

# *Workshop on Dynamics of Pelagic Fish in the North Pacific under Climate Change*

*Yeosu, Korea 16 Oct 2014*

## **An ecosystem and optimization framework for fish population dynamics assessment under the influence of fishing and climate**

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<sup>5</sup> Pacific Islands Fisheries Science Center, Ecosystems and Oceanography Division, Honolulu, USA.

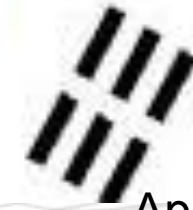
Introduction to SEAPODYM  
(Spatial Ecosystem  
And Population  
Dynamics Model)



Micronekton and  
habitats  
Environmental forcing  
variables



Optimization  
framework for parameter estimation  
(Maximum Likelihood Estimation)  
Stock assessment

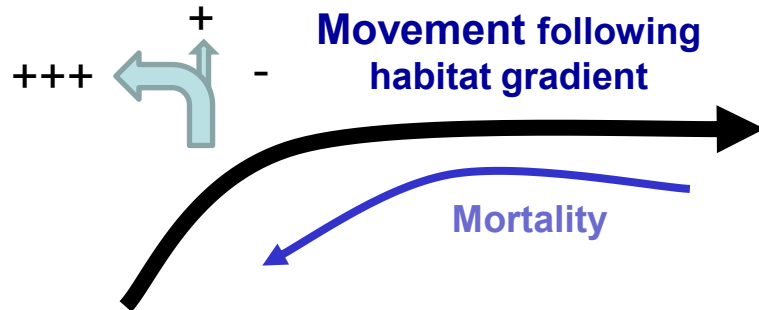


Applications to Climate  
Change impacts

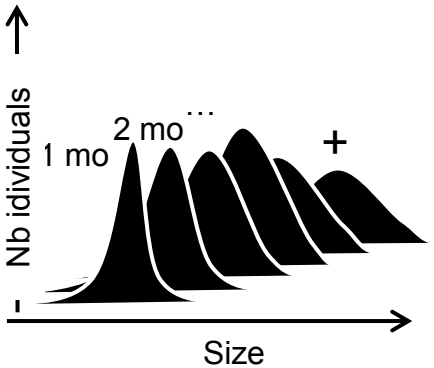
Spatial dynamics based on advection - diffusion equations with movement % to fish size and linked to habitats

Predator (e.g., tuna) population

(Lehodey et al 2008 Prog Oceanog)

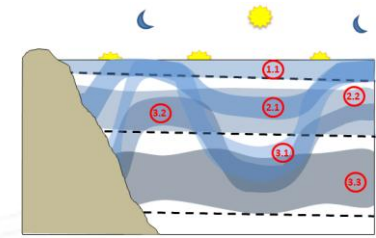


**Feeding Habitat =**  
Prey abundance (micronekton)  
x accessibility ( $T^\circ, O_2$ )



**Growth**  
**mortality** } **By cohort**

**IF MATURE**  
seasonal  
switch



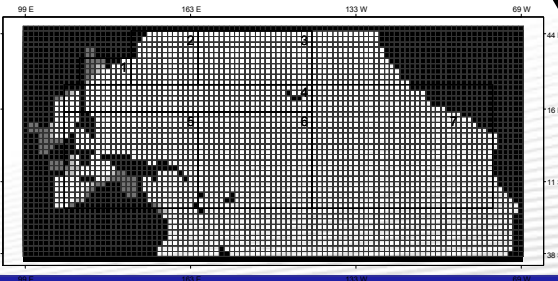
**Simplified 3D vertical structure**

**Population: age structured over a regular grid**

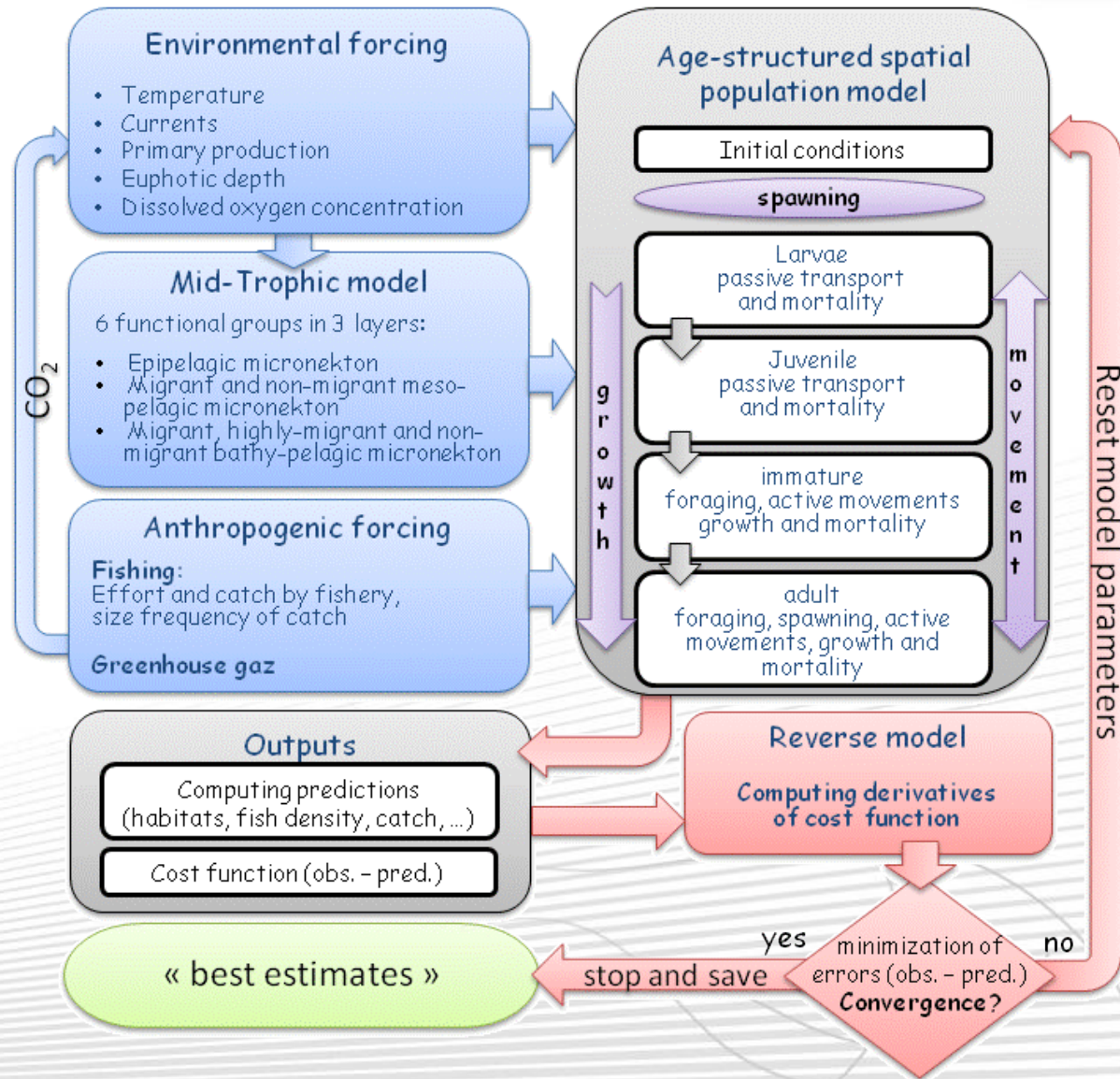
**Spawning Habitat =**  
**Temperature**  
**Prey of larvae (zoopl.)**  
**Impact of predation (micronekton)**  
**Accessibility ( $T^\circ, O_2$ )**

**Local Larval Stock-Recruitment**

**Drift in current**

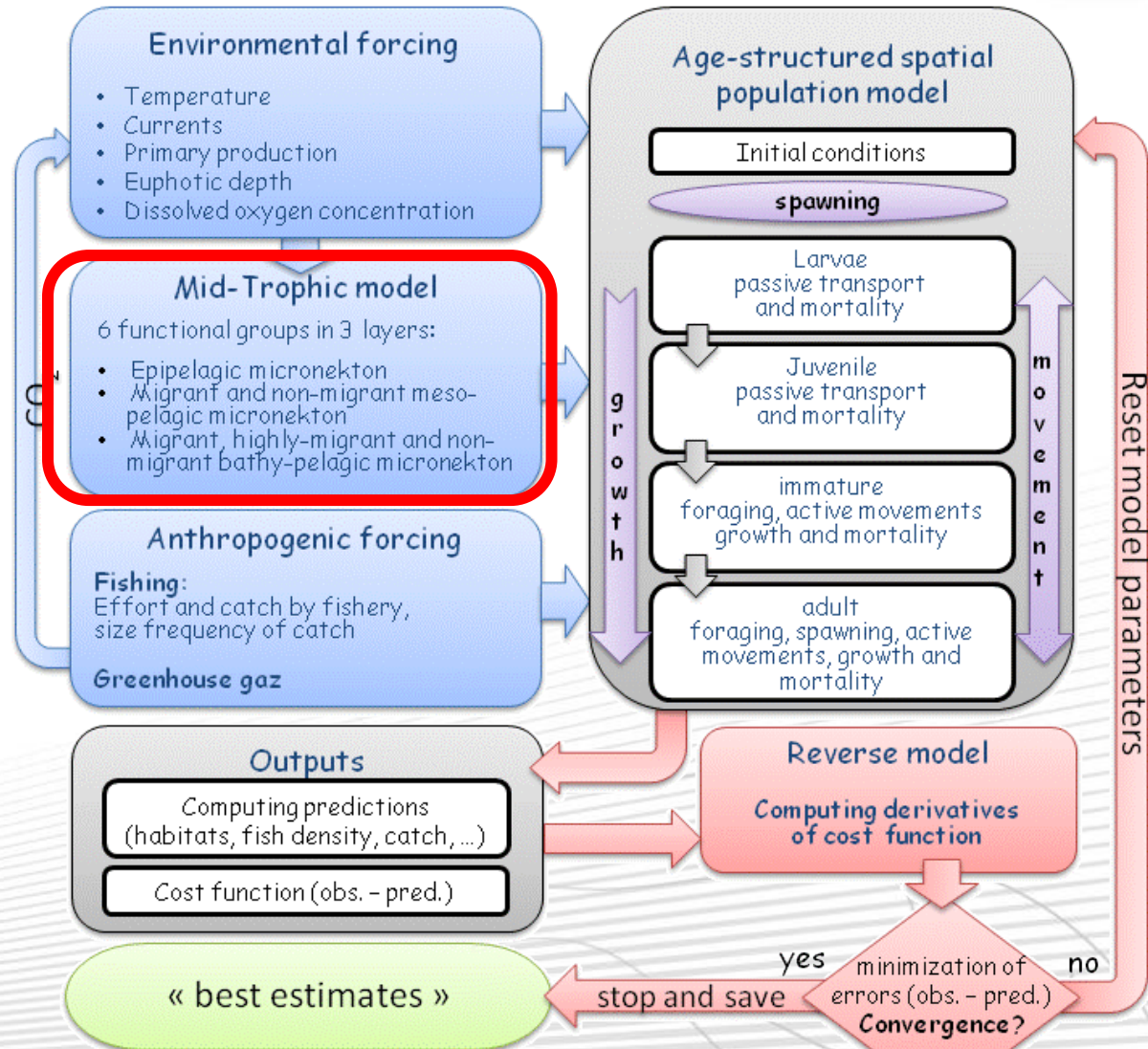


- SEAPOODYM uses ocean bio-physical variables characterizing the environment of predator species to drive the population dynamics of this (these) predator(s).
- An original sub-model describes zooplankton and 6 functional groups of prey organisms (micronekton at the mid-trophic level: MTL)
- Observed effort by fisheries is used to predict catch and Maximum Likelihood Estimation approach fit the catch (and size frequency of catch) to observation to get the best set of parameters





## MICRONEKTON

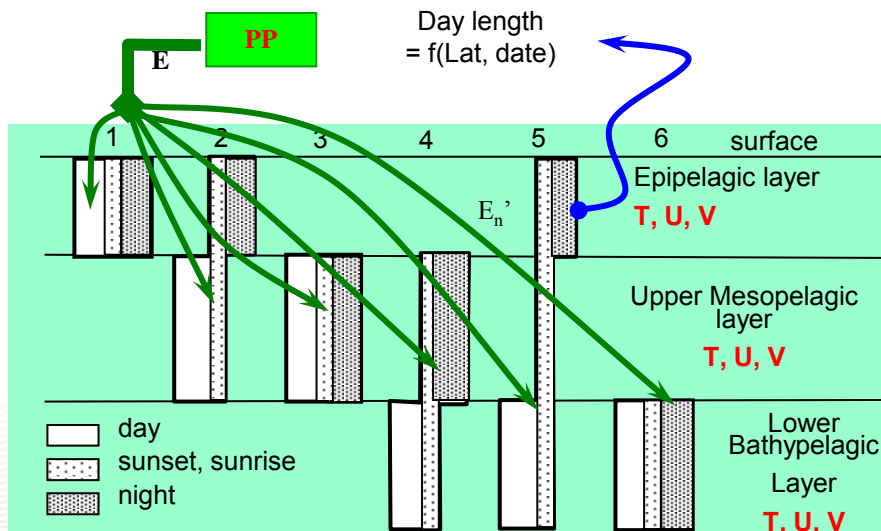
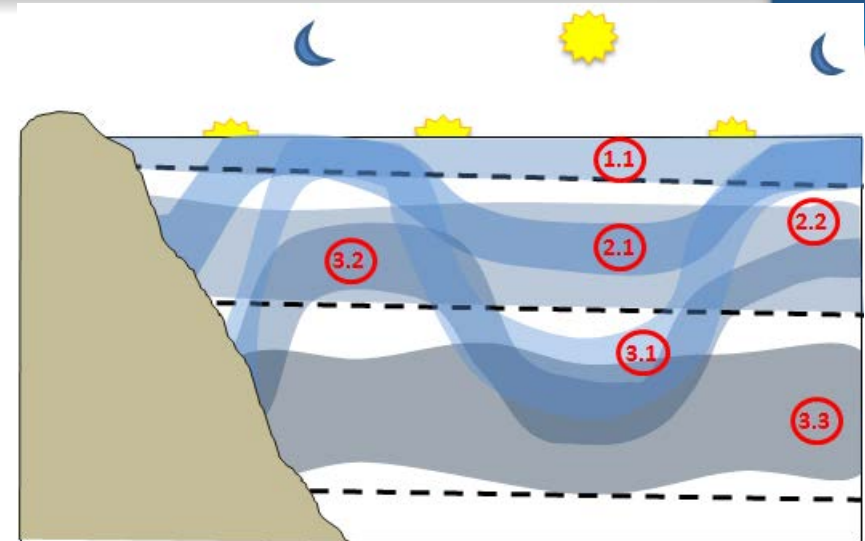
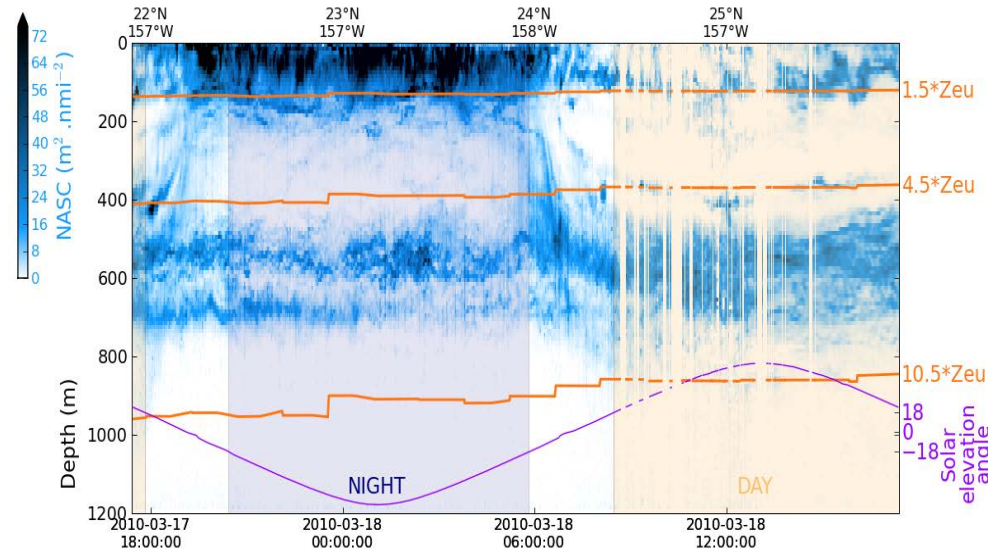


### STRENGTH:

Micronekton (the prey of adults and predators of larvae) is THE KEY component of the system !

### WEAKNESS:

Wrong micronekton modeling = wrong predator simulation and fit to data based on spurious parameterisation

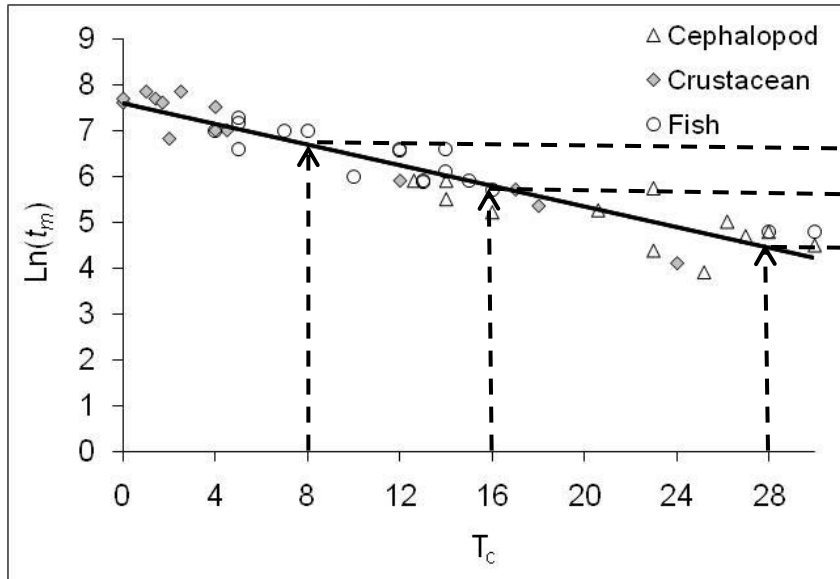


## A model of micronekton

6 functional groups in 3 vertical layers. Three components exhibit diel vertical migrations, transferring energy from surface to deep layers.

The source of energy is the primary production PP.

**6 parameters (  $E$ ;  $E'_n$  ) to calibrate**



Age at recruitment:

Age at maturity:

199 d

797 d

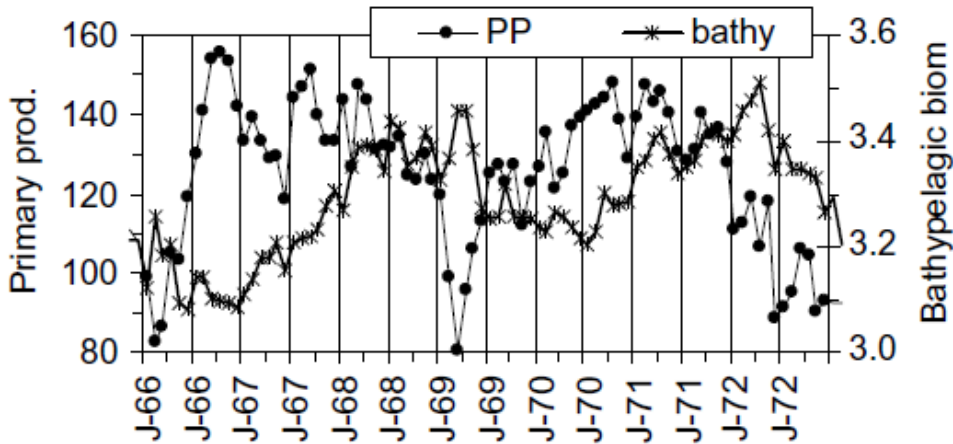
64 d

255 d

22 d

88 d

Time of development in days (Log scale) of mid-trophic organisms until age at maturity ( $t_m$ ) in relation to their ambient habitat temperature  $T_c$

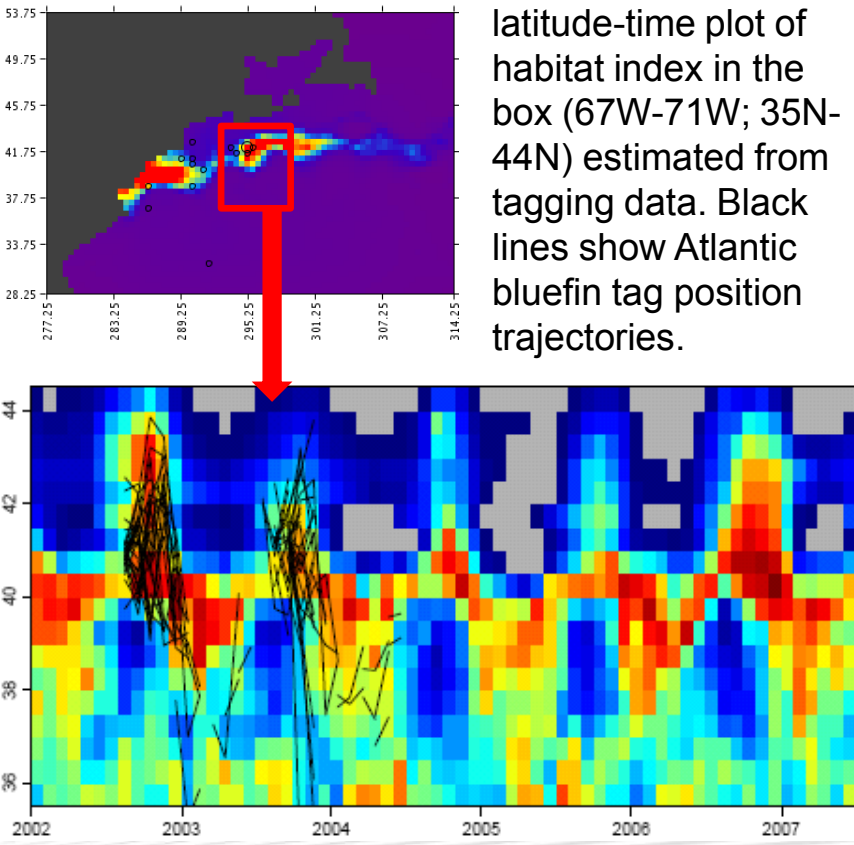


**As a consequence the dynamic of biomass of MTL in deep (cold) layers can be decoupled of surface primary production**

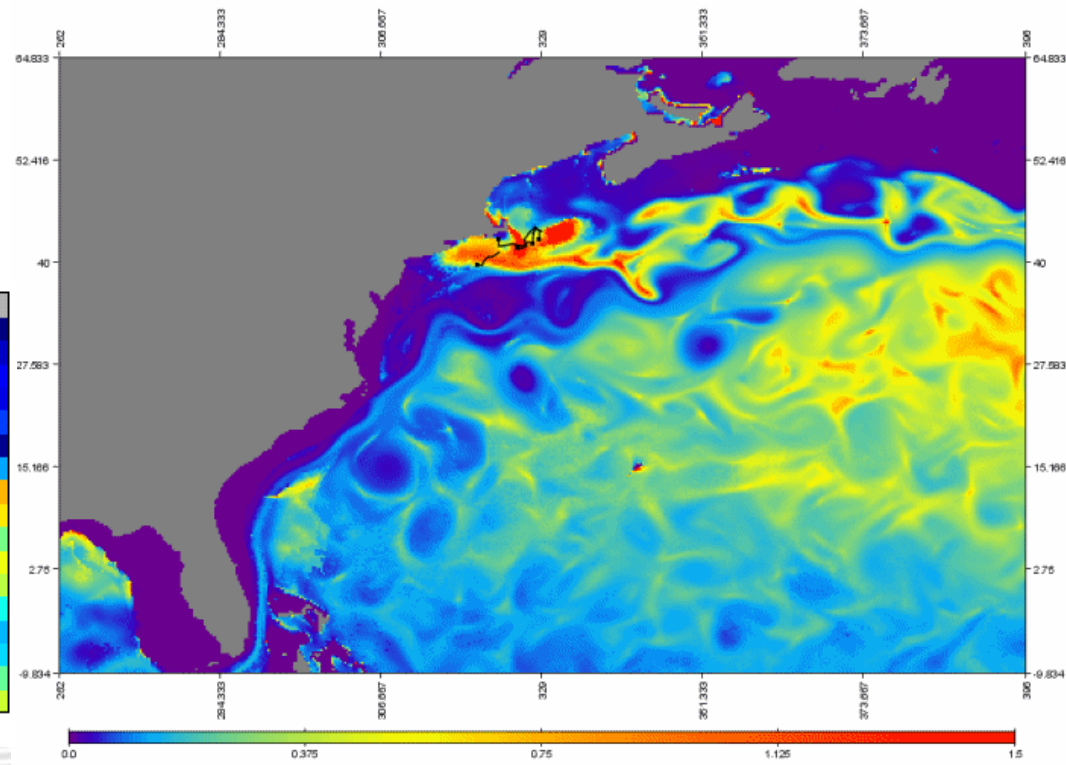
Box: 5°N-5°S; 120°W-100°W

## Ex.: Feeding habitat of Atlantic bluefin tuna

The model predicts the seasonal and interannual variability of feeding habitat in the North-East Atlantic where fish were released with PSATs



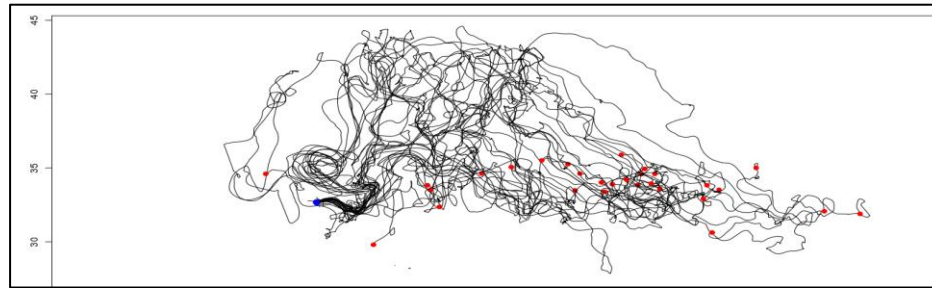
Electronic tagging data kindly from M. Lutcavage



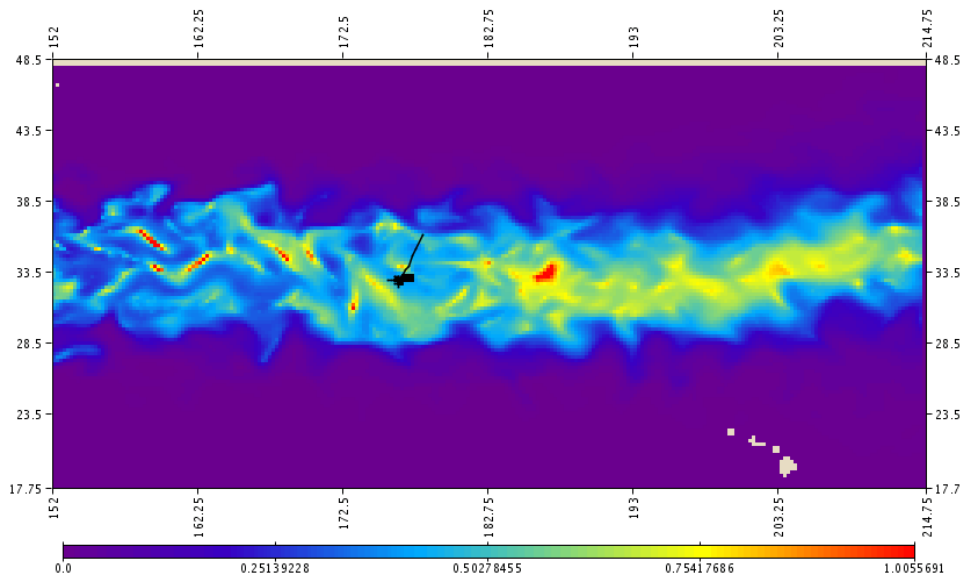
ibM\_HH\_dym  
2002 July 27



## Ex.: Feeding habitat of Loggerhead turtles in the north Pacific



Observed displacements between May 2005 and May 2006



Using temperature and micronekton it was possible to model the feeding habitat and movements of turtles opening the way to possible forecast of maps indicating the risk of turtle bycatch to assist fishermen in their activity.

Abecassis M, Senina I, Lehodey P, Gaspar P, Parker D, et al. (2013) A Model of Loggerhead Sea Turtle (*Caretta caretta*) Habitat and Movement in the Oceanic North Pacific. PLoS ONE 8(9): e73274.

Predicted habitat and observed tracks

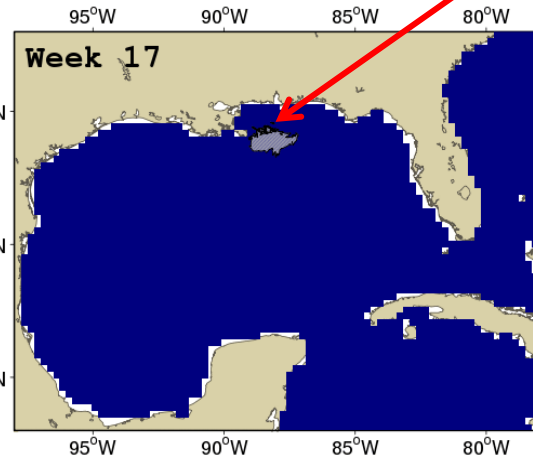
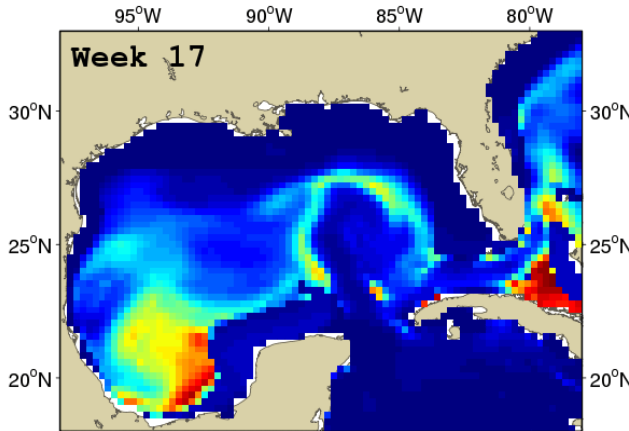
lbt\_Hadym  
2005 May 02

## Ex.: Spawning habitat and larvae dynamics: Atlantic Bluefin tuna

- ❑ Favorable **temperature** for optimal growth
- ❑ Coincidence of spawning with presence of food (**zooplankton**) for larvae (match/mismatch of Cushing, 1975)
- ❑ Coincidence of spawning with absence of predators (**micronekton**) of larvae
- ❑ Retention by **currents** of larvae in favorable areas (i.e., with conditions above)
- ❑ Accessibility to spawners (oxygen, clear waters, deep cold waters)
- ❑ Food for spawners (micronekton)

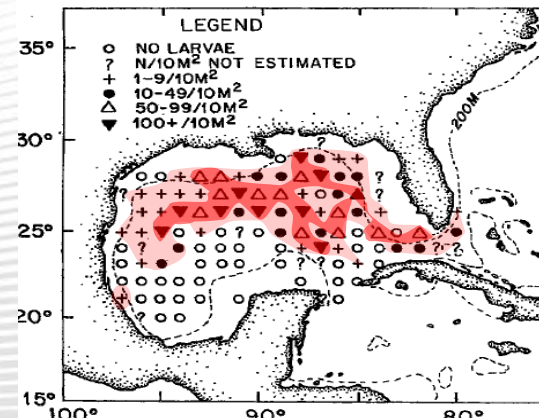
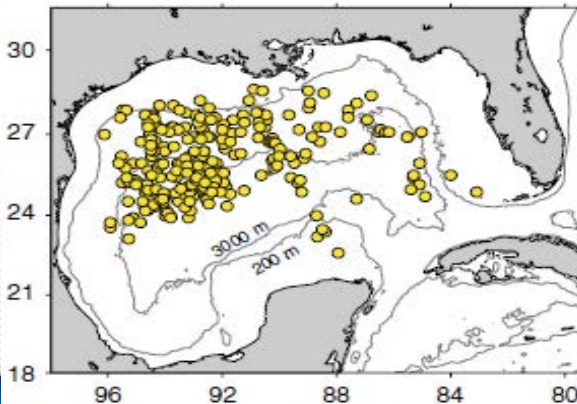
Oil spill from Deep Horizon platform accident

Predicted Bluefin spawning habitat



Predicted larvae density (âge 1-30 d)

Geolocation of 28 bluefin tuna identified from their vertical behaviour as in breeding phase (Teo et al. 2007).



Density of bluefin tuna larvae sampled in the Gulf of Mexico in 1973-74, from Mather et al. (1995)

## STRENGTH:

Parcimonious population model (<20 parameters) to describe complete and spatial dynamics independently of fishing data

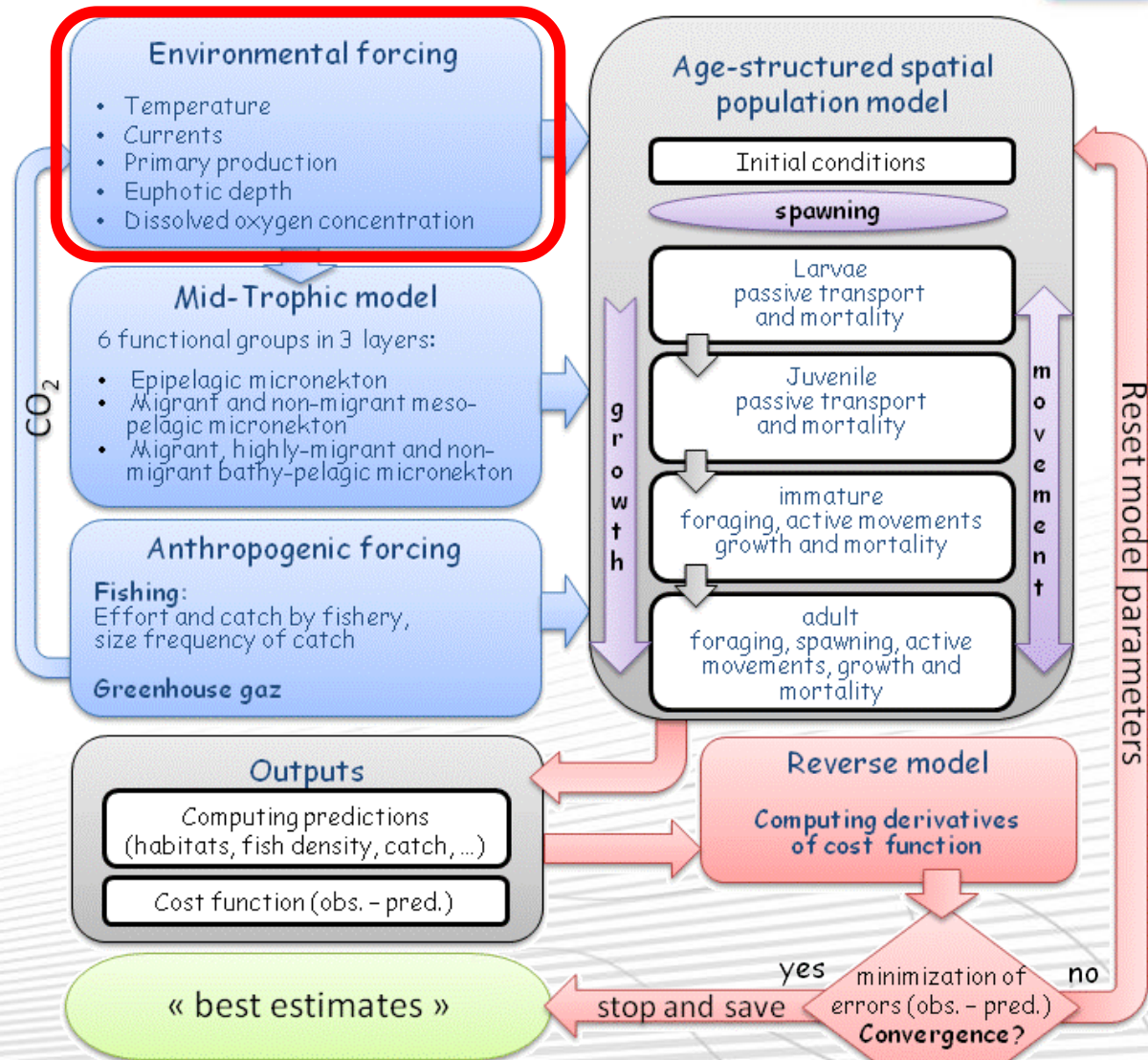
Combining (analyzing) fishing impact and environmental / climate effects.

Climate forecasts

## WEAKNESS:

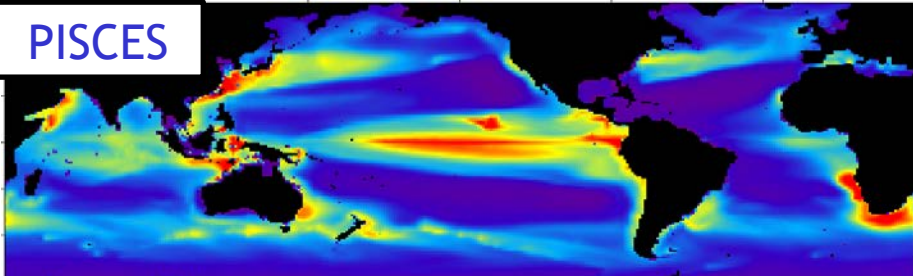
Despite obvious progress, there are still uncertainties, errors, biases in 3D simulations of ocean physics and biogeochemistry...

Problem to access long historical reliable time series

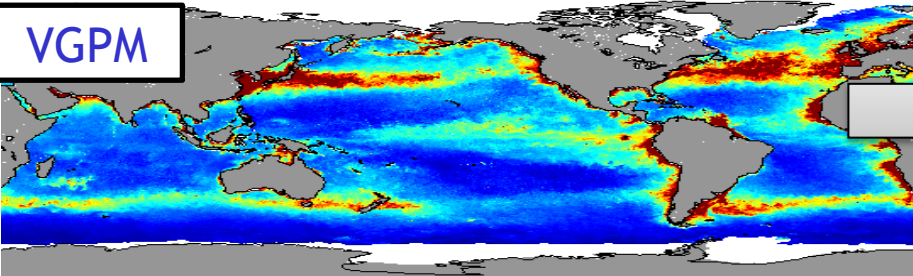


➤ Which Primary production ?

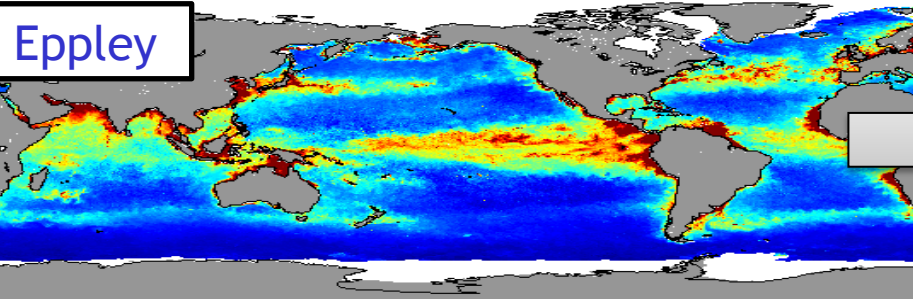
PISCES



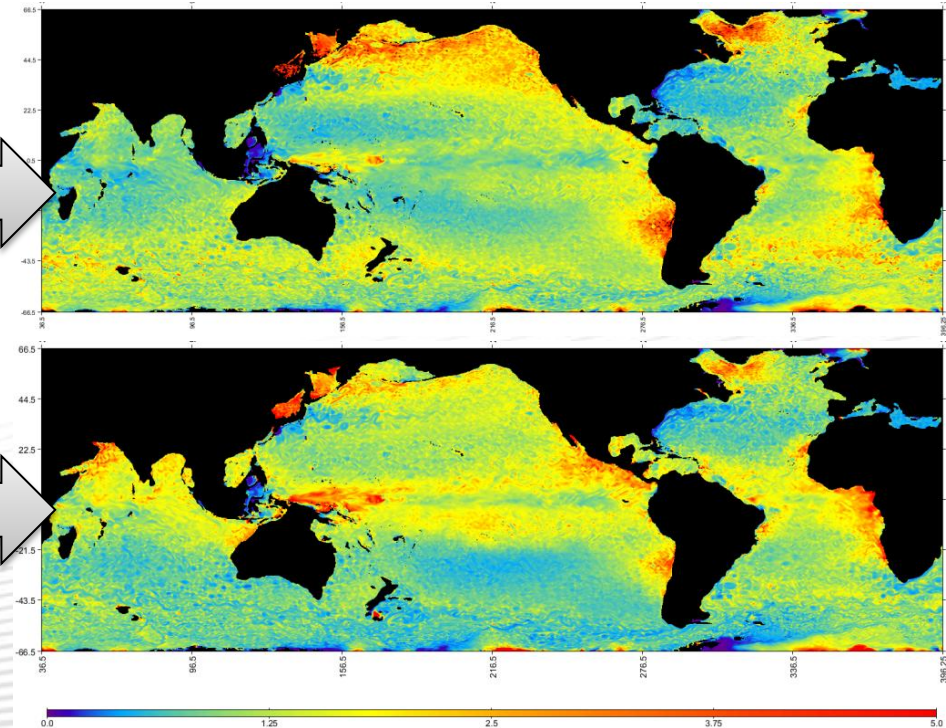
VGPM



Eppley

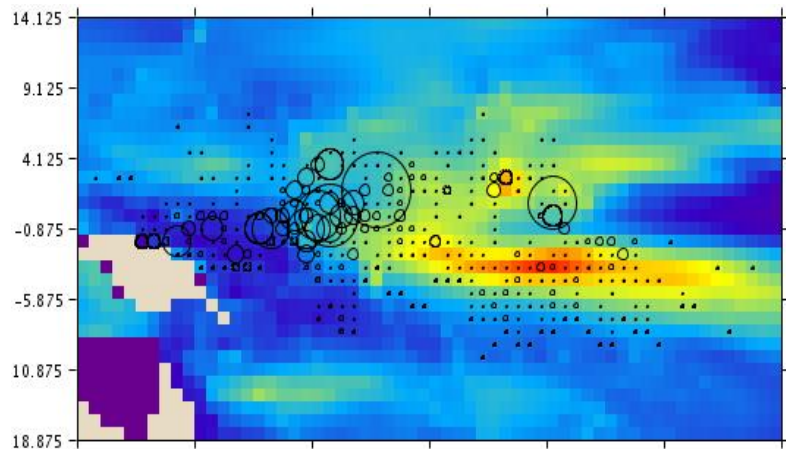


Deep non migrant micronekton

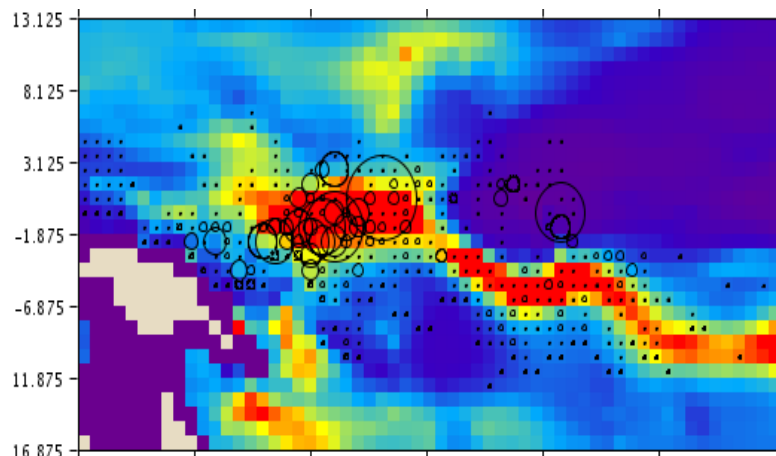


➤ Which currents?

Sep 2008

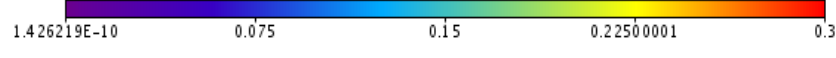
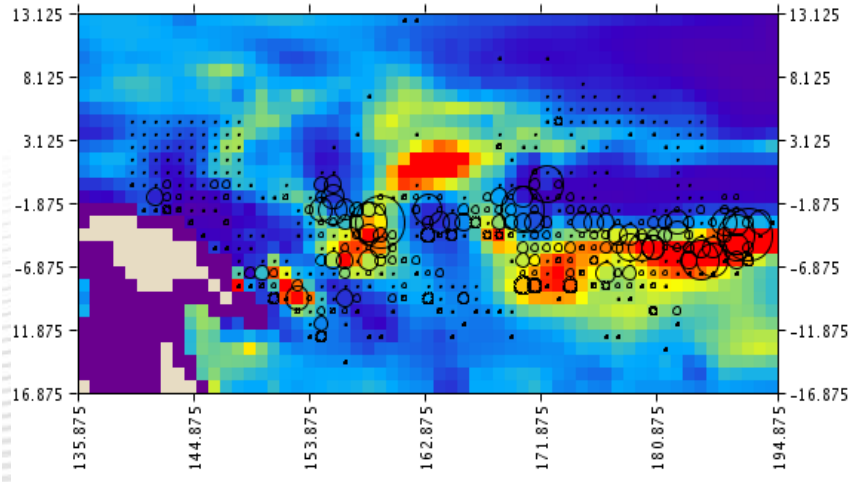
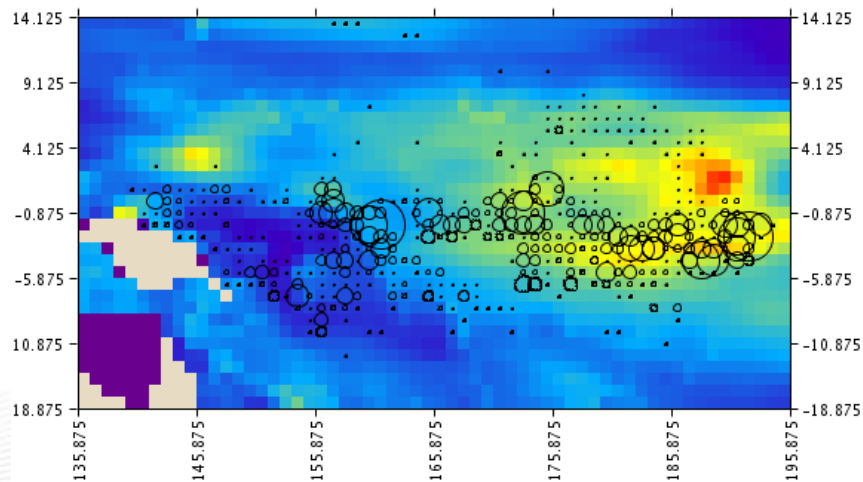


Skipjack with GLORYS



Skipjack with OMEGA

Jan 2010



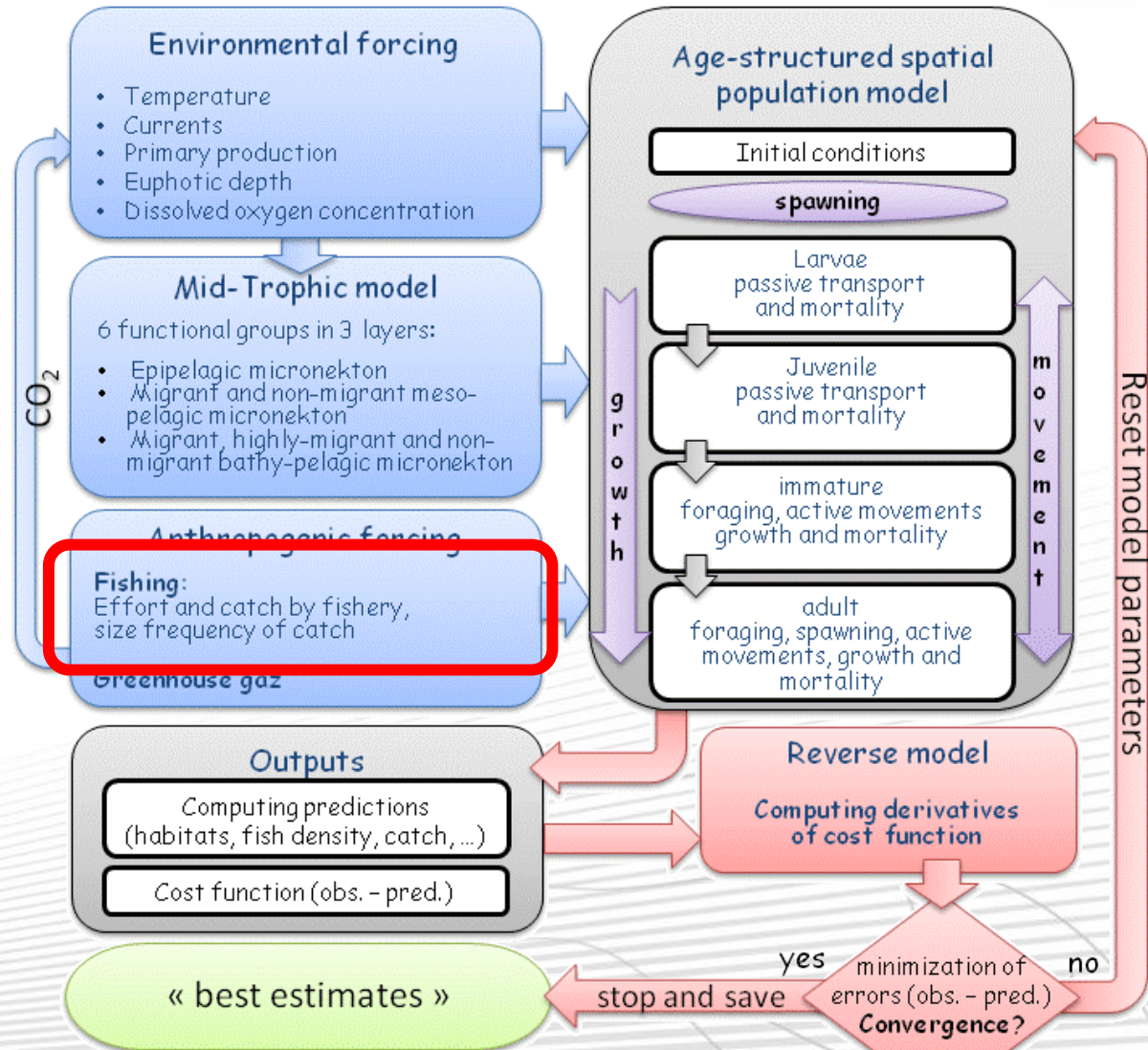
## Geo-referenced fishing data

### STRENGTH:

Take advantage of all information at its highest possible resolution in time and space.  
 Link CPUE variability directly to change in abundance and linear trends in catchability

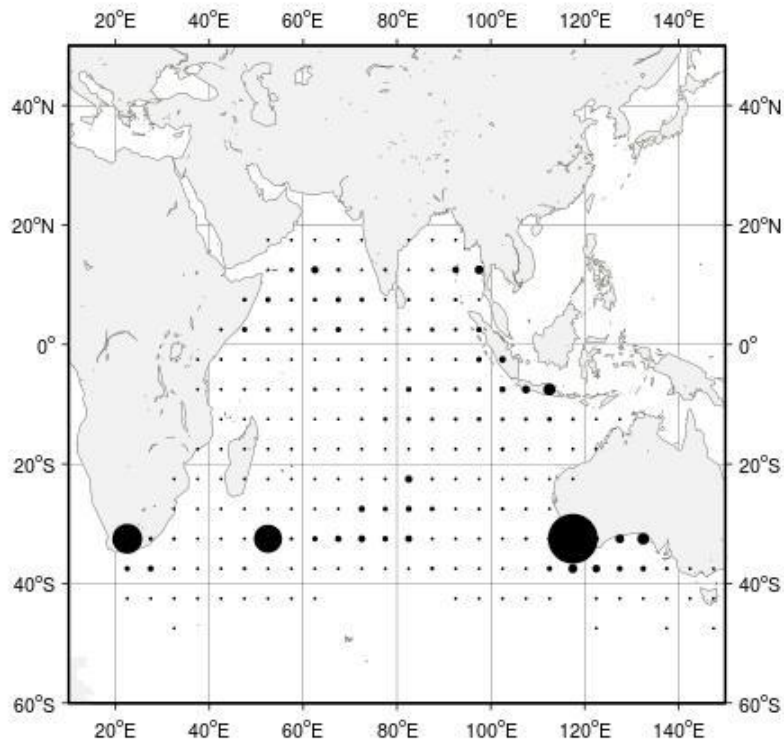
### WEAKNESS:

Historical spatially disaggregated fishing data frequently partial or not available.  
 Require careful homogeneous definition of fisheries and data screening.

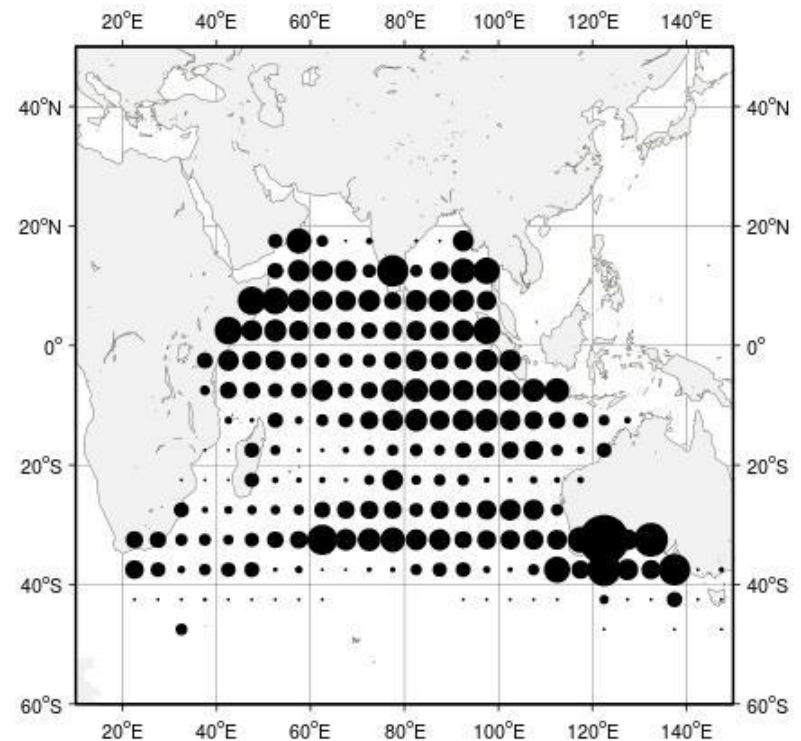


Ex: Detection of outliers following Hampel outlier Identifier method

Raw data



After outlier screening



CPUE variance (bigeye) of Japanese longline dataset

## STRENGTH:

MLE is a powerful robust approach to estimate model parameters

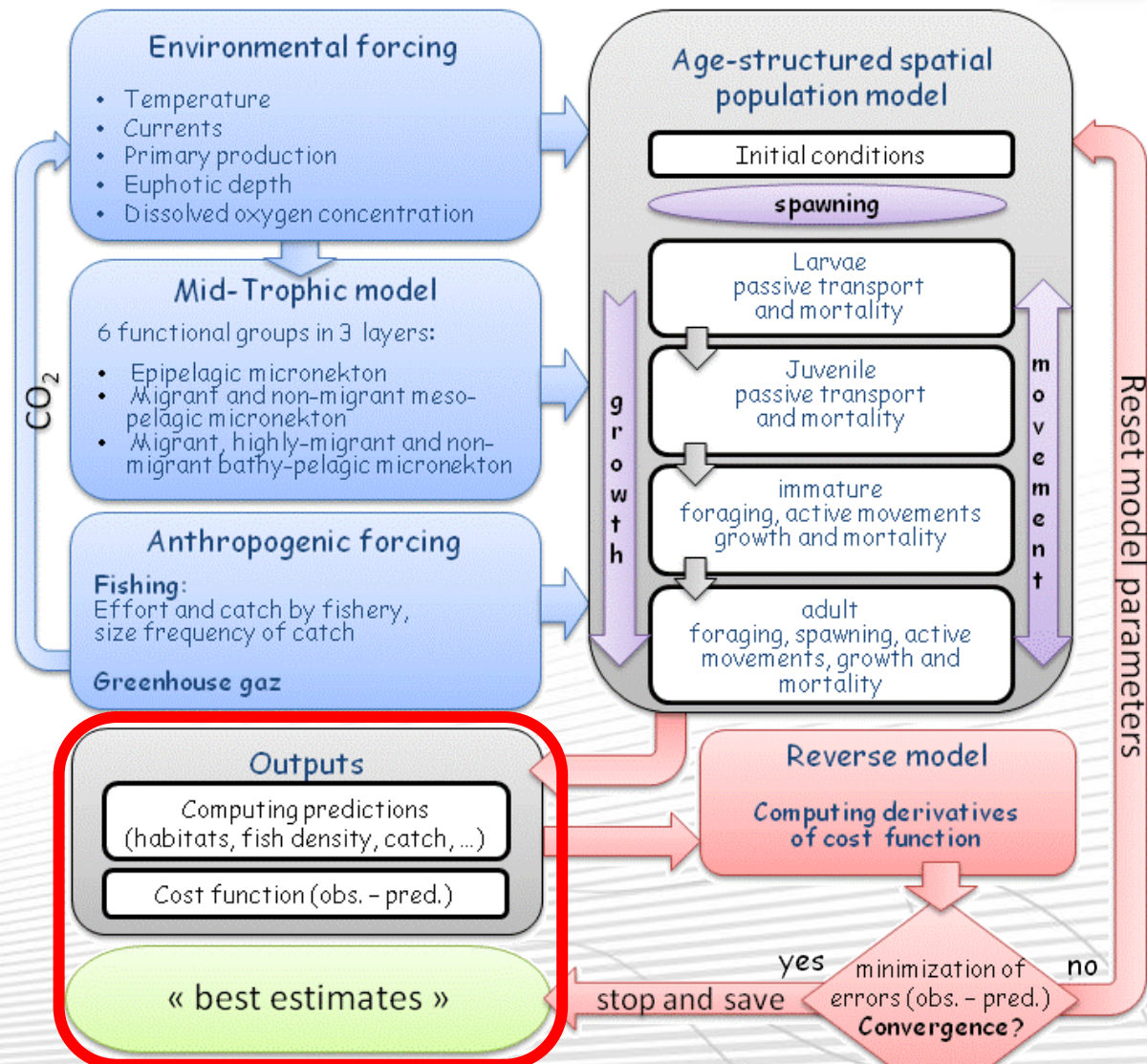
Various kind of data and « a priori » knowledge can be used to construct the cost function (CPUE LF, acoustic data, eggs and larvae densities, tagging data)

## WEAKNESS:

The global minimum of the LL is 'hidden' by a large number of local minima

Sensitivity to error probability distribution selected

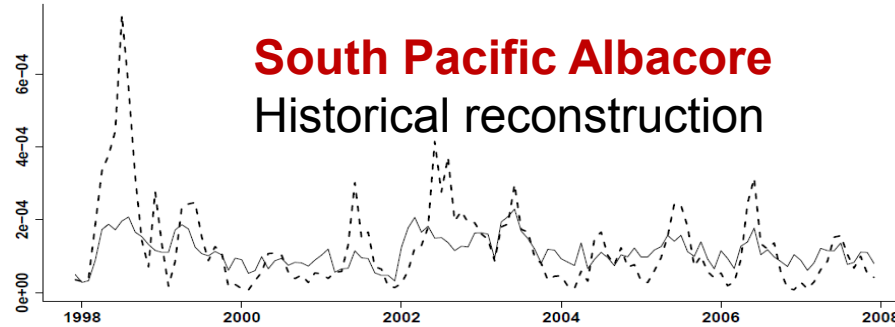
Some parameters cannot be estimated due to lack of informative data



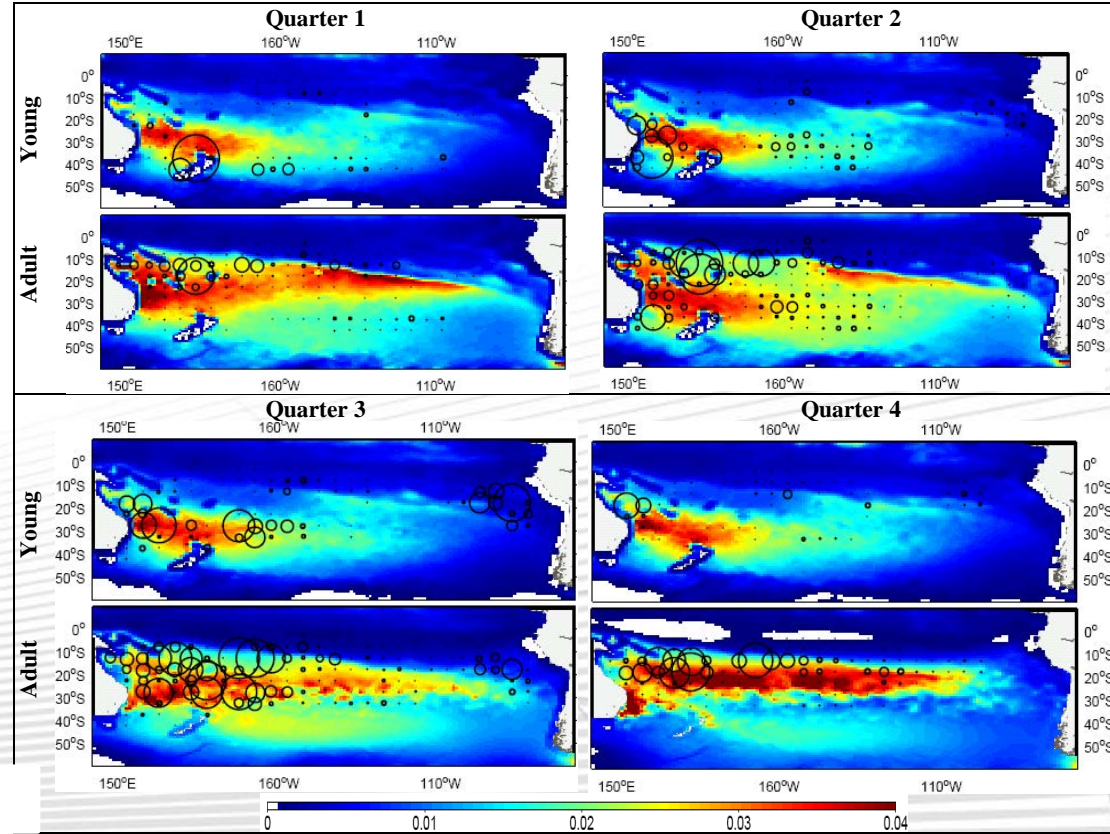
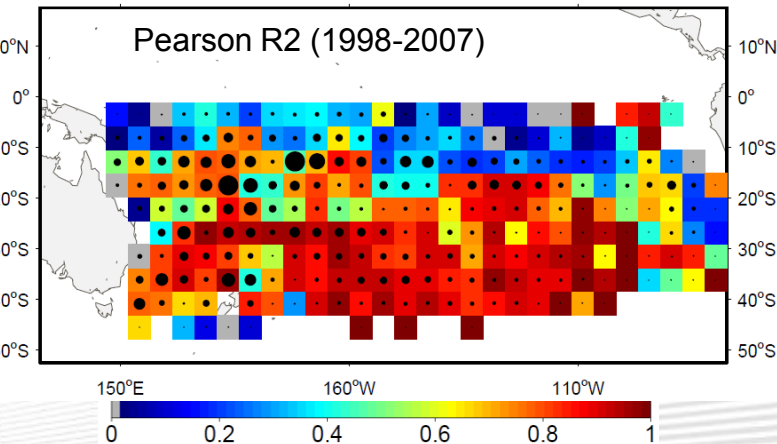
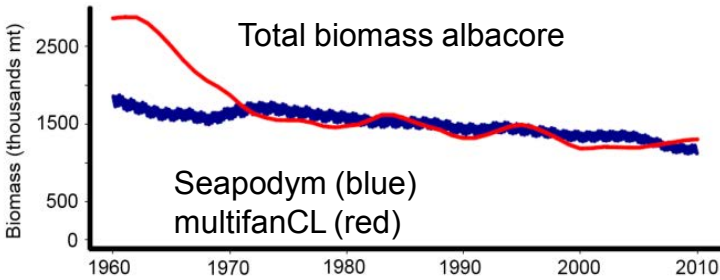
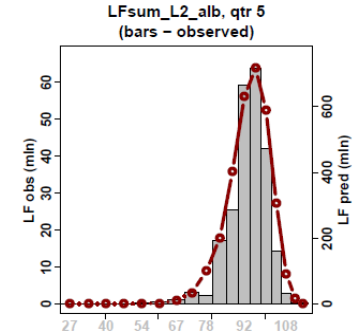


Applications of the model at basin scale at coarse resolution 2°x month, using hindcast simulation from coupled physical-biogeochemical models (OPA-PISCES)

obs CPUE\_alb\_L2 vs pred CPUE\_alb\_L2 , R1 = 0.663



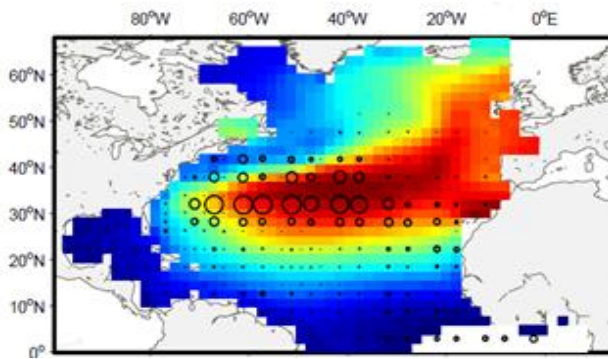
## South Pacific Albacore Historical reconstruction



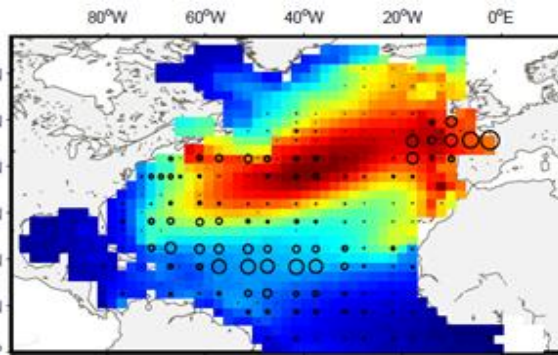
Historical hindcast forced by NCEP reanalysis

## North Atlantic Albacore

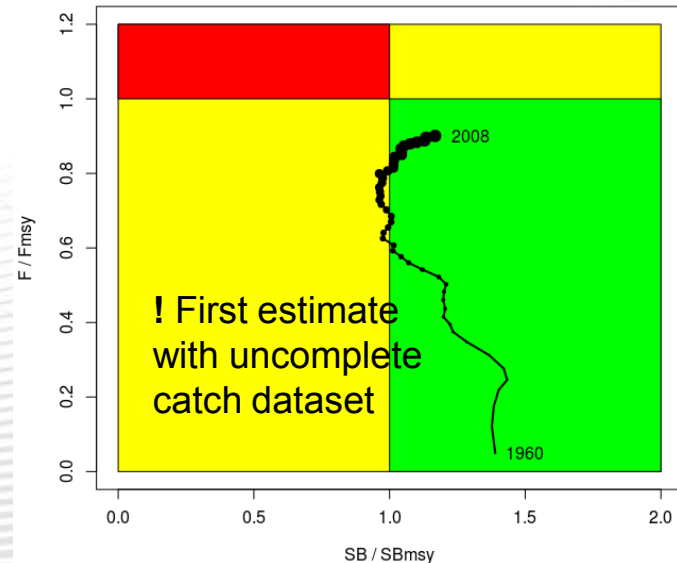
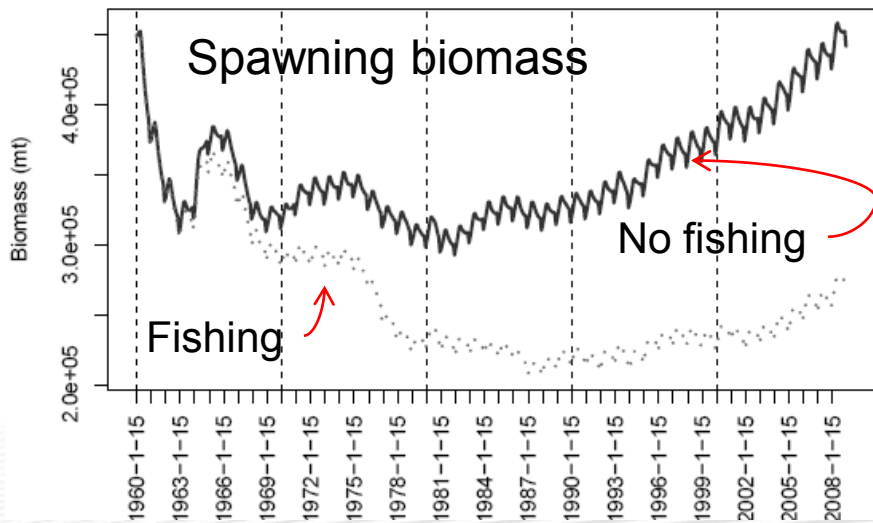
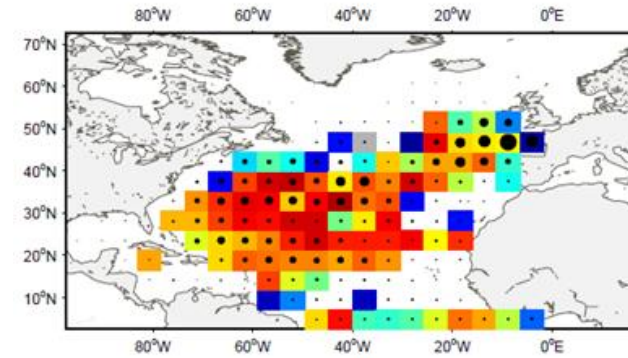
Average 1st Quarter



Average 3rd Quarter



R-squared goodness of fit (Total catch over 1960/1 - 2008/12)  
mean=0.62, nbc.pos=93



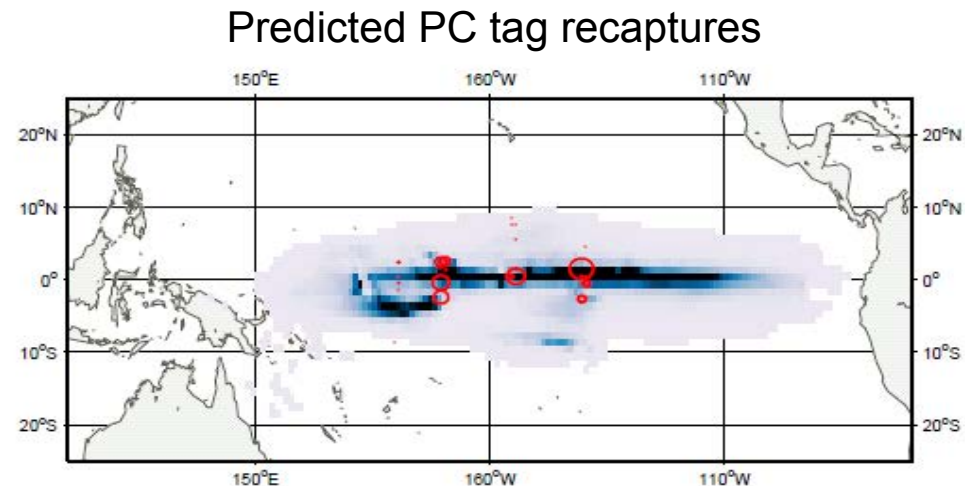
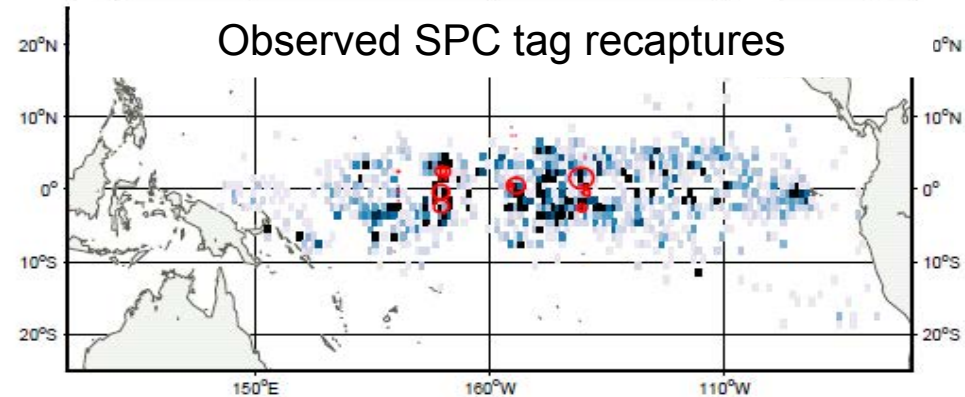
## Pacific Bigeye

A typical 'behaviour' of the model optimization experiments based on catch data only is to increase biomass and diffusion to improve the fit to catch data (largely proportional to observed fishing effort)

1/ we can include an a priori total biomass estimate in the cost function

2/ Starting to use conventional and archival tagging data to constraint the movements parameters (using only recaptures to be independent of fisheries)

3/ if available acoustic biomass estimates can be also added



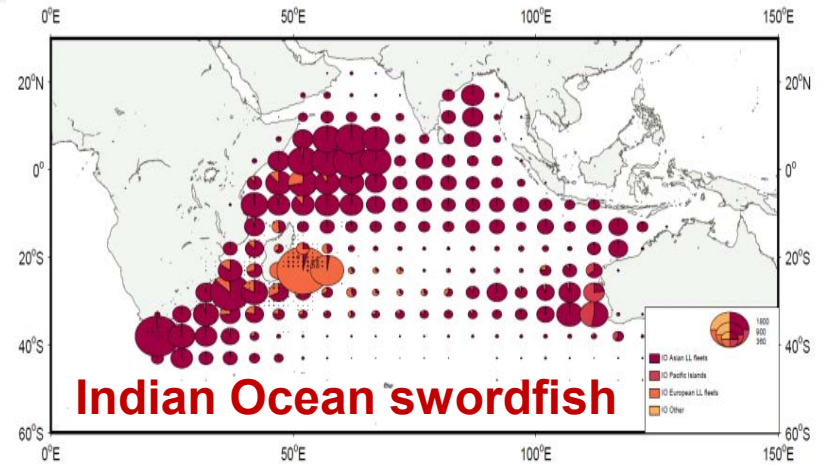
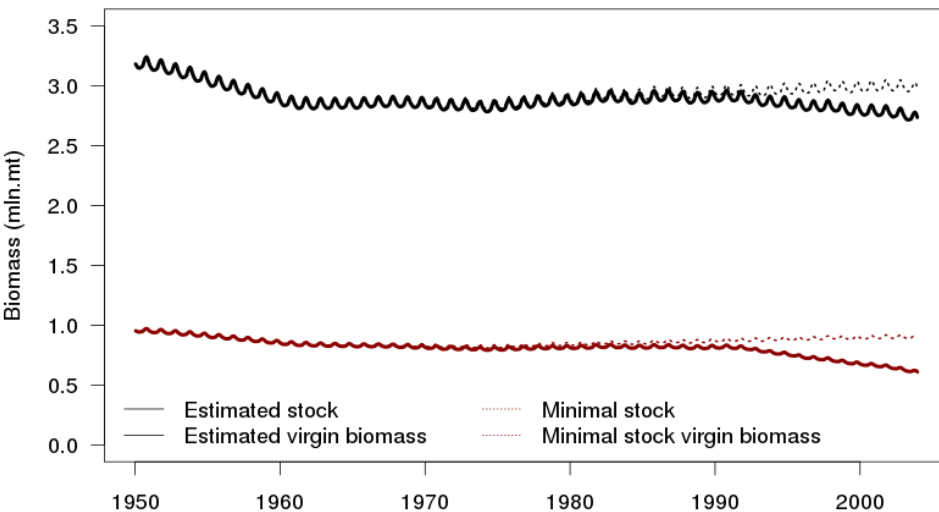
○ Red circles proportional to releases



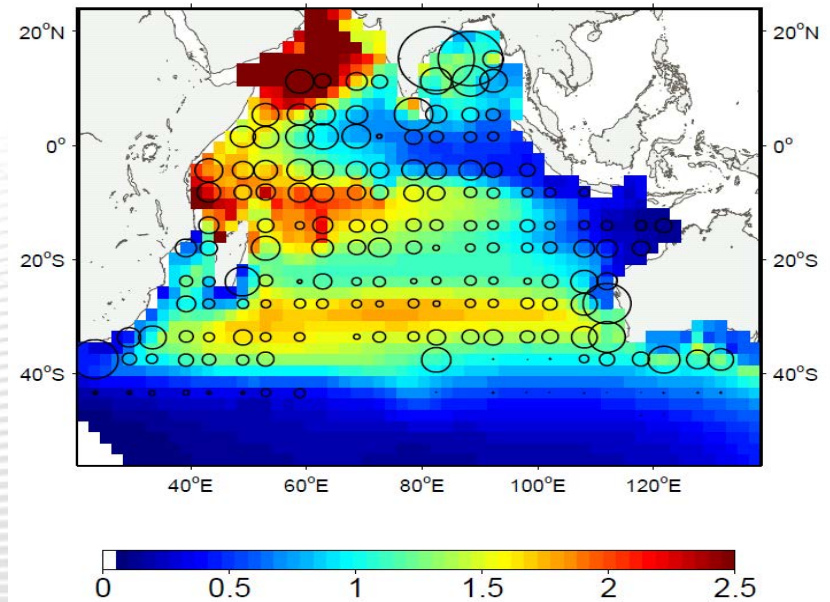
A typical 'behaviour' of the model optimization experiments based on catch data only is to increase biomass and diffusion to improve the fit to catch data (largely proportional to observed fishing effort)

4/ Computing minimum biomass estimate given the spatial and temporal distribution of catch

**Indian ocean swordfish stock estimates**

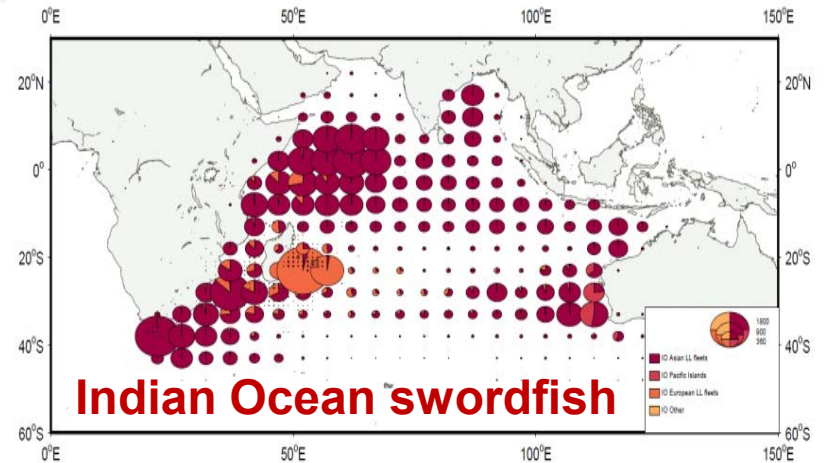


Swordfish density (adult & immature) in Nb. ind. km<sup>-2</sup> with circles proportional to average CPUE of Japanese fleet (Nb.ind/100.hooks)

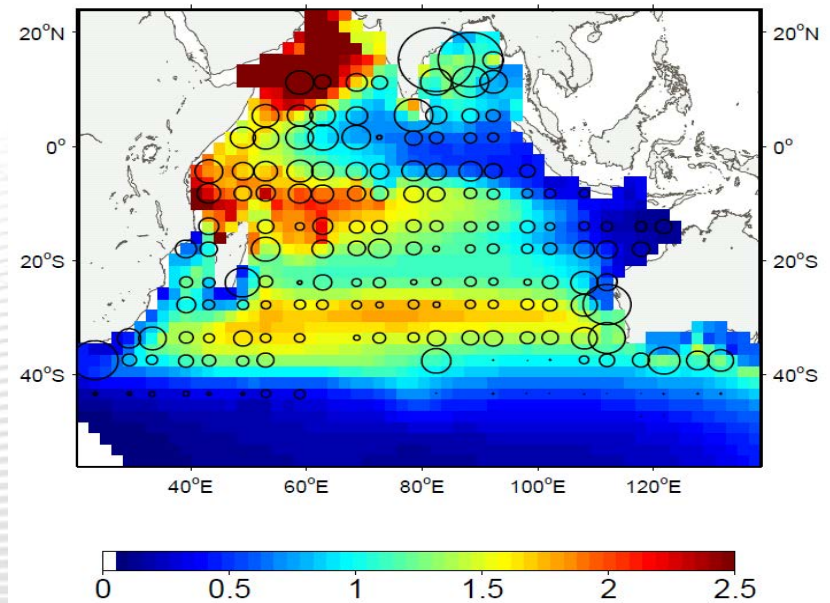
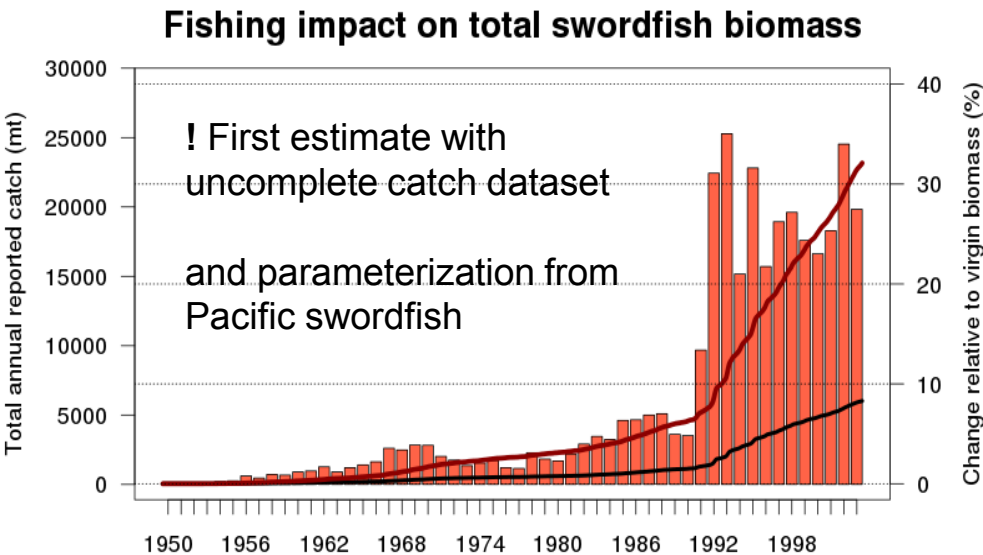


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4/ Computing minimum biomass estimate given the spatial and temporal distribution of catch



Swordfish density (adult & immature) in Nb. ind. km<sup>-2</sup> with circles proportional to average CPUE of Japanese fleet (Nb.ind/100.hooks)



Running SEAPODYM with Climate Earth Model forcing: **Pacific skipjack (IPCC A2)**

Larvae

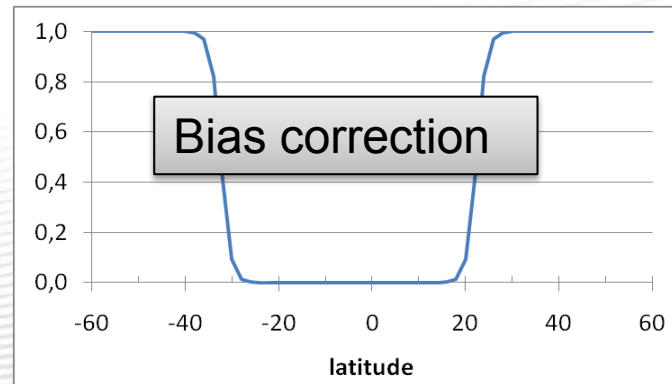
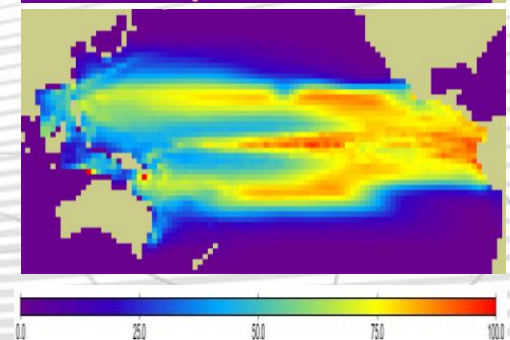
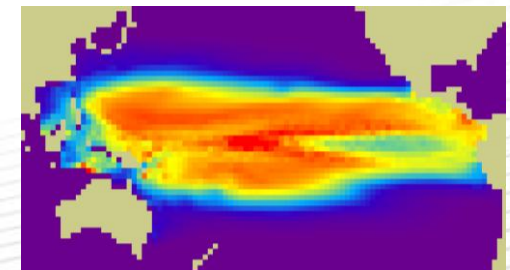
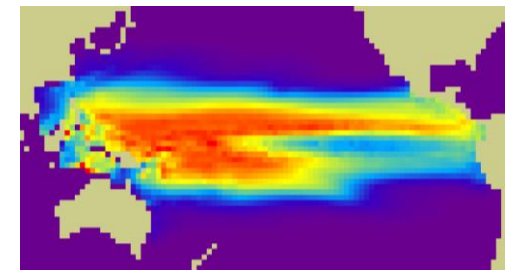
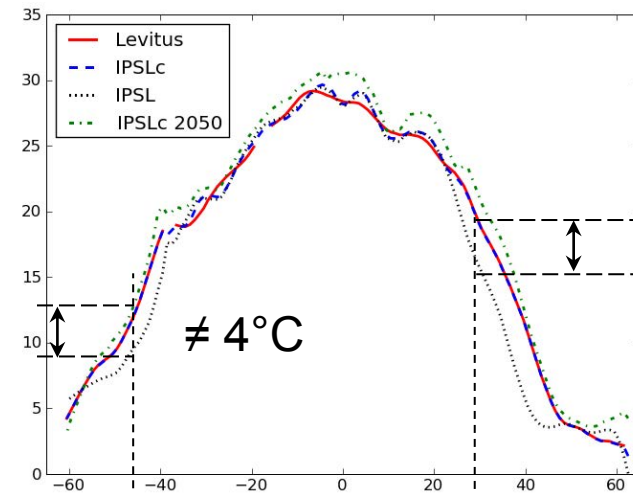
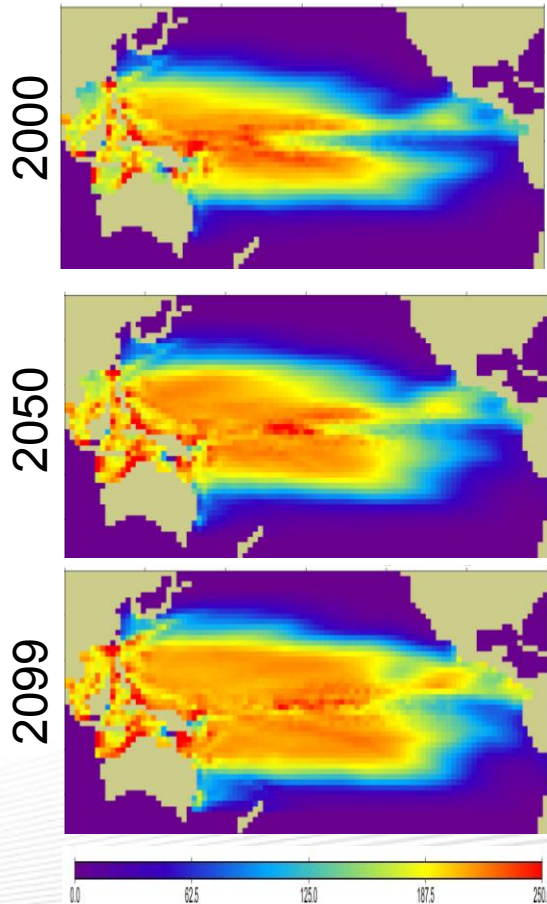
The model has a bias in temperature

Larvae

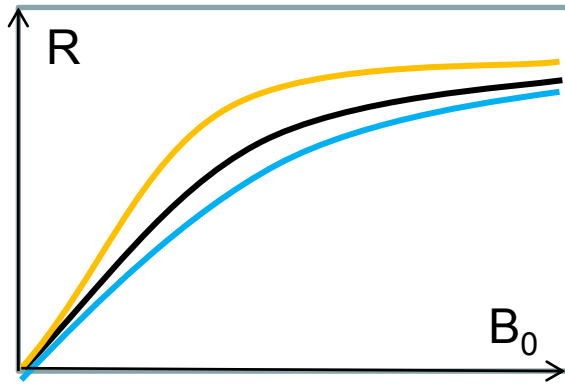
1st Exp with IPSL-CM4

Temperature transect at longitude 180°

2nd Exp after T° correction



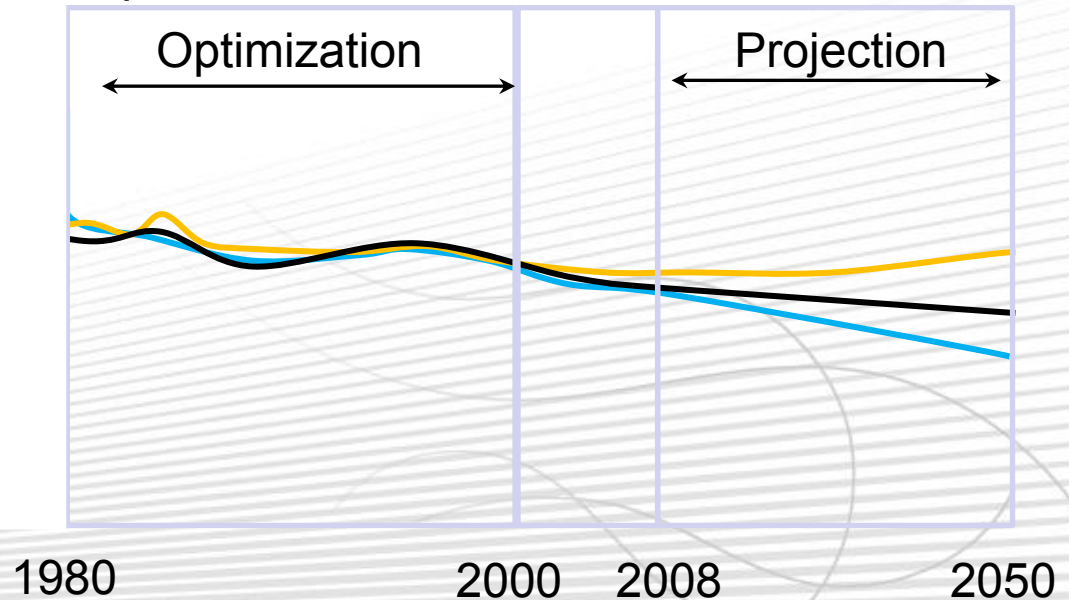
## Sensitivity to Stock-Recruitment parameters: ex. Albacore



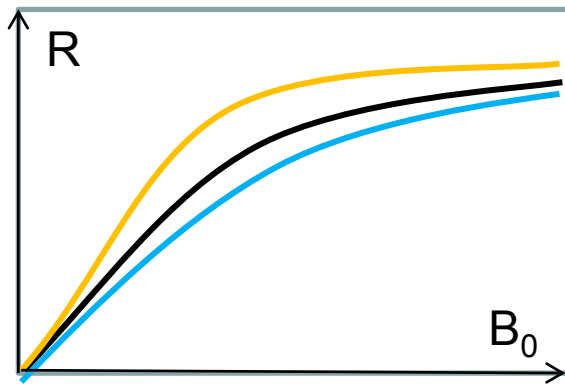
- Fish lifespan = 15 yrs
- Lack of model sensitivity to stock-recruitment relationship.
- Can lead to large error in CC projection



Check the fit of pred vs obs catch

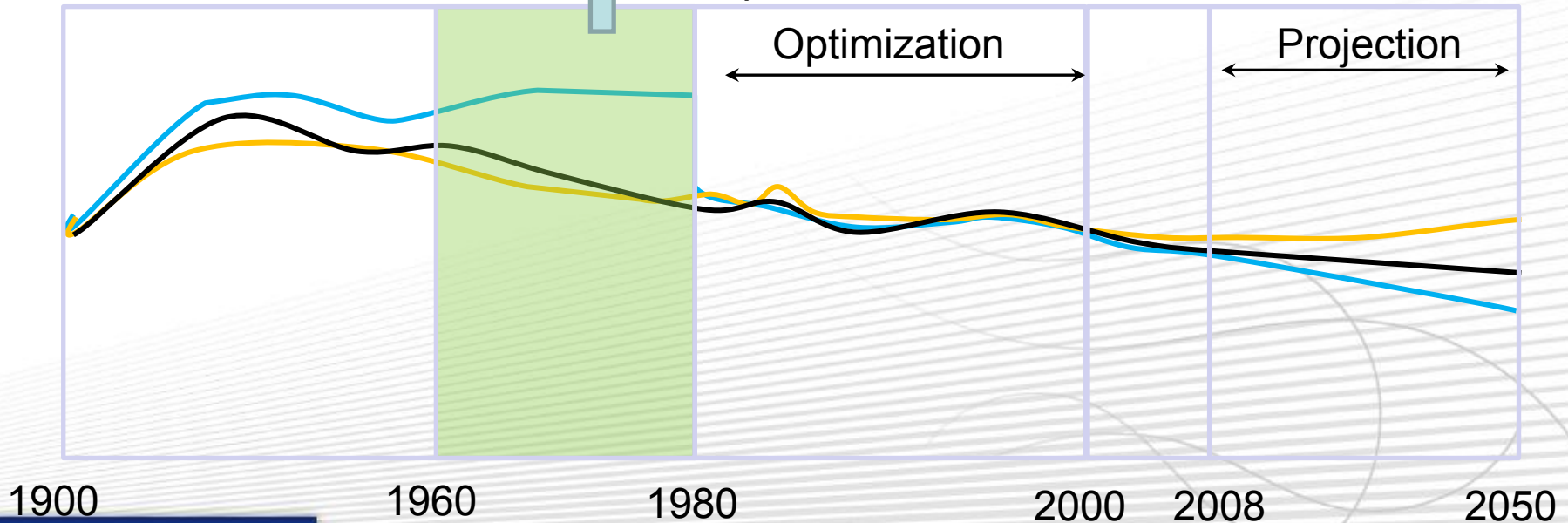


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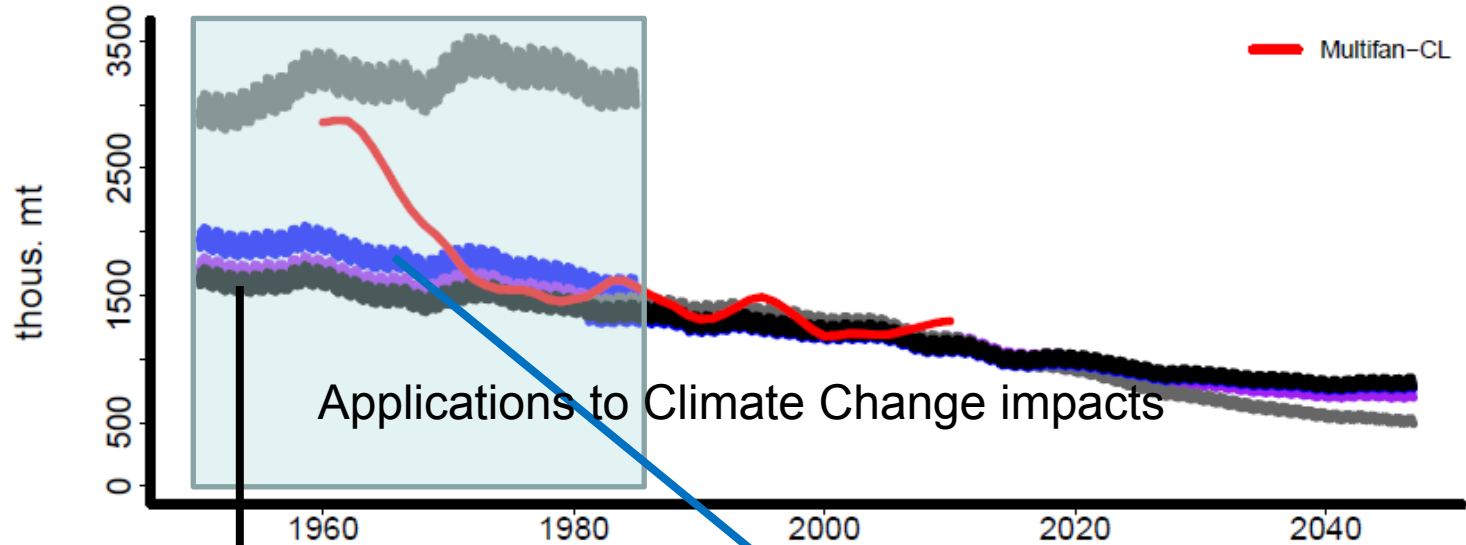
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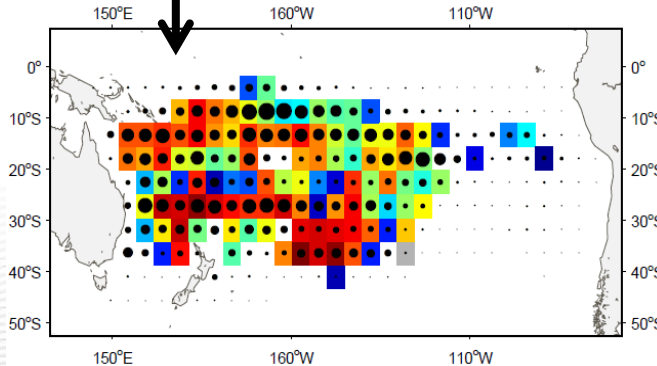


alb B tot.

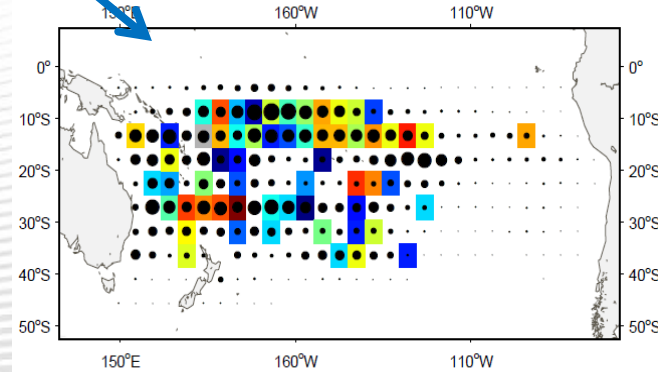


Applications to Climate Change impacts

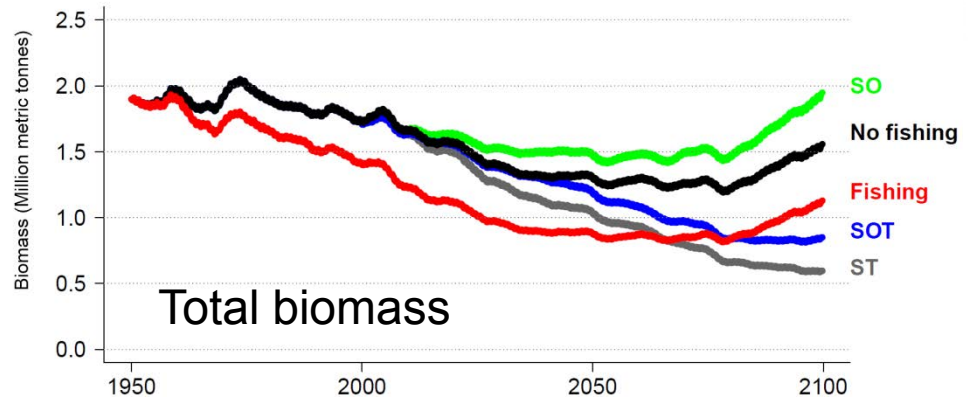
R-squared goodness of fit (Total catch over 1950/1 - 1984/12)  
mean=0.55, nbc.pos=113



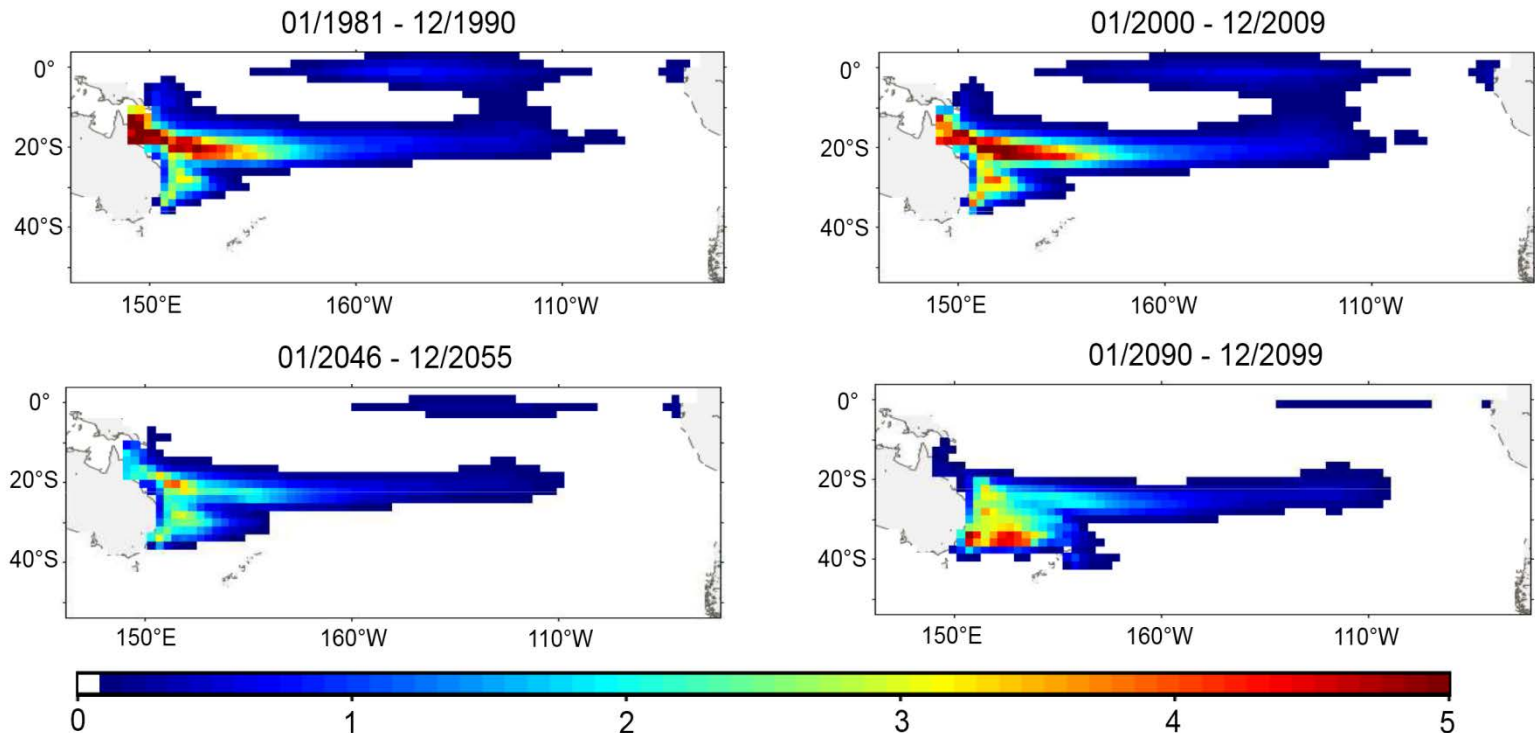
R-squared goodness of fit (Total catch over 1950/1 - 1984/12)  
mean=0.22, nbc.pos=60



Projected change in **South Pacific albacore** under IPCC A2 scenario with IPSL-CM4-PISCES forcing (temperature corrected)



Larvae density

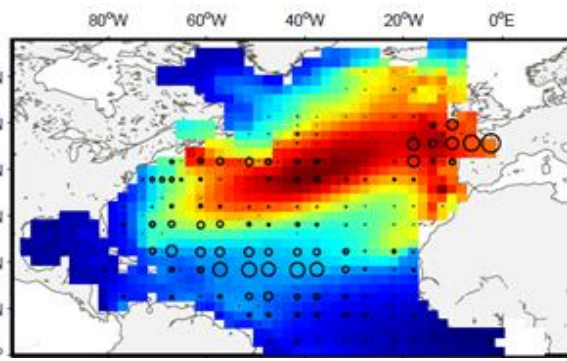
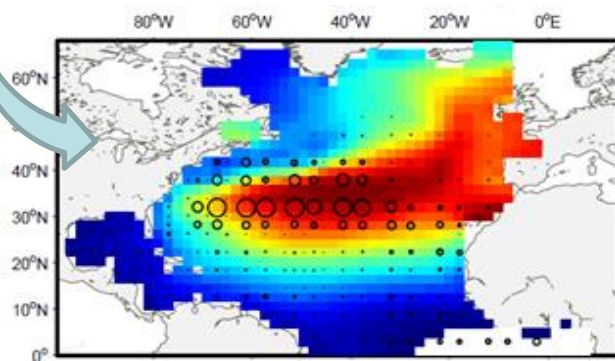


Projected change in **North Atlantic albacore** under IPCC A2 scenario with IPSL-CM4-PISCES forcing (temperature corrected)

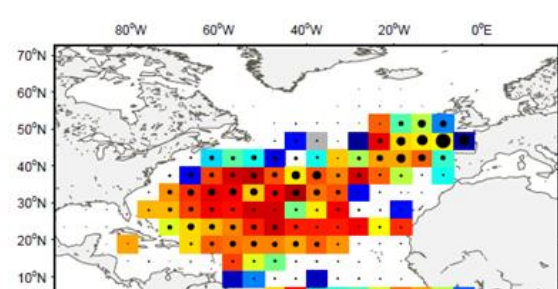
Historical hindcast forced by NCEP-OPA-PISCES reanalysis

Average 1st Quarter

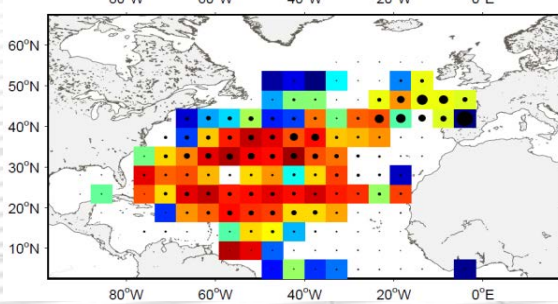
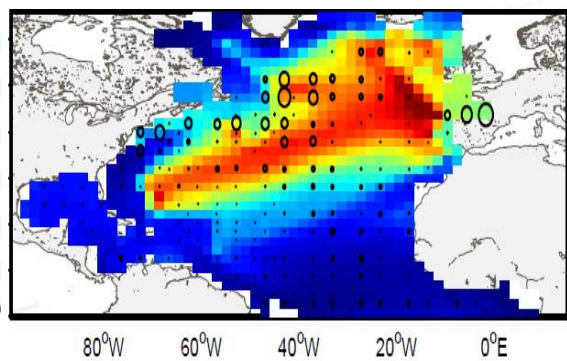
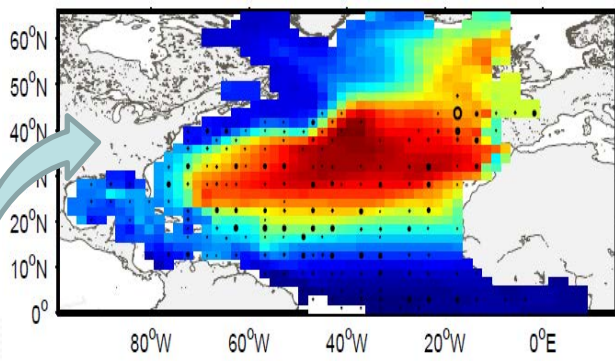
Average 3rd Quarter



R-squared goodness of fit (Total catch over 1960/1 - 2008/12)  
mean=0.62, nbc.pos=93



R-squared goodness of fit (Total catch over 1971/1 - 2000/12)  
mean=0.55, nbc.pos=92



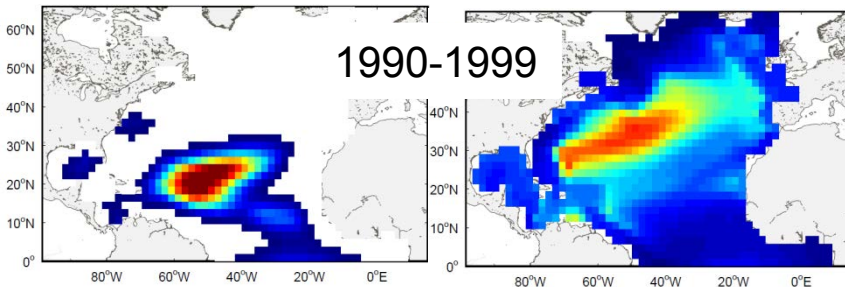
IPSL-CM4- PISCES – A2 IPCC

Projected change in **North Atlantic albacore** under IPCC A2 scenario with IPSL-CM4-PISCES forcing (temperature corrected)

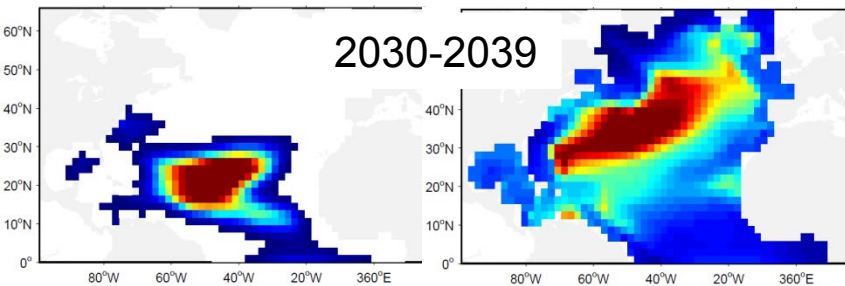
Larvae

Immature

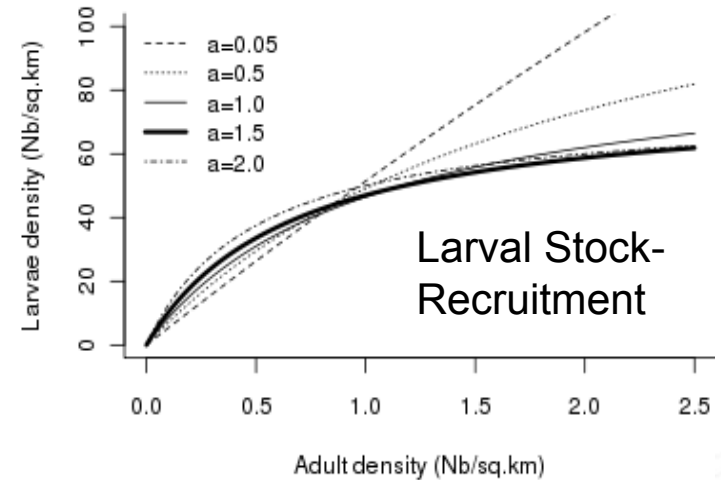
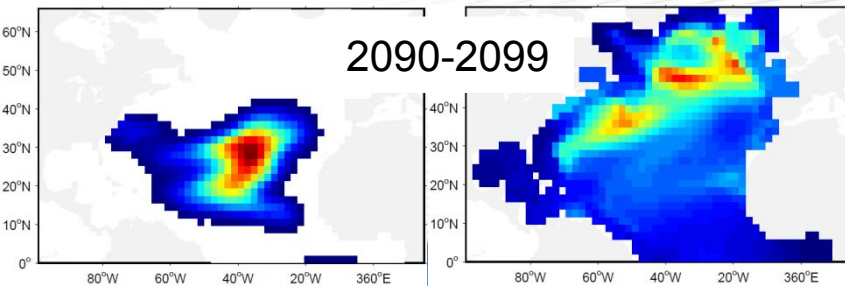
1990-1999



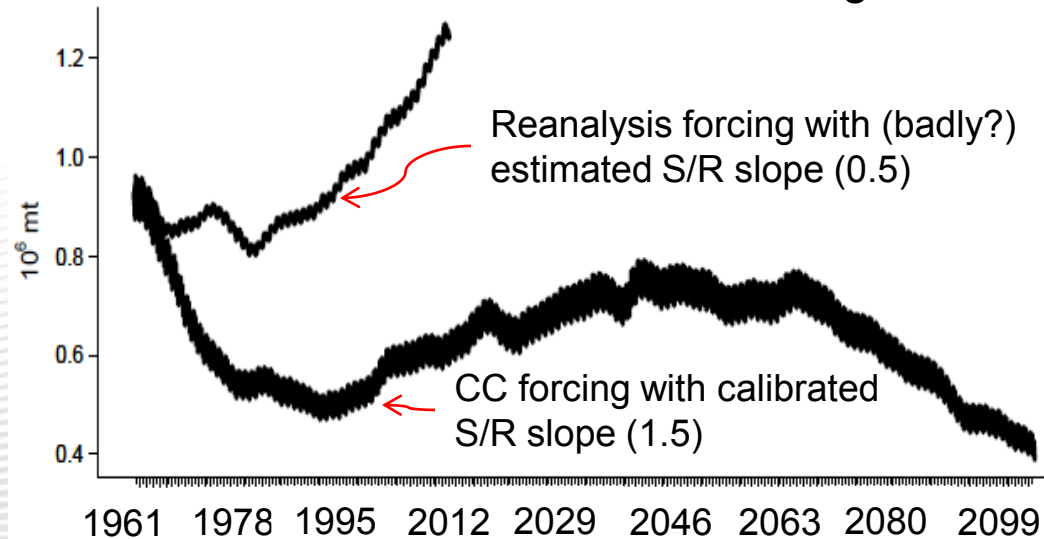
2030-2039



2090-2099



Total biomass without fishing





## Climate change simulations

- Long term simulations are interesting to estimate and test the impact of (larval) S/R relationship. (Will we be able to replace this relationship by a mechanistic detailed modeling approach one day?)
- There is no perfect simulated environmental forcing (especially from Earth Climate Models). This makes the parameter optimization approach necessary for each forcing... This is time consuming!
- To develop ensemble simulation forecast of CC, the idea is to use a single reference model for parameterization over the historical period, then to project the change with forcings from different Earth Climate models but using only their signal anomalies that would be extracted and added to the mean state of the reference model
- Other impacts to be investigated ... pH, competition?
- Fishing is still the main driver!

## Further model developments:

Given the (very) large uncertainty on micronekton biomass, the links to micronekton are developed using relative mechanisms, e.g.:

- gradient for movements
- penalty over mean values of mortality relatively to food availability and intra and multi species competition

But the model does not include absolute parameterization of mechanisms accounting for carrying capacity ( $\Rightarrow$  weakness). In some way it makes the assumption of a large decoupling between prey production and predator consumption at basin scale.

- To answer this question (critical for climate change projection) we need absolute estimates of epi and mesopelagic micronekton biomass (ongoing work to optimize the micronekton model using acoustic data)
- We need to run multi-species optimisation experiments (implying code parallelization) to estimate the level of food competition.

## Further model developments:

Ongoing regional, high resolution, and operational (real-time) applications could bring quick answers to various mechanisms and parameterisation of the model

Thank you very much  
for the invitation to this  
workshop!

Questions?

