

A spatially-explicit population dynamics and stock assessment model driven by environmental variables

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SEAPODYM introduction



Spatial Ecosystem And Population Dynamics Model



An hybrid modeling framework to combine:

- the progress in ecological sciences to describe ocean pelagic ecosystem and species habitats and behaviour;
- The progress in ocean (physical and biogeochemical) modeling;
- the progress achieved in stock assessment models for quantitative estimation of population dynamics parameters models (MLE).



SEAPODYM introduction



(2.1)

(3.1)

Spatial Ecosystem And Population Dynamics Model

Zoo, unekton

T°, U&V, PP, Zeu, O2,



2D – fish population dynamics (ADR)

+ fisheries & parameter estimation (spatial resolution from 2° to 9 km) habitats + movements Integrated over day-night conditions in the 3 vertical layers



Micronekton (forage)

E'₁

E'n



Temperature; currents Primary production













I - SEAPODYM Fish spatial dynamics modelling

- Feeding Habitat
- Movement
- Spawning habitat and recruitment
- II Applications to tuna
 - Parameters and stock estimates
 - · Environmental variability
 - Impact of Climate change



SEAPODYM-Fish







SEAPODYM-Fish Feeding Habitat



Ha= index accounting for the abundance of different groups of forage and their accessibility in the vertical layer inhabited (day and night changes)



Biomass in layer z due to Biomass in layer z due to migrant migrant groups present Biomass in layer z due groups present during the night $(1-\tau)$ during the day (τ) to non-migrant group $\begin{pmatrix} F_{11} & 0 & 0 \\ F_{21} & F_{22} & 0 \\ F_{31} & F_{32} & F_{33} \end{pmatrix}$ $H_{a} = \sum_{z=1}^{3} \Theta_{a,z} \left(F_{zz} + \tau \sum_{k \neq z}^{\Psi} F_{zk} + (1 - \tau) \sum_{k \neq z}^{\Psi} F_{kz} \right)$ $\Theta_{a,z} = P(O_z; \boldsymbol{\theta}_{O,a}) \cdot G(T_z; \boldsymbol{\theta}_{T,a})$ Coefficient of accessibility in the concerned layer as a function of oxygen and temperature 20 Temperatu O2 (ml/l)



SEAPODYM-Fish Feeding Habitat



Feeding habitat of loggerhead turtles





SEAPODYM - Fish Movement

Population resulting



Rules of movements (D_a, U_a, V_a) based on H_a

- Habitat is null (no gradient) Diffusion rate is maximum Advection minimum
- Habitat is high (no gradient) Diffusion rate is minimum Advection minimum
- Habitat is low (high gradient) Diffusion rate is low Advection is high
- Habitat is medium (gradient)
- Diffusion % with Habitat value Advection % with Habitat gradient





Diffusion:



 D_0 : mean theoretical diffusion rate for a null habitat $D_0(\mathbf{a}) = \overline{V}_a^2 \frac{\Delta t}{4} \quad \overline{V}_a = L_a s^{-1}$

 $D_{a} = \sigma D_{0}(1 - cH_{a}^{3})$

2 parameters: σ and c

Advection: Passive (currents) + active (swimming)

$$U_a = \hat{u} + \chi_a \frac{\partial H_a}{\partial x}, \ V_a = \hat{v} + \chi_a \frac{\partial H_a}{\partial y},$$

$$\chi_a = V L_a{}^A$$

2 parameters - V: velocity (BL/s) at maximal habitat gradient - A: slope coefficient

ONLY 4 parameters to define the movement of all cohorts



SEAPODYM - Fish Movement







SEAPODYM Fish - Spawning & Recruits



Holling-III Gaussian Spawning habitat H_s Log-Normal micronk $H_s = f_1(prey) f_2(T^\circ) f_3(predator)$ Χ Χ zoopk Larvae recruits N(0,t,x)Prev Water Predators density Temperature density $N(0,t,x) = H_s \frac{R\widehat{N}}{1+h\widehat{N}}$ with \widehat{N} Number of mature fish

Then recruited larvae drift with currents; Average mortality coefficient + a range of variability $\propto Hs$





Applications: Param. Estimation



θ Description

Spawning habitat and Reproductiv

- $\overline{\sigma}_0\,$ standard deviation in temperature Gaussian function at age 0, $^\circ C\,$
- $T_0^\star~$ optimal surface temperature for larvae, $^\circ C$
- α_P prey encounter rate in Holling (type III) function, day^{-1}
- $\alpha_F~$ Log-normal mean parameter predator-dependent function, g/m^2
- $\beta_F~$ Log-normal shape parameter in predator-dependent function
- R reproduction rate in Beverton-Holt function, mo^{-1}
- -b slope parameter in Beverton-Holt function, nb/km^2

Mortality

- \bar{m}_p predation mortality rate age age 0, mo^{-1}
- β_p slope coefficient in predation mortality
- \bar{m}_s senescence mortality rate at age 0, mo^{-1}
- β_s slope coefficient in senescence mortality
- variability of mortality rate with habitat index $M_H \in (\frac{M}{(1+\epsilon)}, M(1+\epsilon))$

$Feeding \ habitat$

- $T_0~$ optimal temperature (if Gaussian function), or temperature range for the first young cohort, $^\circ C$
- T_K optimal temperature (if Gaussian function), or temperature range for the oldest adult cohort, $^\circ C$
- γ slope coefficient in the function of oxygen)
- \hat{O} threshold value of dissolved oxygen, ml/l
- ${\cal E}_{11}$ contribution of epipelagic for age to the habitat
- E_{22} contribution of mesopelagic forage to the habitat
- E_{21} contribution of migrant mesopelagic forage to the habitat
- ${\cal E}_{33}$ contribution of bathype lagic forage to the habitat
- ${\cal E}_{32}$ contribution of migrant bathypelagic for age to the habitat
- E_{31} contribution of highly migrant bathypelagic forage to the habitat

Movement

- V velocity at maximal habitat gradient and A = 1, BL/s
- A slope coefficient in allometric function for tuna velocity $\overline{V}^2 \wedge T$
- σ multiplier for the theoretical diffusion rate $\frac{\bar{V}^2 \Delta T}{4}$
- c coefficient of diffusion variability with habitat index

26 parameters + 3 or 4 par by fishery (growth, age at maturity from independent studies).

The Maximum Likelihood Estimation approach has been implemented in SEAPODYM to include:

- spatially disaggregated catch and size frequency data (Senina et al. 2008)
- Spatially disaggregated catch (Senina et al 2016)
- Acoustic biomass estimates, by analogy to a pseudo fishery without mortality (Dragon et al. 2015)
- Tagging data (Senina et al 2017; in rev.)

 10^{-10}



Applications: Param. Estimation









For tagging data

- the choice was to use only data from recaptures to be independent of fisheries
- this information contributes mainly to the estimation of movement and habitats parameters
- Data are aggregated by time strata, e.g. quarter, and distributions smoothed using Gaussian kernels to reduce the errors related to recapture date and positions.
- More weight to longer time of liberty at sea
- The same model of A&D is used without considering mortality.

Senina I., Lehodey P., Calmettes B., Nicol S., Caillot S., Hampton J. and P. Williams (2016). Predicting skipjack tuna dynamics and effects of climate change using SEAPODYM with fishing and tagging data. WCPFC, 12th Regular Session of the Scientific Committee, Bali, Indonesia 3–11 August 2016, WCPFC-SC12-2016/EB WP-01: 71 pp. http://www.wcpfc.int/node/27443



Applications: Stock estimates



Pacific Skipjack

Physical & biogeochemical forcing:

NEMO-PISCES **2° x month** driven by atmospheric reanalysis INTERIM **1979-2010**

Age structure:

36 monthly cohorts + one (age+). Age-length and age-weight from MULTIFAN-CL estimate (Rice et al., 2014).

Fishing:

Revised SPC + IATTC (2013) datasets (6 pole-and-line, 9 purse-seine and 2 longline fisheries)

Tagging data:

Tags recaptured between May 2008 and Dec 2010





Applications: Stock estimates







Applications: Stock estimates



Pacific Skipjack Impact on habitats & dynamics

Recruited larvae (nb/ km²) 1980-2010







Applications: Environmental Var.









CC projections: Ensemble simulation to account for uncertainty





Applications: Climate Change



I - Optimization

Comparison of predicted distributions for 1st and last decade of the historical time series. Total observed catches are shown with catch proportional to circles (same scales between decades).

Skipjack: WCPFC SC12 & Senina et al. (sub.)

Yellowfin: WCPFC SC13

Bigeye: WCPFC SC 14

South Pacific Albacore: WCPFC SC14









II – Ensemble simulation

Uncertainty explored in the simulation ensembles produced for this study

Uncertainty in atmospheric forcing		Structural uncertainty	
Code	CMIP5 model	in biogeochemical model	SEAPODYM
IPSL	IPSL-CM5A-MR (Institut Pierre Simon Laplace, France)	 Primary production: Increase of PP by 10% (PP10) in tropical waters (defined by SST >27°C) Dissolved Oxygen: No change (O2clim) = Use of climatological fields 	 Genetic adaptation: Regular increase in optimal spawning temperature to reach + 2°C at the end of the Century Ocean acidification: Additional mortality on larvae based on laboratory experiments with low medium and high sensitivity to pH (available only for yellowfin).
MIROC	MIROC-ESM (Model for Interdisciplinary Research on Climate, Japan)		
NorESM	NorESM1-ME (Norwegian Climate Centre, Norway)		
MPI	MPI-ESM-MR (Max Planck Institute for Meteorology, Germany)		
GFDL	GFDL-ESM2G (Geophysical Fluid Dynamics Laboratory, USA)		

Ensemble of 20 members for SKJ, BET and ALB projections and 35 for YFT (with additional Ocean acidification scenarios)



Applications: Climate Change



II – Ensemble simulation (skipjack)



Senina et al. (2018) Impact of climate change on tropical tuna species and tuna fisheries in Pacific Island waters and high seas areas 14th Scientific Committee of the WCPFC, Busan, South Korea, 8-16 Aug. 2018. WCPFC-SC14-2018/EB-WP-01, 44 pp. <u>https://www.wcpfc.int/node/30981</u>



Conclusions



- SEAPODYM simulates spatial structure & spatial dynamics of exploited pelagic fish species
- MLE approach at the resolution of the model => hundred thousands of data are assimilated
- Recent integration of tagging data in the MLE has greatly improved skipjack optimization
- With fish dynamics driven by environmental variables, the model can help to understand (forecast?) natural climate variability (e.g., ENSO) and impacts of climate change
- The results are sensitive to the quality of environmental forcings
- Ensemble simulations help to account for uncertainty
- Multiple possible applications, spatial management, connectivity study, operating model, ...







www.seapodym.eu

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