposed to highest for the deepest depth level (30 - 45 m) without mortality
333 (Fig. 5c).

334 At the surface layer (0 - 15 m), winter spawning months favored *E. ringens*
335 recruitment when using an age-criterion (Fig. 6a) and a size-criterion in case 1 (Fig. 6d) whereas
336 in case 2 (Fig. 6g) recruitment tended to be uniform over months. By contrast, in intermediate
337 layers (15 - 30 m) summer months for spawning favored recruitment when using an age-
338 criterion (Fig. 6b) and a size-criterion in case 2 (Fig. 6h), whereas in case 1 (Fig. 6e) recruitment
339 tended to become uniform. In deeper layers (30 - 45 m), summer favored recruitment in all
340 cases (Fig. 6c, f, i). When mortality was included, case 1 at the surface layer (0 - 15 m, Fig. 7d)
341 lead to fairly low and uniform recruitment whereas all other cases showed highest recruitment
342 in summer (Fig. 7a, b, c, e, f, g, h, i).

343 When a size criterion was used for recruitment (criterion 2), the corresponding age
344 at which individuals recruited was very variable, ranging from 20 to 90 days (Fig. 8). In Sim 5
345 (case 1), the coastal zone from 6° S to 9° S was the most favorable to early recruitment (Fig. 8a),
346 while in Sim 6 (case 2) the northernmost zone (2 - 3° S) showed the lowest ages at recruitment
347 (Fig. 8b). In case 1, recruitment started at an age of ~ 35 days for all spawning depth levels, and
348 peaked at a similar age of ~ 50 days (Fig. 8c). In case 2, individuals in the northernmost part of
349 the study domain recruited as early as ~ 20 days (Fig. 8b) and recruitment peaked at ages ~ 25 ,
350 ~ 35 and ~ 45 days for depth levels 0 - 15, 15 - 30 and 30 - 45 m, respectively (Fig 8d).

351 *Limitation by food*

352 A food limitation sensitivity test between Sim 7 (case 1) and Sim 5 (case 1) showed
353 that food acted as a growth limiting factor. This is particularly true during winter when mortality
354 was included (Supplementary material, Fig. S4a) and in the 14° - 16° S zone (Supplementary
355 material, Fig. S4b). Similar patterns were observed for case 2 (Sim 6 and Sim 8, Supplementary
356 material, Fig. S5).

357 The amount of larvae recruiting according to their spawning location was also very
358 variable along the coast, ranging from 0 to 150 ind/m² without mortality (Fig. 9a, c) and from
359 0 to 2 ind/m² cell with mortality (Fig. 9 c, d). For case 2 there were three spawning spots

360 favorable to recruitment in the north, center and south of the domain (Fig. 9c, d). For case 1 the
361 northern zone was no longer favorable but the central and southern zones remained (Fig. 9a, b),
362 which is more consistent with the spatial distribution of Peruvian anchovy egg density
363 (eggs/m²) derived from field surveys (Fig. 9e).

364 **4 Discussion**

365

366 We studied larval retention and growth of the Peruvian anchovy (*E. ringens*) in the
367 northern Humboldt Current system (NHCS) using a biophysical model. This model was first
368 forced by currents from a more modern configuration of a hydrodynamic model used previously
369 at the same horizontal resolution (10 km; Brochier et al. 2008). We were able to replicate the
370 general modeled patterns of larval retention obtained previously. Indeed, our results were
371 consistent with another study aiming at answering the same scientific question but using a
372 different dataset, thus proving the replicability of these results (*sensu* National Academies of
373 Sciences Engineering and Medicine, 2019). This emphasizes the robustness of the results despite
374 the stochastic variability inherent to hydrodynamic model configurations. It was also crucial to
375 replicate previous results before assessing the effects of new forcing products and other model
376 components in order to avoid generating false conclusions (Baker 2016). Here, in particular,
377 we found the same opposite seasonal pattern relative to spawning depth (Fig. 3) as Brochier et
378 al. (2008). However, we obtained slightly higher coastal retention values during summer
379 months for the three spawning depths considered. This result could be due to a greater
380 stratification of the water column and to a higher spatial resolution of the wind stress forcing
381 (weaker at the coast) in the new hydrodynamic simulations compared to the old one. Retention
382 within the most coastal spawning zone (bathymetry 0 -100 m) was up to 20% higher in summer
383 than in Brochier et al. (2008).

384 We then used a configuration at higher spatial resolution (2 km) and found that the
385 simulated patterns of coastal retention remained essentially the same (Supplementary material,
386 Fig. S1). The change in bathymetry source slightly impacted the retention values, more, in fact,
387 than increasing the resolution. This may be due to shrinking of the continental shelf retention
388 area in the lower resolution grid. However, these changes were too small to alter the general
389 spatio-temporal patterns, or the general relationship between simulated retention and spawning
390 depth or isobaths. Thus, studies focusing at the scale of the whole Peruvian continental shelf as
391 ours can be conducted with a 10 km grid resolution without risking to miss key hydrodynamic
392 features influencing the retention patterns. This result contrasts with previous studies, which
393 suggested that downscaling models in coastal ecosystems may lead to significantly higher
394 simulated retention rates (Swearer et al. 2019). However, it is in line with Vic et al. (2018) who

395 found stable dispersion patterns across model resolutions in the open ocean. The retention area
396 considered here extended quite offshore with a relatively shallow continental shelf and a
397 straight coast exposed to the open ocean, which might explain this result. Thus, despite
398 increasing resolution might impact retention very near the coastline, the mean retention over
399 the larger area considered here was not impacted which is in line with Garavelli et al. (2014),
400 who showed that between 3 km and 7.5 km hydrodynamic forcings, no difference in dispersion
401 distance was observed and both experiments also demonstrated that the closer to coast, the
402 greater the success of the individuals. Because our results were the same between 2 and 10 km
403 resolutions, we decided to keep the 10 km resolution in the subsequent simulations, which
404 allowed reduced computing time of the hydrodynamic model especially when coupled to the
405 biogeochemical module.

406 After including larval growth into our model based on DEB theory, we explored
407 simulations using a size criterion for retention (Fig. 4 and Fig. 5), as opposed to an age criterion
408 as before. Using a size criterion for retention means considering the impact of environmental
409 variability on the planktonic life duration (PLD), which is crucial in biophysical modelling
410 studies (Lett et al. 2010). Indeed, a shorter PLD, resulting from faster growth, is expected to
411 increase local retention and therefore recruitment. Larvae that grow quicker may also escape
412 predators, swim more efficiently and have therefore a better chance to survive (Houde 2008),
413 which was also explored in our results including mortality. In our simulations, larvae
414 experienced temperatures ranging from $\sim 17^{\circ}\text{C}$ in winter to $\sim 23^{\circ}\text{C}$ in summer. The effect of
415 temperature on growth depended on the hypothesis we made on the C_T function (Eqn. 5) as we
416 considered two temperature correction curves. Under the hypothesis of a max C_T at $\sim 23.4^{\circ}\text{C}$
417 (Case 1), the PLD could be as low as 20 days and the largest recruitment was found in summer
418 in the Guayaquil Bay. However, this bay lies at the northern limit of *E. ringens* distribution
419 (Calderón-Peralta et al. 2020), and large recruitment of Peruvian anchovy has not been observed
420 there to our knowledge. Tuning model parameters in order to fit a known distribution is a way
421 to study the ecological niche limits. For our model prediction to fit the spatial extent of the
422 observed spawning area (thus excluding the Guayaquil Bay, Fig. 9e), we had to change the C_T
423 function such that its maximum value occurs at $\sim 19^{\circ}\text{C}$. In this case the average PLD of
424 simulated recruited larvae was ~ 50 days, which is in the order of *in situ* and laboratory
425 observations (Palomares et al. 1987). Furthermore, Castillo et al. (2022) showed that the main
426 habitat temperature range of adult anchovy population was $16\text{-}24^{\circ}\text{C}$, which is consistent with
427 an optimal larval growth temperature around the middle of this range. However, the hypothesis

428 that temperature would be the main factor limiting larval growth for individuals in the
429 Guayaquil Bay should be challenged by new laboratory experiments designed to identify the
430 upper temperature limit for larval growth. Indeed, current experiments found the fastest growth
431 at 19°C for larval stages but did not investigate higher temperature values (Rioual et al. 2021).
432 Some preliminary results tend to indicate for juvenile stages reduction of ingestion rate from
433 21°C (*unpublished data*), which would impact the growth rate. So, more laboratory experiments
434 should be designed specifically to identify the shape of the C_T function for *E. ringens*. Our
435 results obtained with a 19°C maximum C_T (case 1) are also in line with Xu et al. (2013) who
436 found a rather adverse effect of inter-annual variability, specifically during the El Niño period,
437 where the number of days to reach recruitment increased and survival decreased considerably.

438 In simulations where food was considered as not limiting larval growth, we found
439 similar results as in simulations where both food and temperature were limiting. This result
440 contrasts with Thomas et al. (2016) who used a similar bio-energetic approach as ours to study
441 oyster larvae growth and recruitment in Polynesia. In a context where temperature variability
442 was much smaller (~28-29°C) they found that food limitation explained most of recruitment
443 variability. In our case, larval food limitation did not impact the seasonal pattern but it had a
444 small impact on the spatial pattern, suggesting an average higher food limitation south of the
445 Pisco upwelling cell (~14 – 15 °S), which is in line with a lower upwelling productivity
446 (Espinoza-Morriberón et al. 2017). Validation of simulated zooplankton fields is notoriously
447 difficult because the corresponding data is rare. In the northern Humboldt current system off
448 Peru, Aronés et al. (2016) studied zooplankton biomass data from 1961 to 2012 and reported a
449 maximum biomass occurring in spring, not fully matching our simulation that predict a
450 maximum in summer, but the data showed a considerable spatial and inter-annual variability,
451 and the seasonal differences were strongest during the first period of the study than after. Thus,
452 further zooplankton observations and models are needed to get a more precise idea of food
453 limitation for larval growth in the Humboldt.

454 The confirmation by laboratory experiments of a “smooth” temperature correction
455 function (as in case 1 of the present simulations) for *E. ringens* would be consistent with the
456 widely accepted idea that temperature is a limiting factor for anchoveta blooming (Chavez et
457 al. 2008). However, a steeper temperature correction (case 2) function would challenge this
458 idea. In this latter case, other factors responsible for the northern limit of *E. ringens* habitat,
459 possibly correlated with temperature, should be identified, as water masses (Bertrand et al.

460 2004; Swartzman et al. 2008), oxygen (Bertrand et al. 2011), or food quality (Ayón et al. 2008;
461 Calderón-Peralta et al. 2020) as food abundance was not found as a key limiting factor in our
462 study. The change in species dominance shown in sediment records, corresponding to periods
463 of environmental changes (Salvatteci et al. 2018, 2019), would then be more associated to
464 changes in stratification and circulation leading to a decrease in oxygen availability and/or
465 decrease in ichthyoplankton retention (Brochier et al. 2013; Espinoza-Morriberón et al. 2021),
466 affecting larval vertical distribution.

467 In Peru, small pelagic fish monitoring is based on spawning biomass estimation and
468 egg and larvae surveys (Pauly and Soriano 1987; Ayón 2000; Gutiérrez et al. 2012) without
469 explicitly accounting for spatial features (e.g. cross-shore and vertical). However, our results
470 shows that spatial and vertical distribution also largely impact the success of recruitment. We
471 suggest this information should be included in coupled model and observation operational
472 system, which allows to forecast the seasonal success of recruitment. Thus, spatial monitoring
473 of ichthyoplankton distribution should include assessment of vertical distribution. This can be
474 achieved using multinet or, for a faster processing of the information, in situ imaging system
475 that may allow a rapid processing (Orenstein et al. 2020).

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486

487 Highlights

488

- 489 • We developed an individual-based model including larval retention and a
490 Dynamic Energy Budget bioenergetic module.
- 491 • Results show that Peruvian anchovy larval growth accelerates with increasing
492 temperature, but the upper threshold is still not properly defined.
- 493 • Food availability limits anchovy growth and recruitment in southern Peru, only
- 494 • Spawning depth has a significant effect on Peruvian anchovy recruitment with a
495 seasonal modulation.

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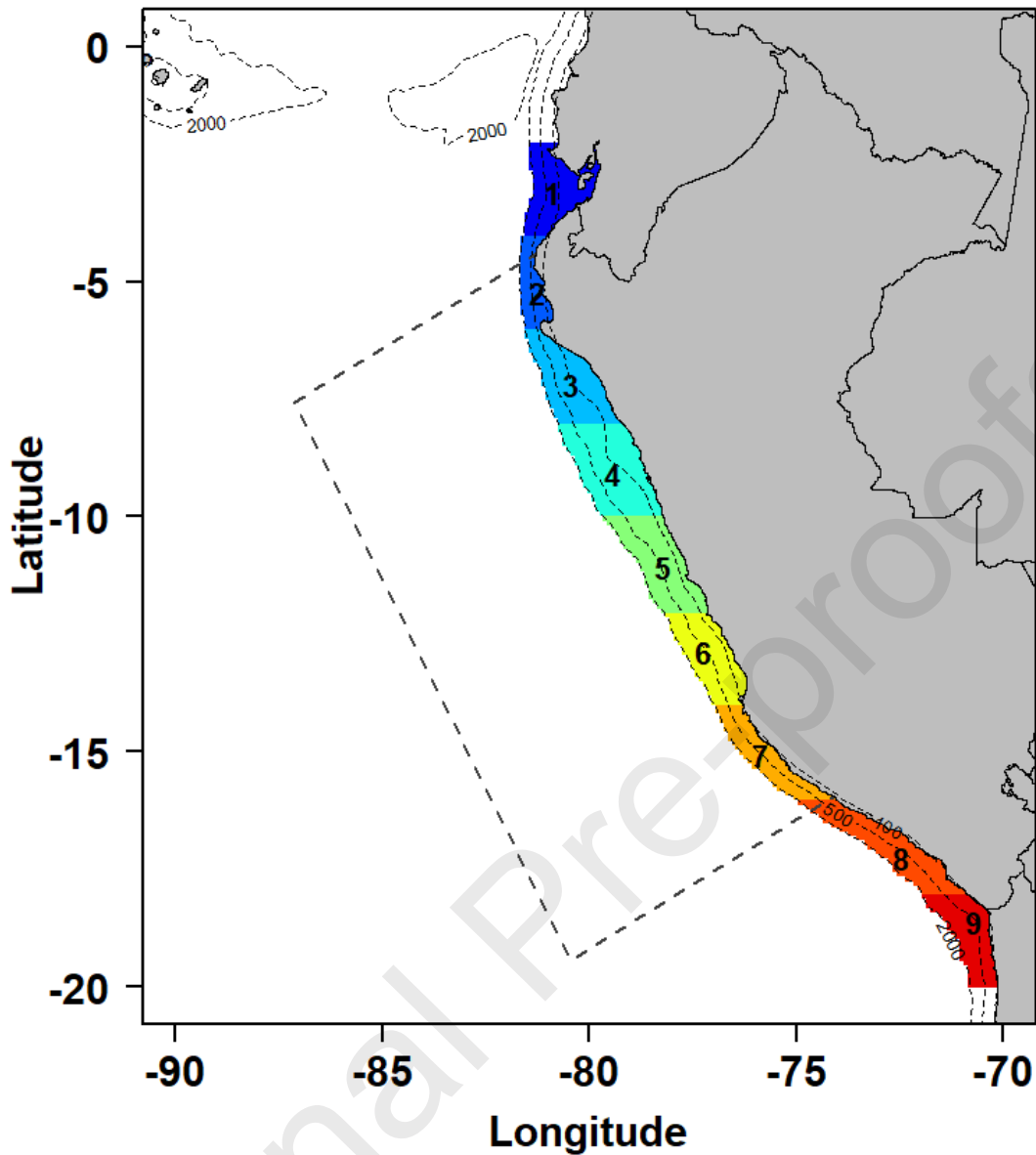
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747 **Highlights**

- 748 • We developed an individual-based model including larval retention and a
749 Dynamic Energy Budget bioenergetic module.
- 750 • Results show that Peruvian anchovy larval growth accelerates with increasing
751 temperature, but the upper threshold is still not properly defined.
- 752 • Food availability limits anchovy growth and recruitment in southern Peru, only
- 753 • Spawning depth has a significant effect on Peruvian anchovy recruitment with a
754 seasonal modulation.

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757 Fig. 1 Model domain at 10 km of spatial resolution (D01). The dotted rectangle represents the
758 nested model domain (D02) at 2 km resolution. Spawning areas (1 to 9) are every 2 degrees of
759 latitude between 2° S and 20° S. Three isobaths (100 m, 500 m and 2000 m) are shown.

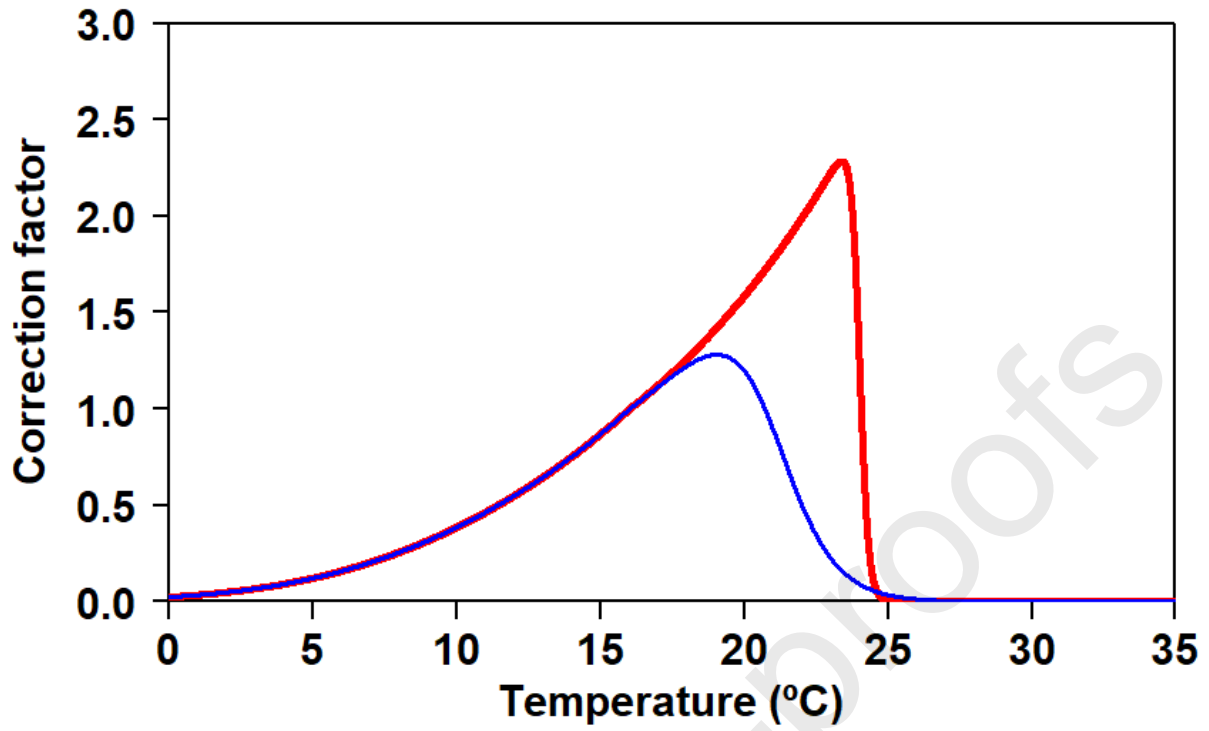


Fig. 2 Temperature correction curves for the metabolic flux in the dynamic energy budget model (equation 5); blue and red curve correspond respectively to case 1 and case 2 in Table 1.

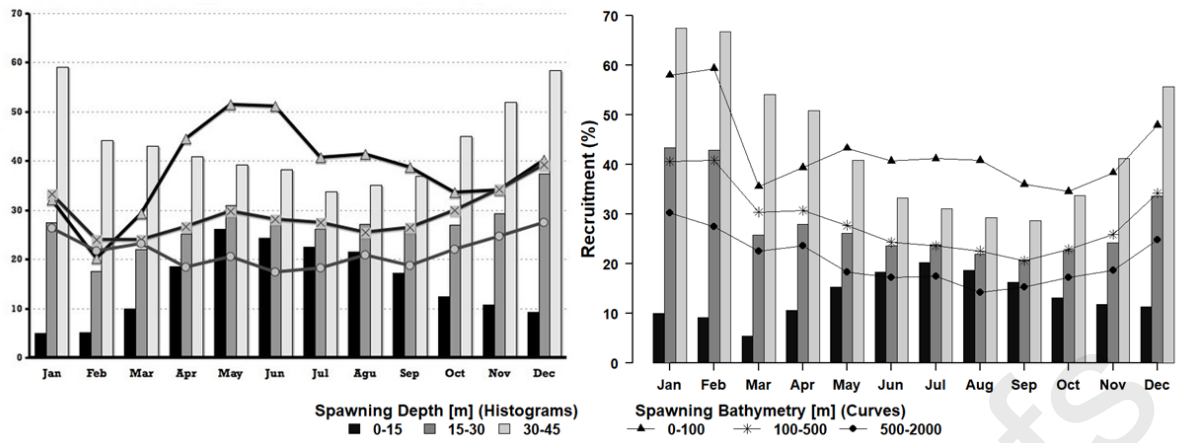


Fig. 3 Percentage of recruited larvae of Peruvian anchovy obtained for different spawning months, spawning depths, and isobaths delimiting spawning areas horizontally from (Brochier et al. (2008) (left) and from Sim 4 (right).

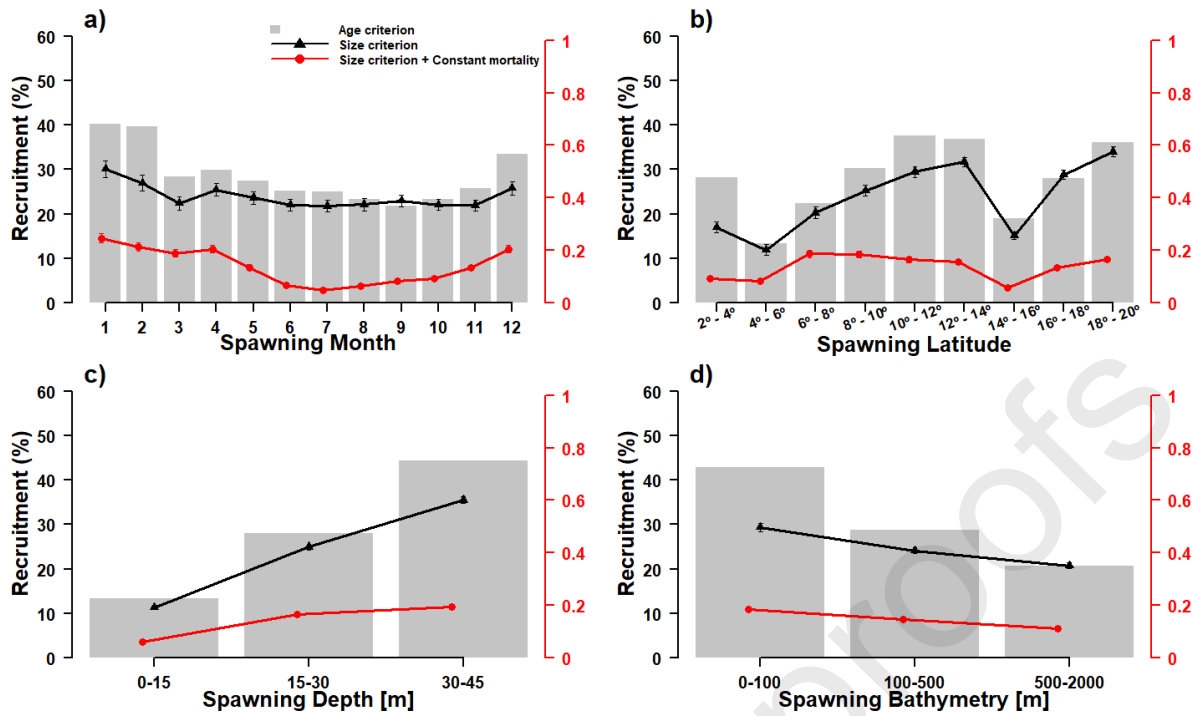


Fig. 4 Percentage of recruited larvae of Peruvian anchovy obtained for different a) spawning months, b) spawning latitudes, c) spawning depths and d) isobaths delimiting spawning areas horizontally using different criteria for recruitment (size criteria -black lines-, size criteria plus constant daily mortality -red lines-) in Sim 5. Recruitment values based on age criteria -grey bars- were taken from Sim 4.

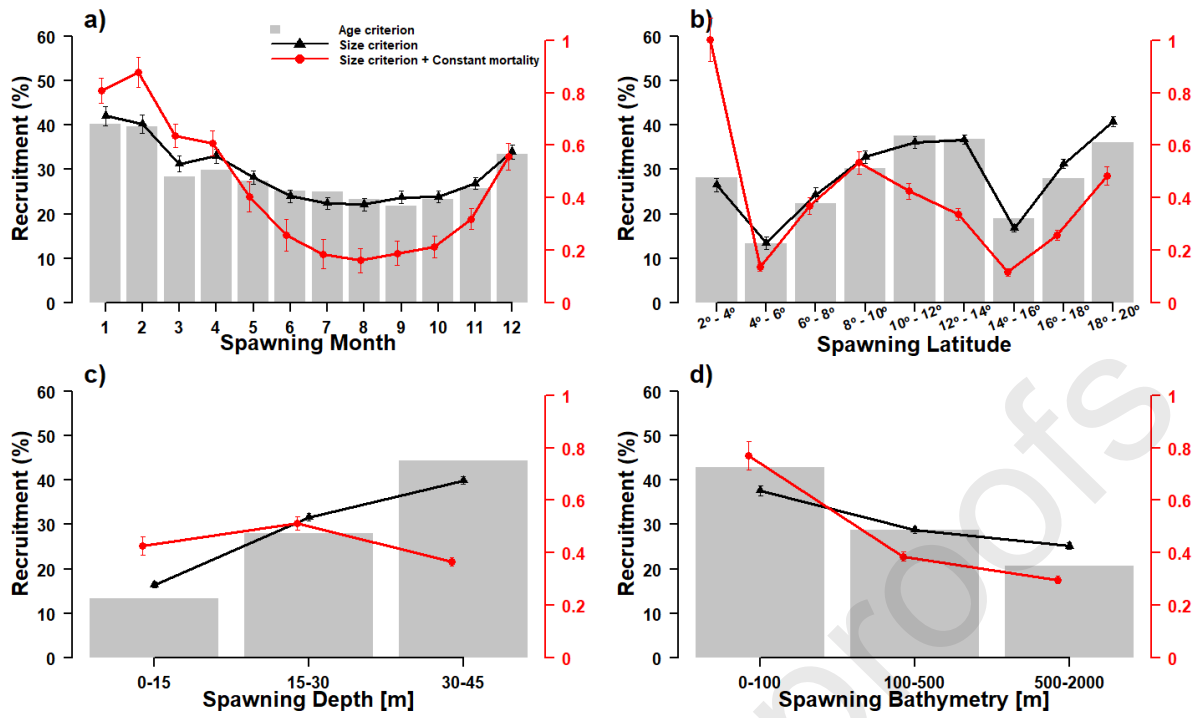


Fig. 5 Same as Fig. 4 in Sim 6.

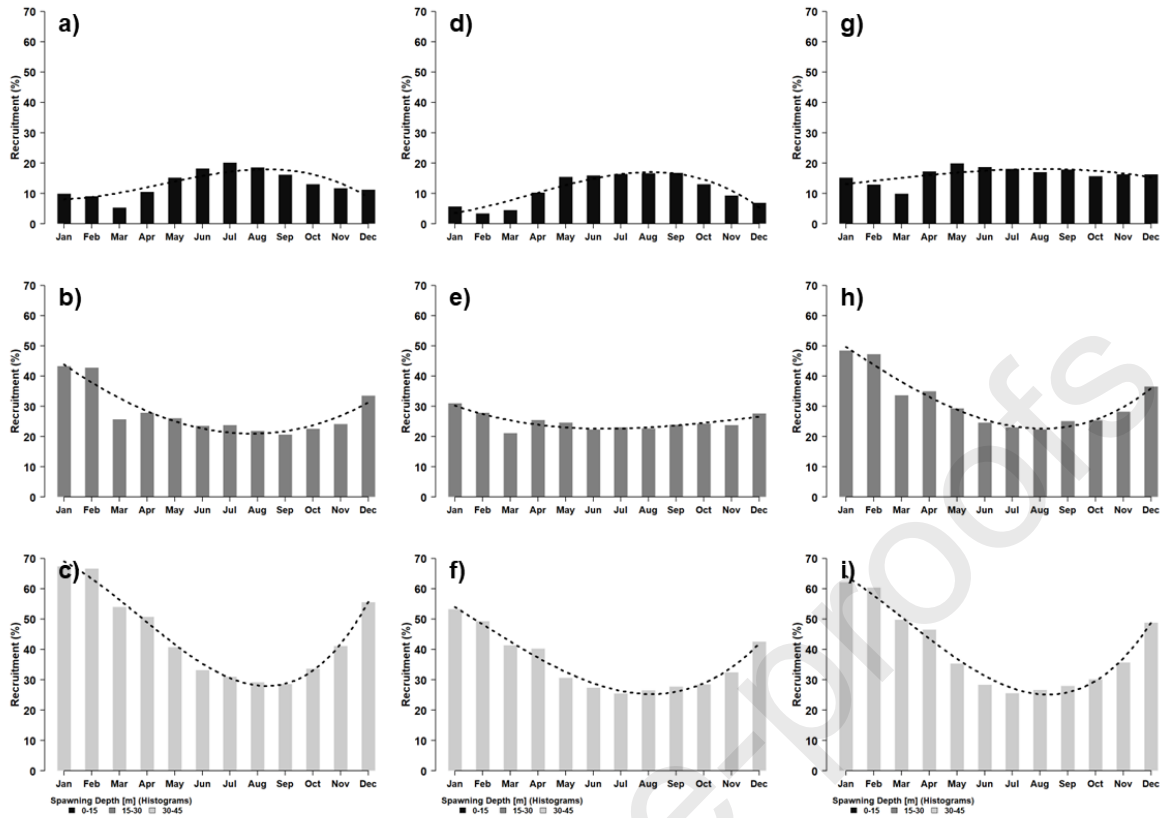


Fig. 6 Percentage of recruited larvae of Peruvian anchovy obtained for different spawning depths in Sim 4 criterion 1 (a, b, c), Sim 5 criterion 2 (d, e, f) and Sim 6 criterion 2 (g, h, i). Spawning depth is (a, d, g) 0 - 15 m, (b, e, h) 15 - 30 m, (c, f, i) 30 - 45 m. The dotted curves are third degree polynomial models fitted to the recruitment patterns.

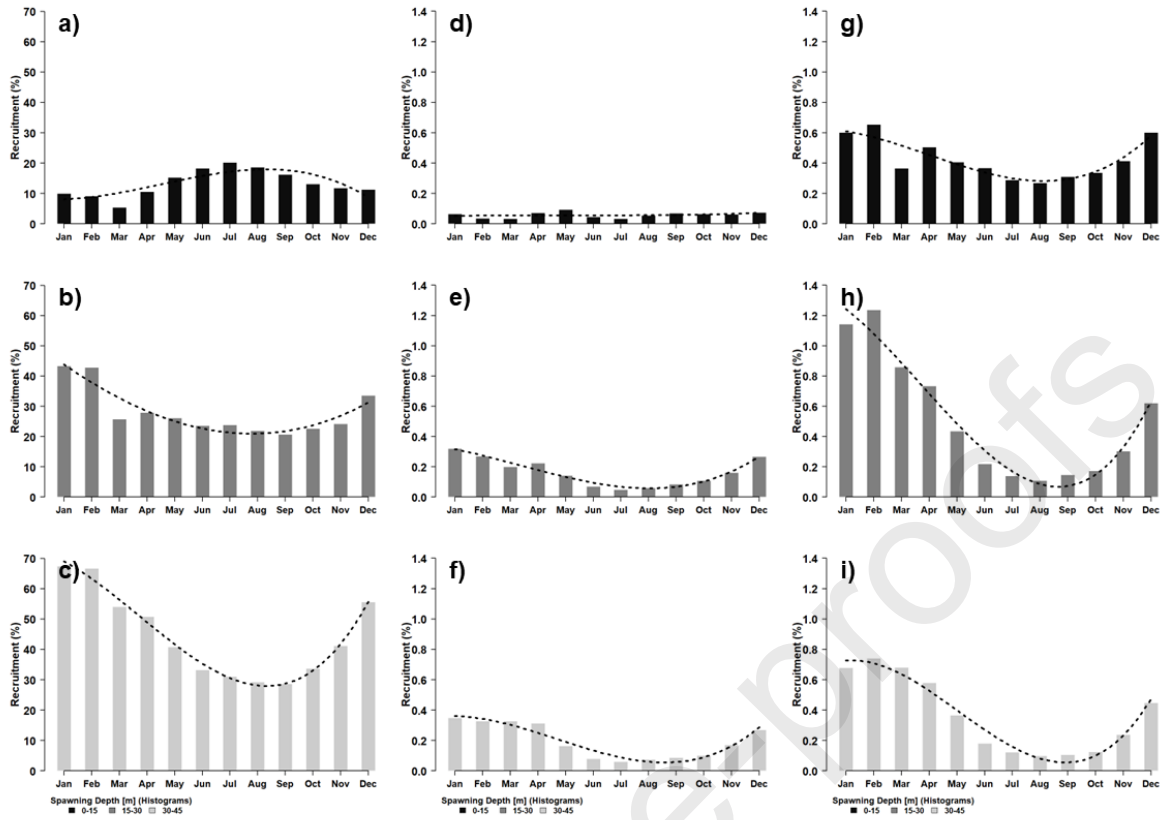


Fig. 7 Same as Fig. 6 but with mortality included in Sim 5 and Sim 6 (criterion 3).

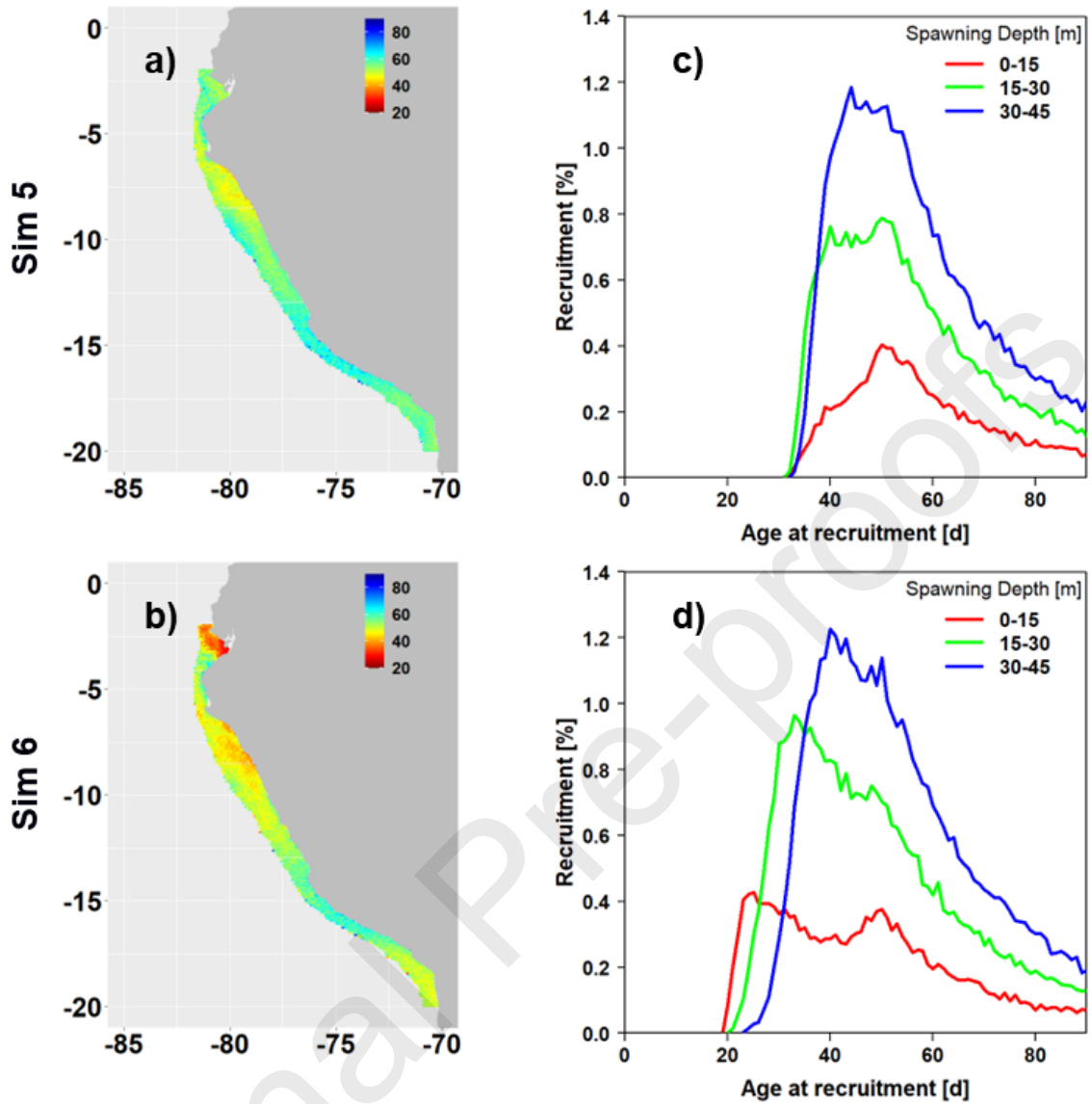


Fig. 8 Spatial distribution of average age at recruitment (a, b) and mean daily recruitment depending on age (c, d), for 0 - 15 m (red line), 15 - 30 m (green line) and 30 - 45 m (blue line) spawning depth. Results obtained with (a, c) Sim 5 and (b, d) Sim 6 using recruitment criterion 2 (no mortality).

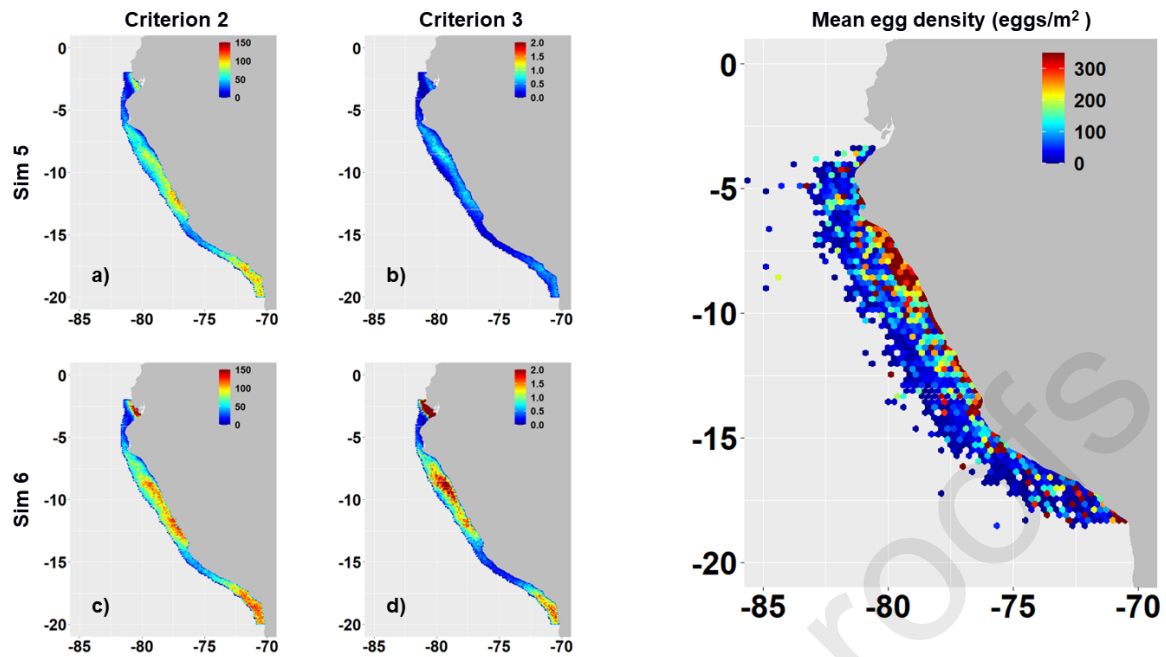


Fig. 9 Spatial distribution of the average number (ind/m²) of simulated Peruvian anchovy larvae recruiting according to their spawning location obtained with (a, b) Sim 5 and (c, d) Sim 6 using recruitment criterion 2 (no mortality) (a, c) and 3 (with mortality) (b, d). (e) Spatial distribution of Peruvian anchovy mean egg density (eggs/m²) derived from IMARPE field surveys from year 1961 to 2016. Note that cell grid was 0.1° x 0.1° in a - d and 0.25° x 0.25° in e).

Table 1: Summary of simulations performed to study recruitment predictions sensitivity. This table list all parameters that differ between simulations.

	Sim 1	Sim 2	Sim 3	Sim 4	Sim 5	Sim 6	Sim 7	Sim 8
Figures	S2	S2	S2	3,4,5,6,7, S3	4,6,7,8,9, S4	5,6,7,8,9, S5	S4	S5
Configuration domain	D01	D02s	D02r	D01	D01	D01	D01	D01
Forcing type	physical	physical	physical	physical & biogeochemical	physical & biogeochemical	physical & biogeochemical	physical & biogeochemical	physical & biogeochemical
Bathymetry	STRM30	Interpolated from Sim1	STRM30	STRM30	STRM30	STRM30	STRM30	STRM30
Horizontal grid resolution	10 km	2 km	2 km	10 km	10 km	10 km	10 km0	10 km
Latitudinal spawning range	6° S - 14° S	6° S - 14° S	6° S - 14° S	2° S - 20° S	2° S - 20° S	2° S - 20° S	2° S - 20° S	2° S - 20° S
Growth sub-model	No	No	No	No	Yes	Yes	Yes	Yes
Recruitment criterion evaluated*	1	1	1	1	2 and 3	2 and 3	2 and 3	2 and 3
Correction Temperature**	--	--	--	--	Case 1	Case 2	Case 1	Case 2
Food half saturation constant (K)	--	--	--	--	1.6 μmolCL^{-1}	1.6 μmolCL^{-1}	0 μmolCL^{-1}	0 μmolCL^{-1}
Mortality	--	--	--	--	Yes	Yes	Yes	Yes

*Recruitment criterion 1: retention at 30 days; 2: retention at 20 mm; 3: retention at 20 mm with constant mortality.

**Temperature correction factor case 1: $T_H = 294 K (= 21^\circ C)$ and $T_{AH} = 95\ 000 K$; case 2: $T_H = 297 K (= 24^\circ C)$ and $T_{AH} = 570\ 000 K$

760 Table 2: Parameters used for the bioenergetic model describing larval growth. These values
 761 were estimated by Pethybridge et al. (2013) for *Engraulis encrasicolus*, except half saturation
 762 constant (K), estimated for the current configuration and fixed at $1.6 \mu\text{mol C L}^{-1}$. Comparison
 763 with data showed that these parameters allowed to reproduce *Engraulis ringens* larval growth.
 764 The values of T_L , T_H , T_{AL} , T_{AH} are detailed for case 1 and case 2 respectively (see section 2.7).

Primary parameters (rates are at reference temperature $T_1 = 289 \text{ K} (= 16^\circ\text{C})$)			
Symbol	Value	Unit	Definition
L_h	0.28	cm	Hatch length
L_b	0.35	cm	Yolk-sac to feeding larva threshold
T_A	9800	K	Arrhenius temperature
T_L	279/279	K	Lower temperature boundary
T_H	294/297	K	Upper temperature boundary
T_{AL}	20 000/20 000	K	Arrhenius temperature for lower boundary
T_{AH}	95 000/570 000	K	Arrhenius temperature for upper boundary
K	1.6	$\mu\text{mol C L}^{-1}$	Half-saturation constant
κ_X	0.71	-	Fraction of food energy fixed in reserve
$\{\dot{p}_{Xm}\}$	325	$\text{J.cm}^{-2}.\text{d}^{-1}$	Maximum surface-specific ingestion rate
$[E_m]$	2700	J cm^{-3}	Maximum reserve density
$[E_G]$	4000	J.cm^{-3}	Volume-specific costs of structure
$[\dot{p}_M]$	48	$\text{J.cm}^{-3}.\text{d}^{-1}$	Volume-specific somatic maintenance rate
κ	0.7	-	Fraction of mobilized reserve allocated to growth and somatic maintenance
Auxiliary and Compounds parameters			
Symbol	Value	Unit	Definition
δ_M	0.152	-	Shape coefficient
$\{\dot{p}_{Am}\}$	$\kappa_X\{\dot{p}_{Xm}\}$	$\text{J.cm}^{-2}.\text{d}^{-1}$	Maximum surface-area-specific assimilation rate

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768 **Declaration of interests**

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770 The authors declare that they have no known competing financial interests or personal
771 relationships that could have appeared to influence the work reported in this paper.

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773 The authors declare the following financial interests/personal relationships which may be
774 considered as potential competing interests:

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