

A spatial Ecosystem And Populations Dynamics Model (SEAPODYM)

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Oceanic Fisheries Programme

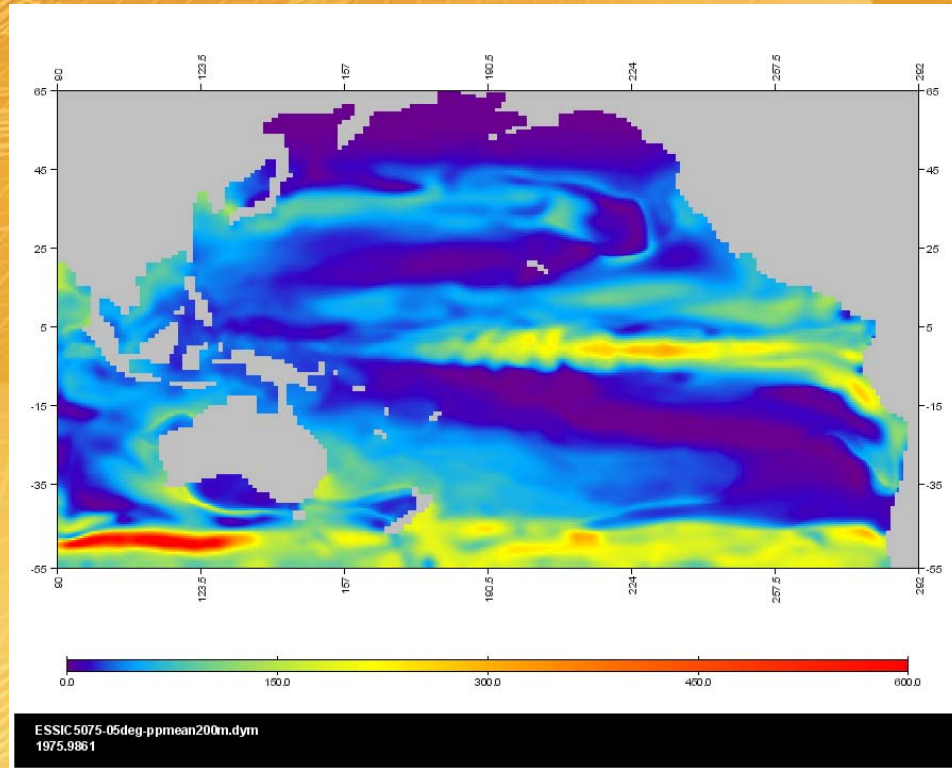
Secretariat of the Pacific Community

Noumea, New Caledonia

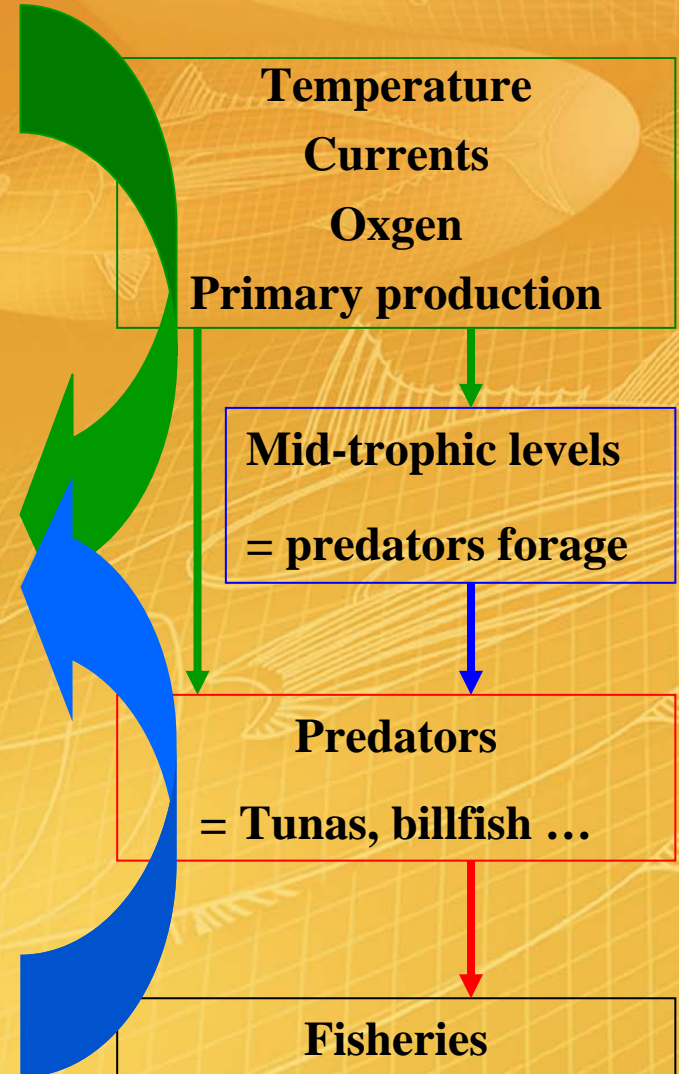


SEAPOODYM: Spatial model driven by physical and “simplified” food-web interactions

Climate/environment variability



Fishing impact



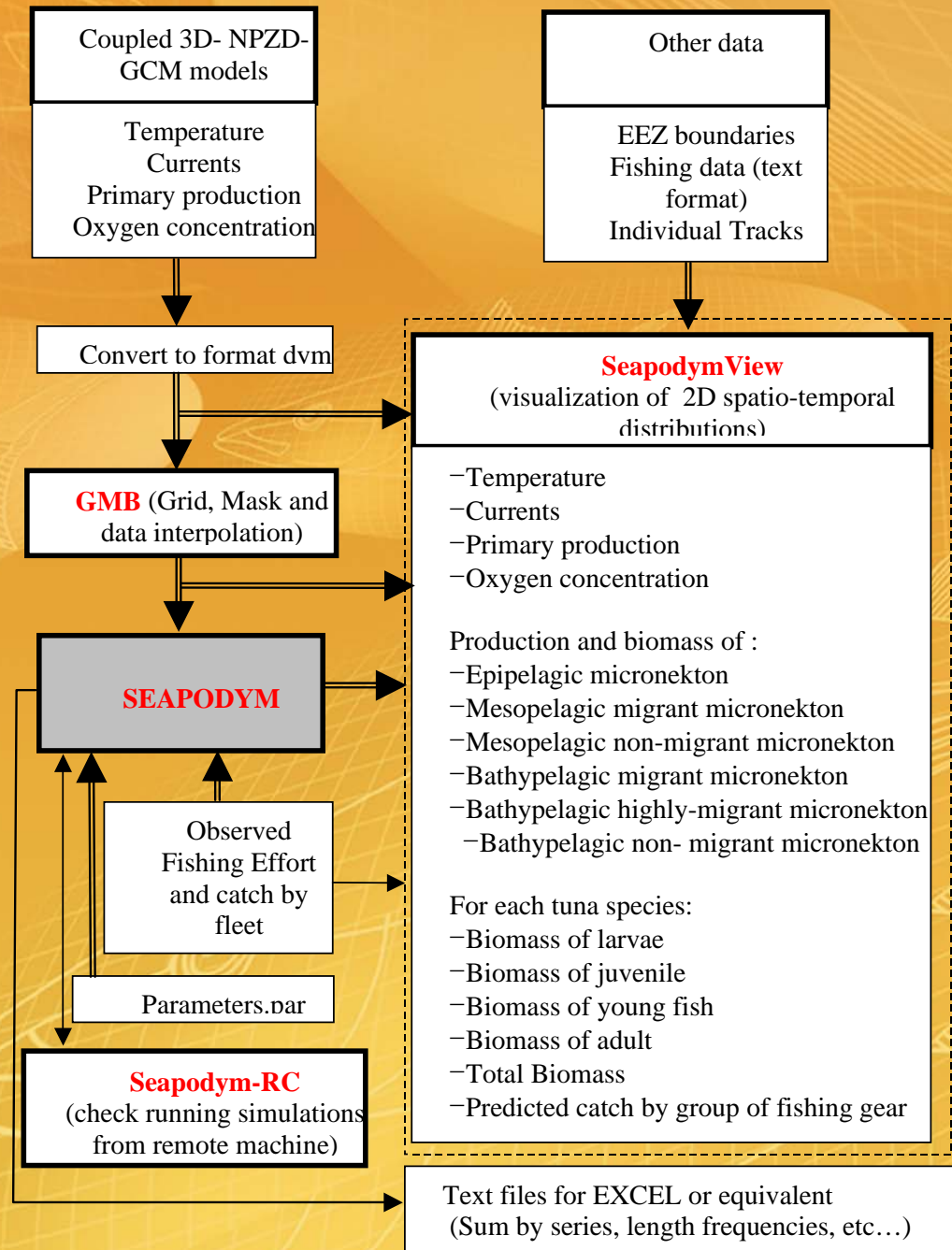
SEAPODYM software environment:

Reference manual :

-> Information Paper **ME IP-1**

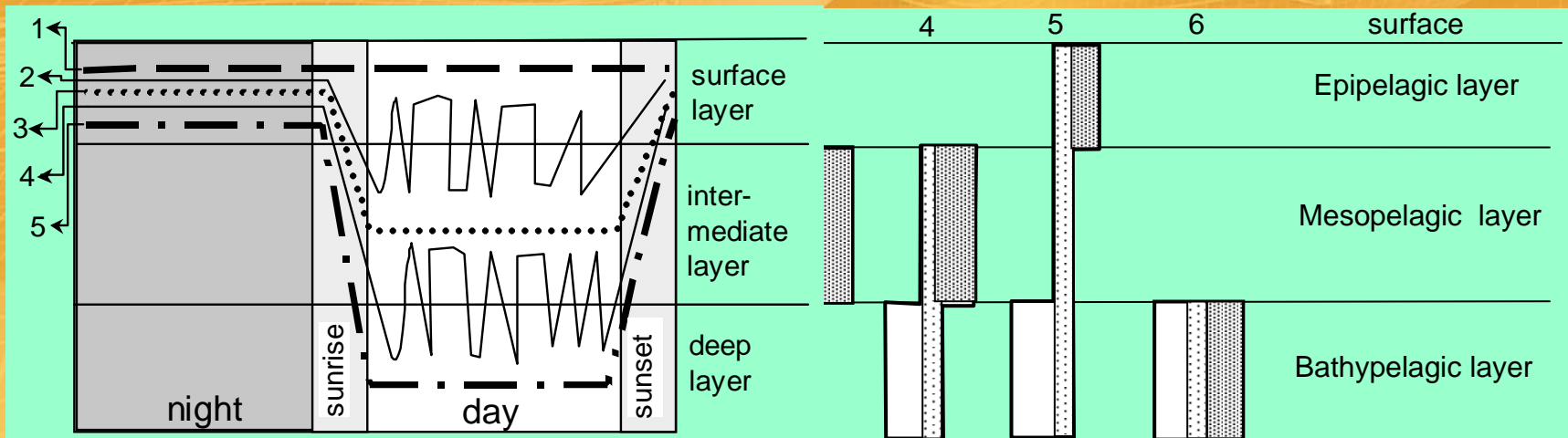
Web site :

-> www.seapodym.org



Vertical structure of the forage-predators pelagic food web

The different daily vertical distribution patterns of the micronekton in the pelagic ecosystem. 1, epipelagic; 2, migrant mesopelagic; 3, non-migrant mesopelagic; 4, migrant bathypelagic; 5, highly-migrant bathypelagic; 6, non-migrant bathypelagic.

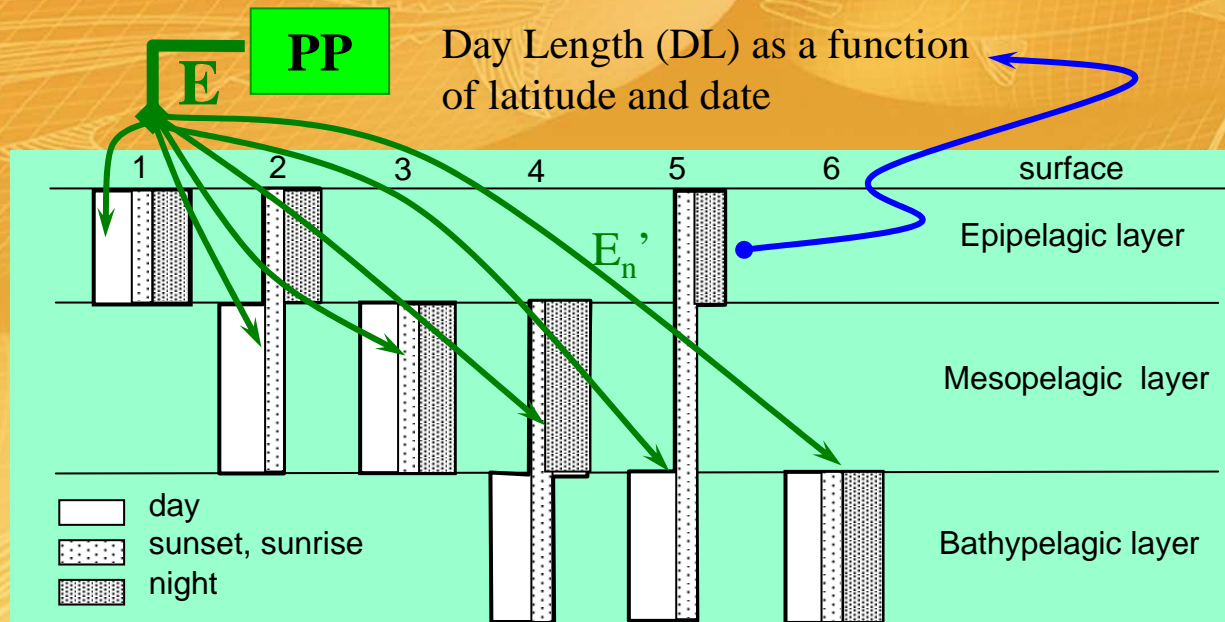


Five typical vertical movement behaviours simulated using a 3-layer and 2-type of prey pelagic system (adapted from Dagorn et al. 2000):

- 1- epipelagic predators (e.g., skipjack, marlins and sailfish);
- 2- predators moving between the surface and intermediate layers during the day (e.g., yellowfin tuna);
- 3- predators mainly in the intermediate layer during the day (e.g., albacore tuna);
- 4- predators moving between deep and intermediate layer during the day (e.g., blue shark);
- 5- predators mainly in the deep layer during the day (e.g., bigeye tuna and swordfish).

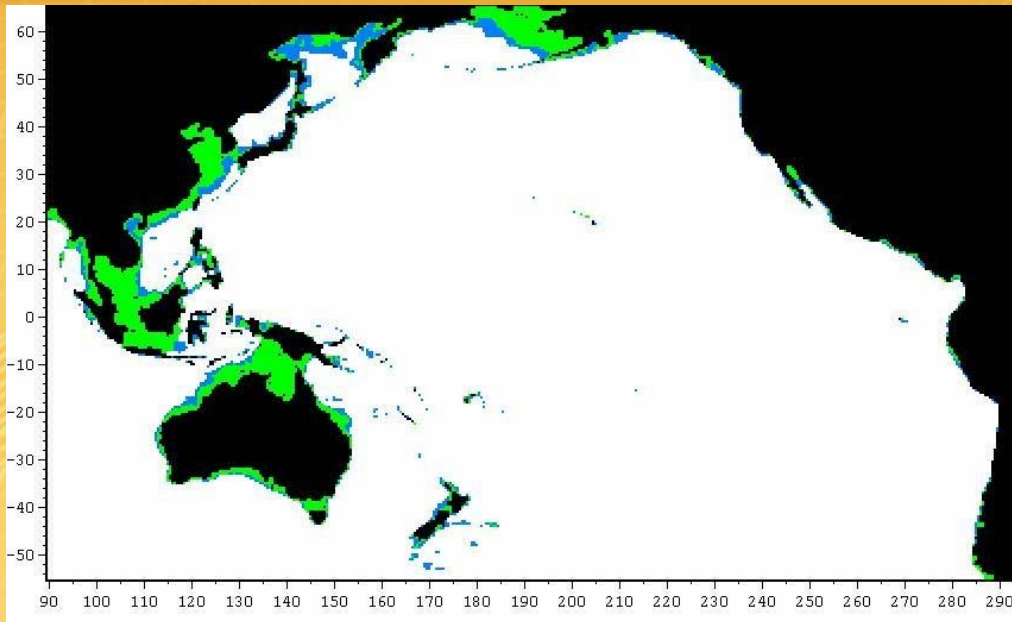
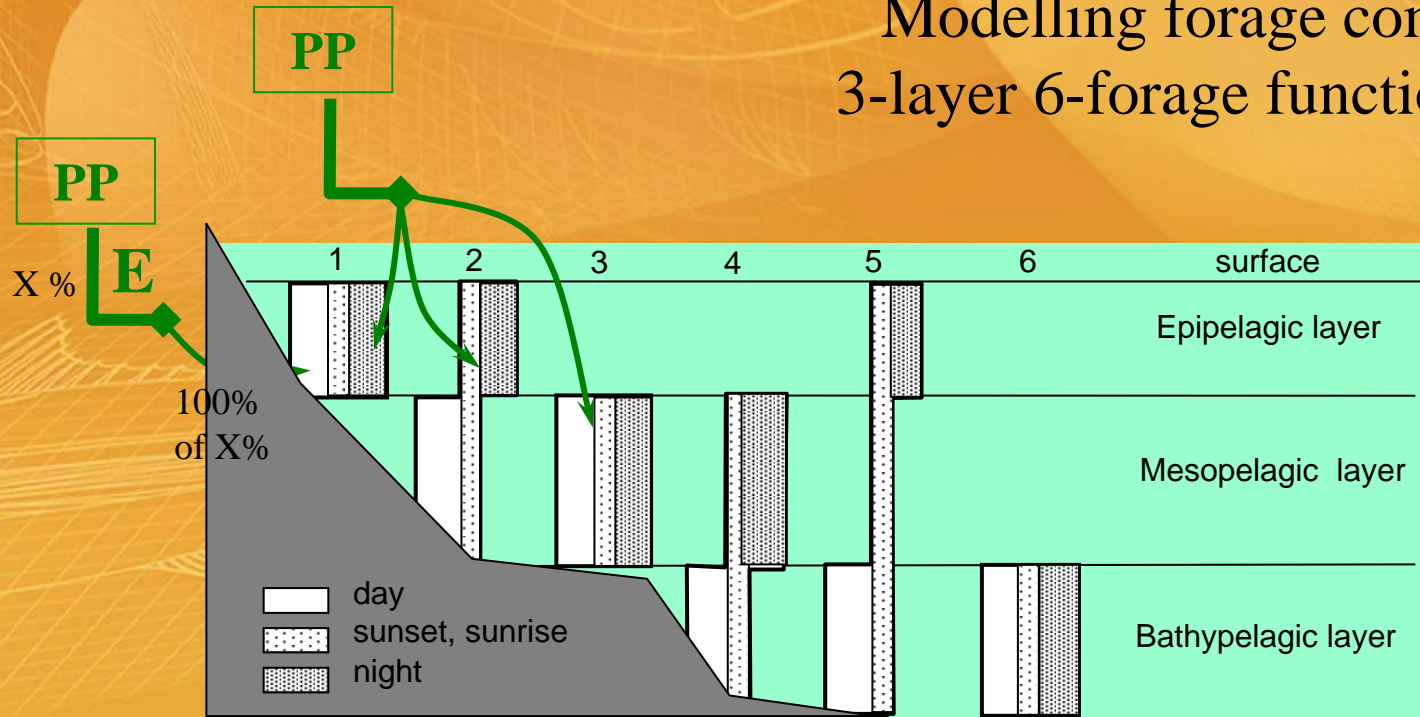
Modelling forage components:

3-layer 6-forage functional groups



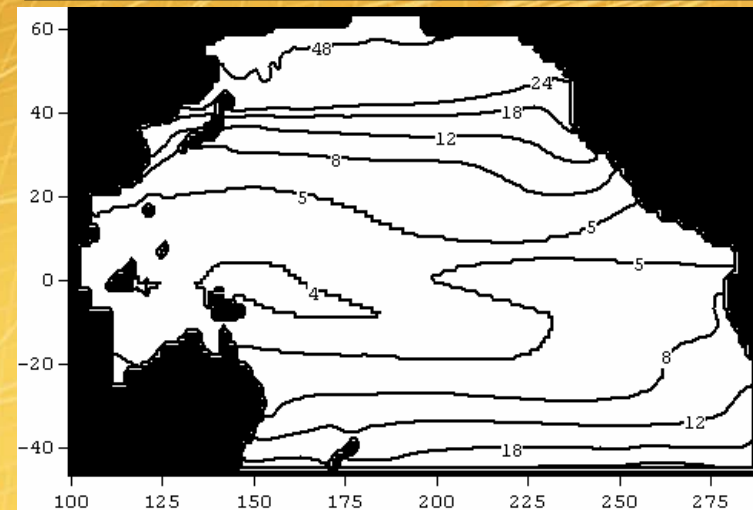
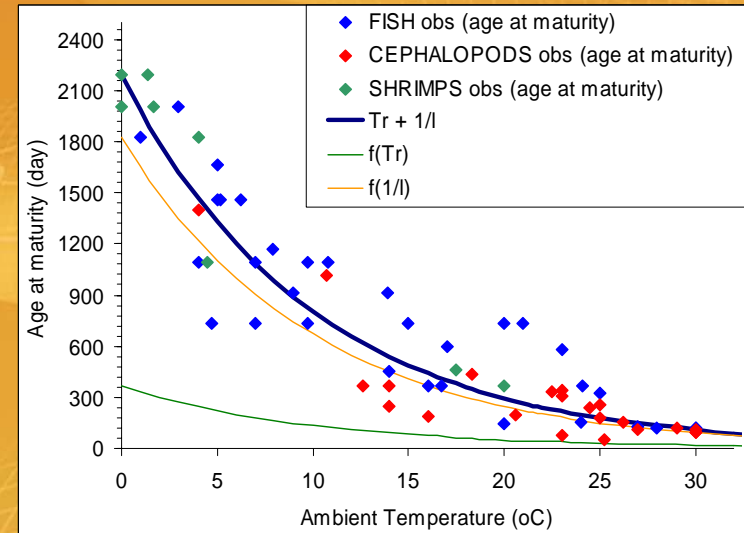
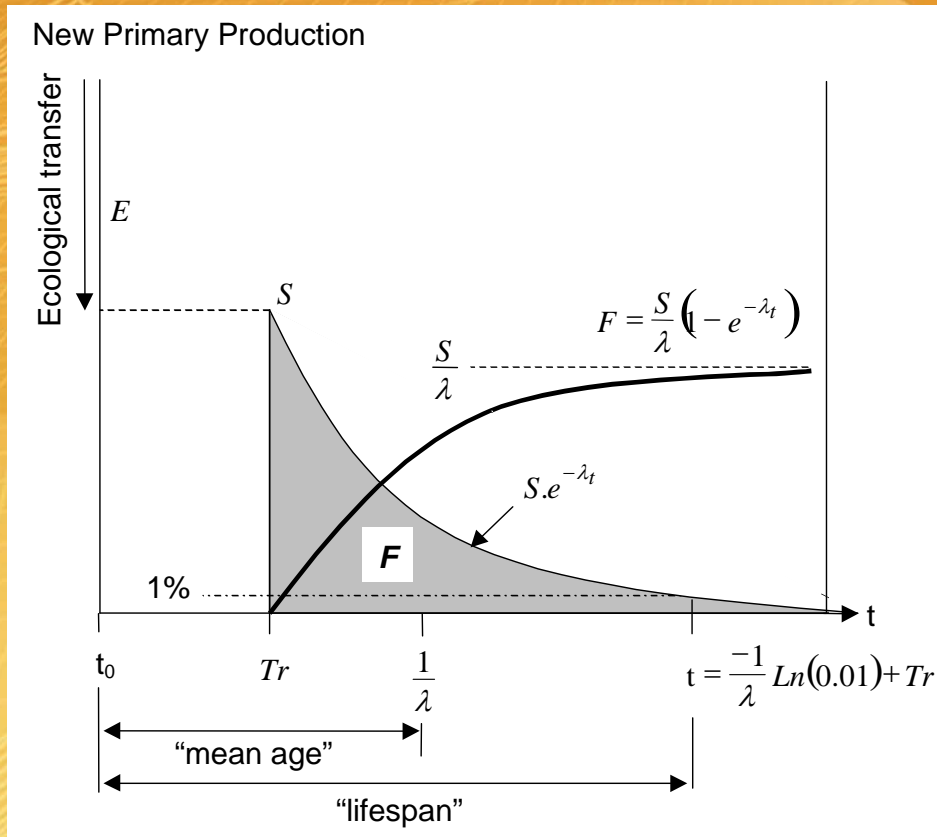
To each component is associated a coefficient of energy transfer from primary production (PP)

Modelling forage components: 3-layer 6-forage functional groups



		Forage component					
		epi	meso	m-meso	bathy	m-bathy	hm-bathy
Vertical layer	Epi-pelagic	1	0	0	0	0	0
	Meso-pelagic	0.307	0.237	0.456	0	0	0
	Bathy-pelagic	0.17	0.1	0.22	0.18	0.13	0.2

Forage functional groups: Dynamics is based on the 3T: Time, Temperature and Transport (currents)



Lehodey P. et al., 1998. *Fisheries Oceanography* 7(3/4): 317-325.

Lehodey P. 2001. *Progress in Oceanography* 49: 439-468.

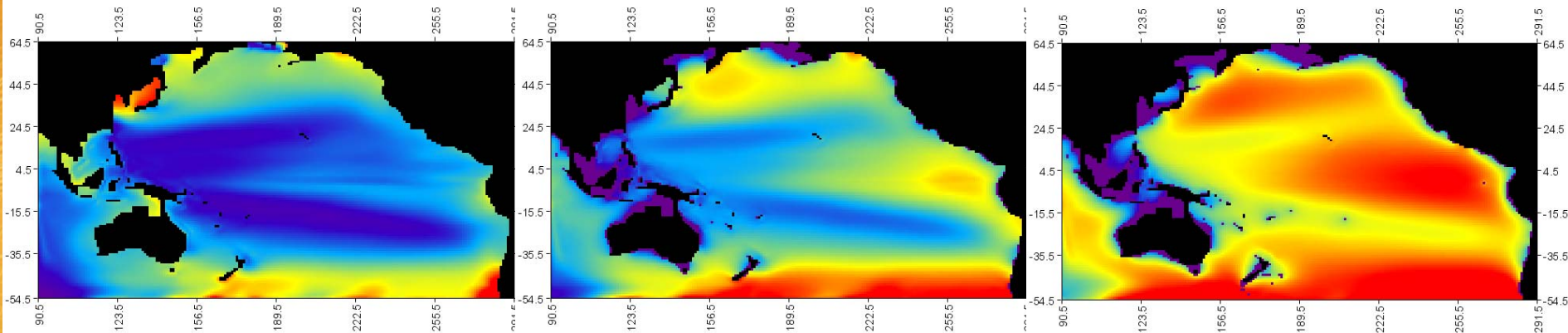
Lehodey P., Chai F., Hampton J. 2003. *Fisheries Oceanography* 12(4): 483-494

Simulation outputs

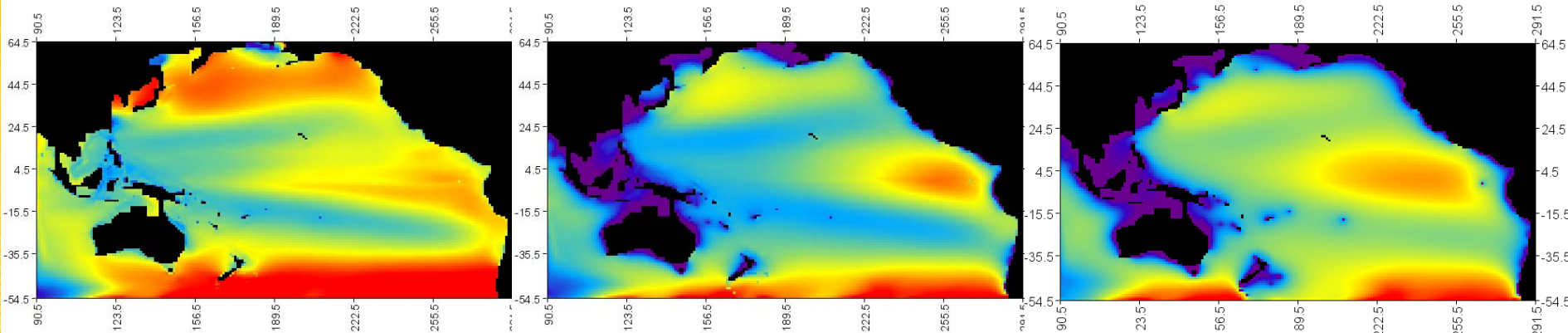
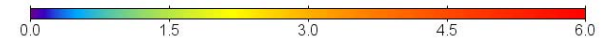
Epi-pelagic layer
(0-100m)

Meso-pelagic layer
(100-400m)

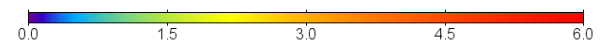
Bathy-pelagic layer
(400-1000m)



Day



Night



Predators dynamics modelling



Structure and dynamic of tuna populations

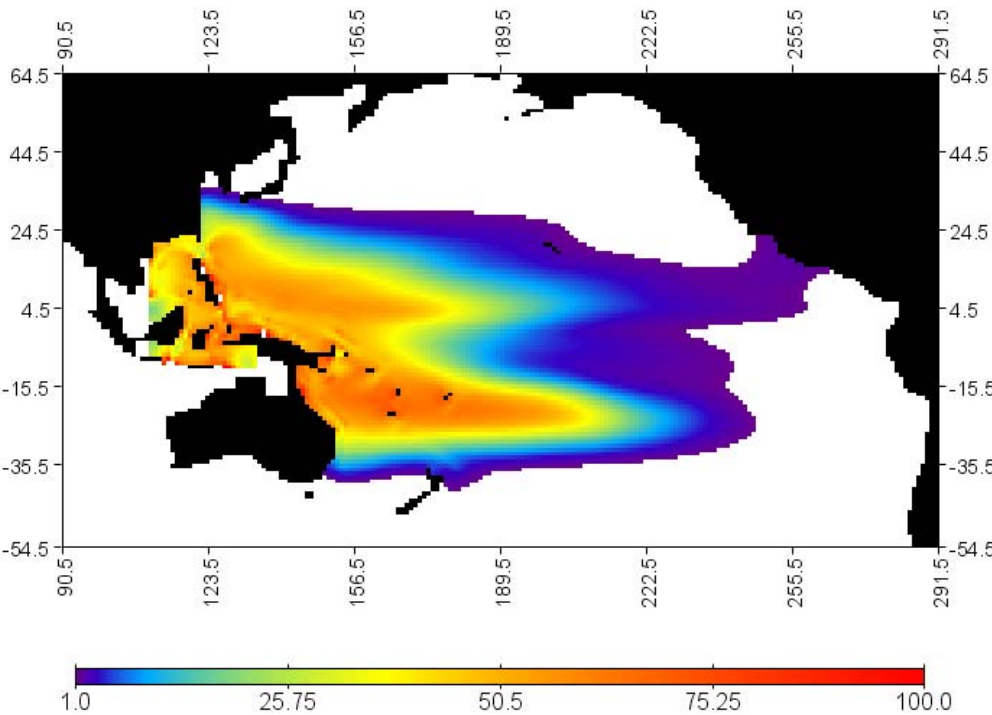
	Spawning	Larvae	Juvenile	Young	Adult
Time / age structure	t ₀	1 st month	2 nd and 3 rd month	2 nd quarter to age of 1 st maturity	1 st maturity to last quarter
Size	2 mm	2 mm -5 cm	5-15 cm	15 - > 40 cm	> 40 cm
Transport / movement (advection-diffusion)		Currents in upper layer		1- Proportional to fish size 2- Random movement (Diffusion) decreasing with increasing habitat 3- Directed movement (Advection) following increasing gradient 4- impact of currents	
Habitat factors	T°, Food (P), Predators (F) in the epi-pelagic layer		T°, Food (Zpk), Predators (all young and adult tuna)	T°, oxygen, Food (F), Predators (all adult tuna) in all layers	T°, oxygen, Food (F) in all layers, spawning seasonality
Natural mortality		Independent estimates + habitat-related variability			
Growth		Independent estimates			

highlights

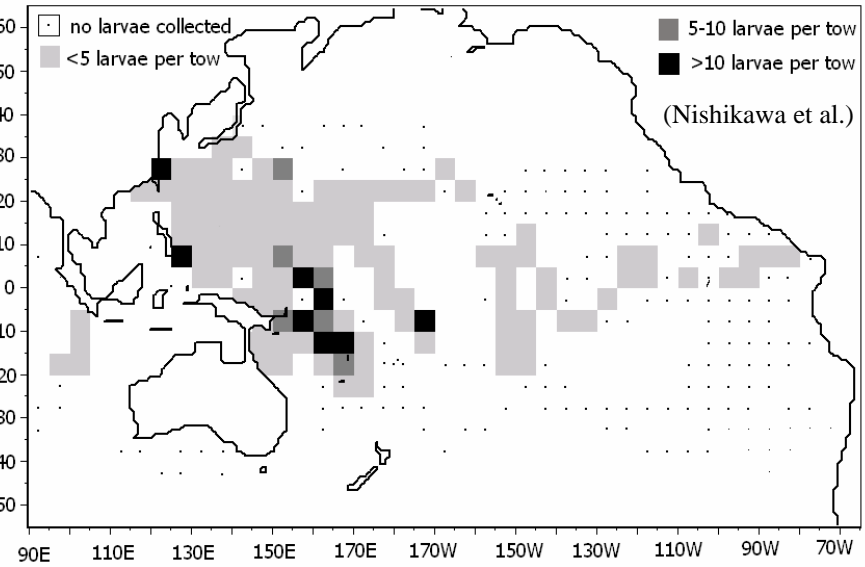
- Habitats
- Movement
- Spawning seasonality
- Variability of natural mortality
- Prey-predator coupling

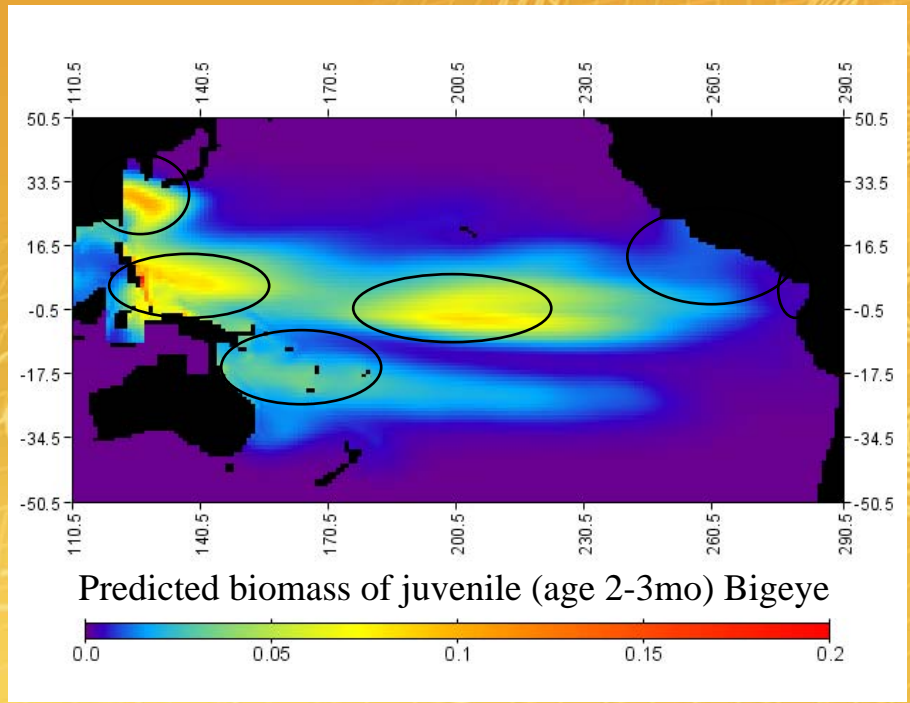
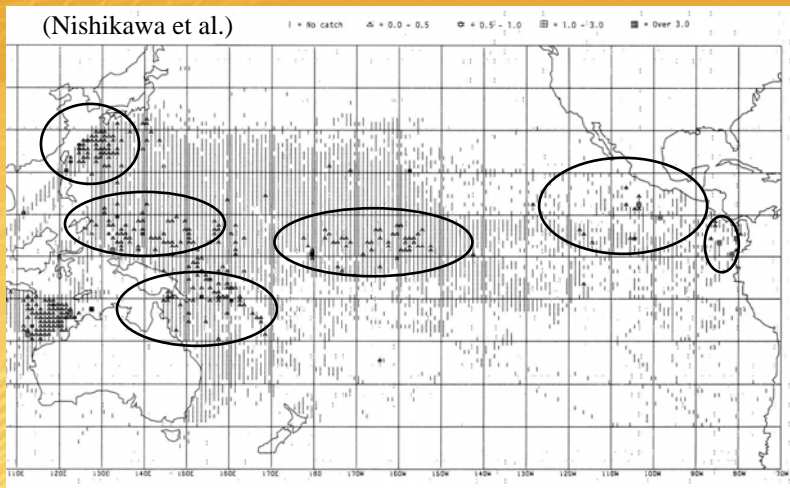
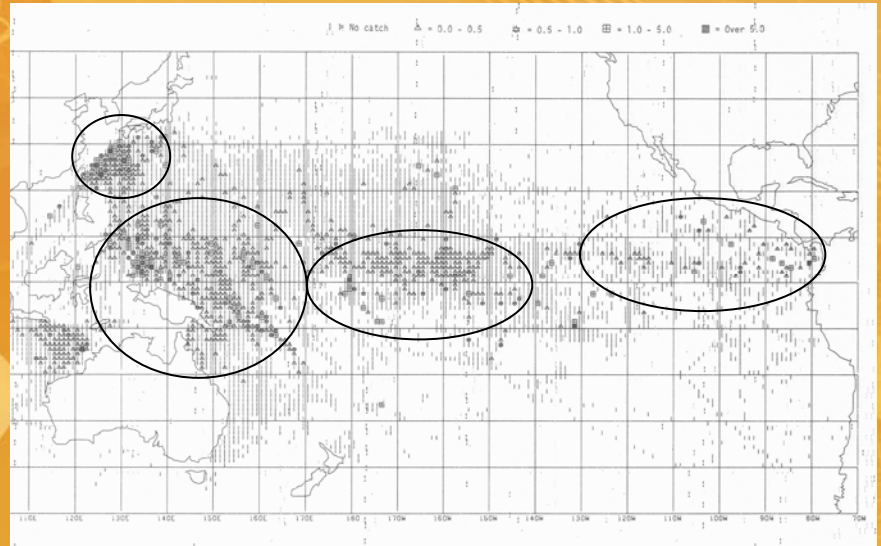
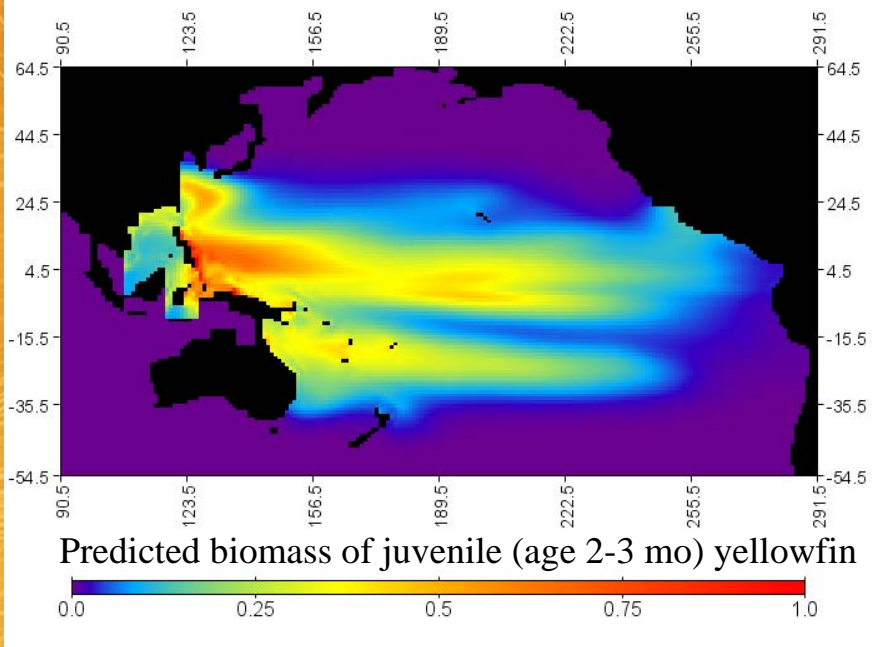
Spawning Habitat

$$H_s = R_s \cdot I_{\theta_s} \cdot e^{\alpha \cdot \log \left[1 + \frac{P}{F_l} \right]}$$



distribution of skipjack larvae

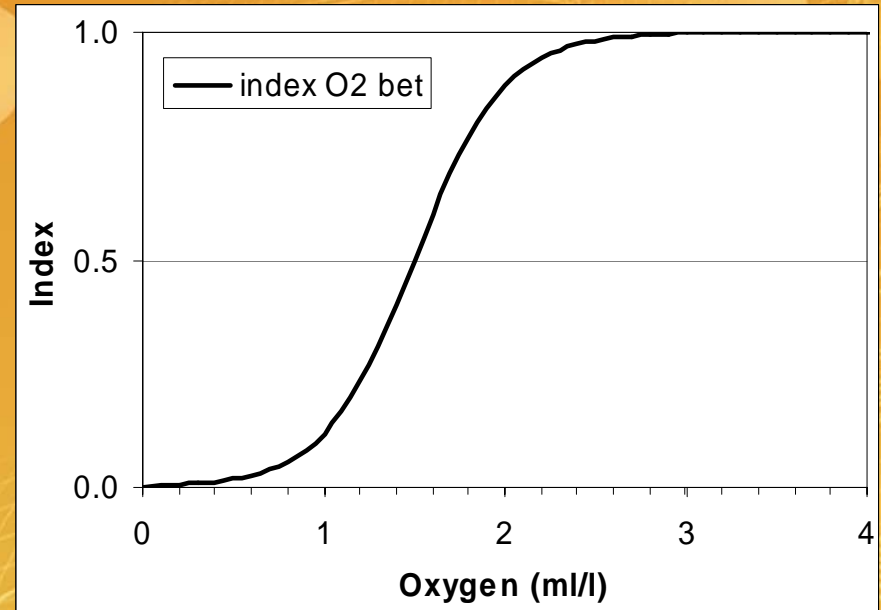
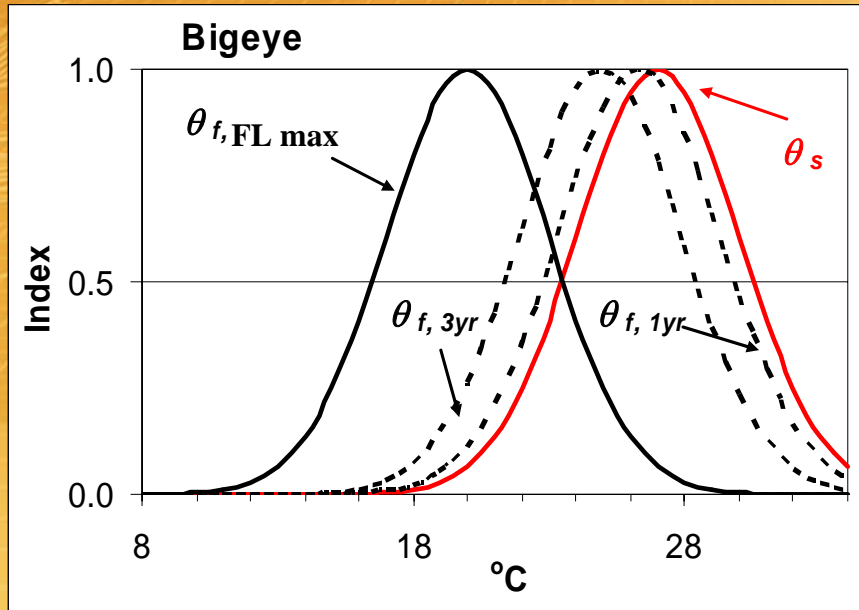




Simple match-mismatch mechanism, ie PP/ F, embedded in a dynamic system (currents, temperature) creates complex but realistic results

Feeding habitat

Defined by the accessibility to the different forage components according to preference of species (by age)



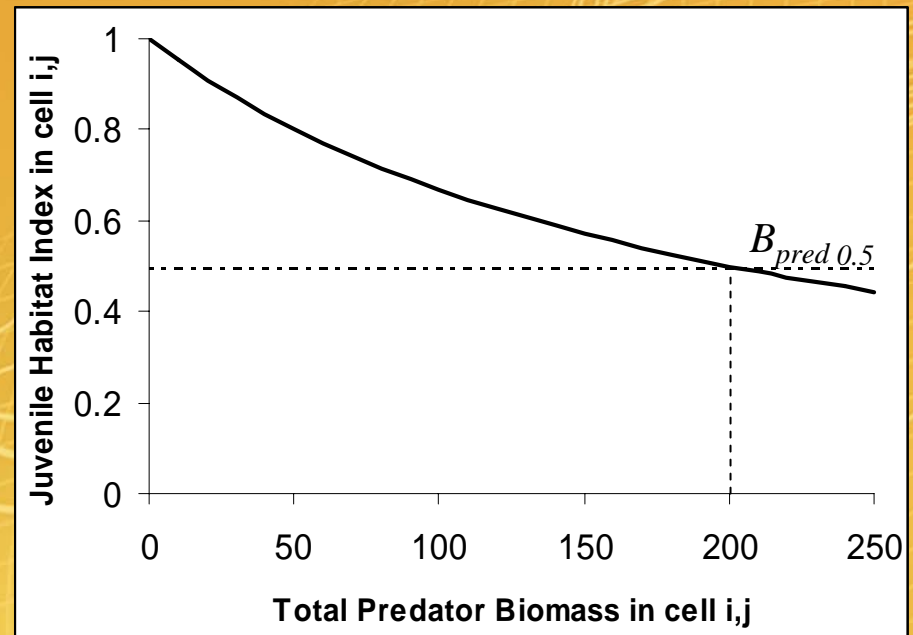
Change in temperature function with age from age 0 (spawning) to maximum age (left) and habitat function for the oxygen (right).

Juvenile habitat

$$H_{juv\,i,j} = I_{\theta\,juv\,i,j} \cdot \left[1 - \frac{B_{pred\,i,j}}{B_{pred\,0.5} + B_{pred\,i,j}} \right]$$

$$B_{pred\,0.5} = 250 \text{ t deg}^{-2}$$

~ max value of biomass of
SKJ+YFT+BET



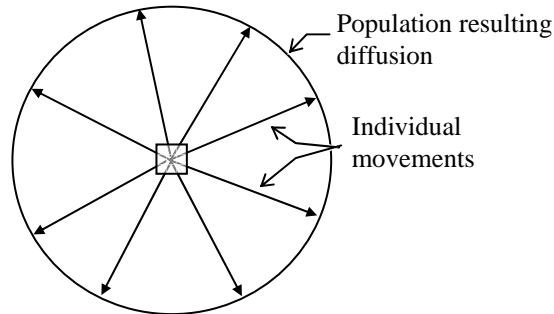
Movement

Advection – directed movement + current effect

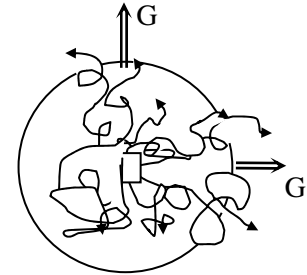
Maximum ($MSS * FL$) for maximal value of gradient of standardized (0-1) adult habitat

Diffusion – random search behavior; maximum if both habitat and gradient of habitat is low

Low if habitat is high or if advection is high



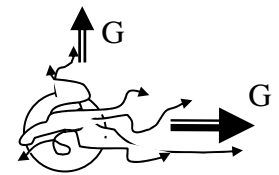
Habitat = null (no gradient)
All displacement is due to kinesis with individuals escaping at MSS in any straight direction. Population diffusion is maximal



Habitat = medium (medium gradient)
Displacement is due to both kinesis and klinotaxis. Population diffusion and advection are medium



Habitat = high (no gradient or negative gradient)
All displacement is due to kinesis, but population diffusion is low since individuals stay in this favorable area



Habitat = low (high gradient)
Displacement is mainly due to klinotaxis. Population diffusion is low and advection is high

Movement

Advection = directed movements along Habitat gradient
(Klinotaxis)

In x direction: $A = u + X \cdot G_x$

Current effect % to time spent in different layers

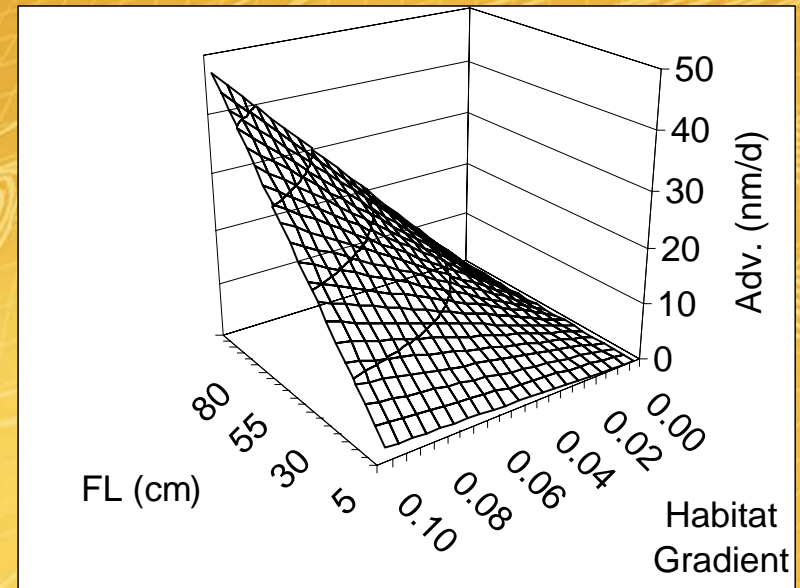
Gradient of Habitat

$MSS = 1 \text{ BL} \cdot \text{s}^{-1}$

$$X = \frac{1}{G_{\max}} \cdot MSS$$

MSS = Maximum Sustainable Speed (in body length. s^{-1})

G_{\max} = max gradient of the standardised Habitat

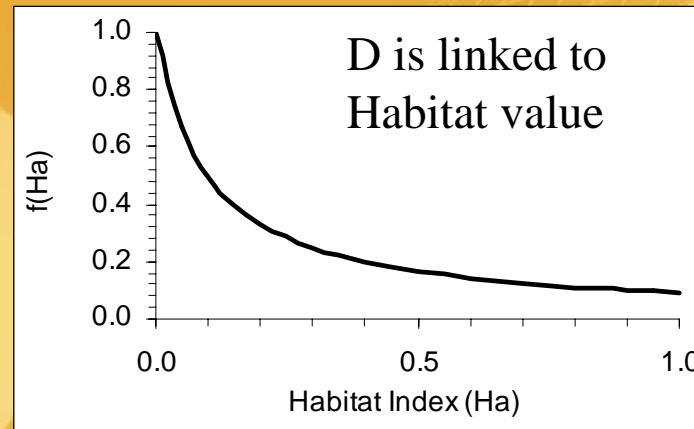


Movement

Diffusion = random movements

$$D = \left(\frac{1}{4} (MSS \cdot FL)^2 t \right) \cdot \left(1 - \left[\frac{H_a}{\beta + H_a} \right] \right) \cdot \left(1 - 0.9 \cdot \left| \frac{G}{G_{\max}} \right| \right)$$

Dmax

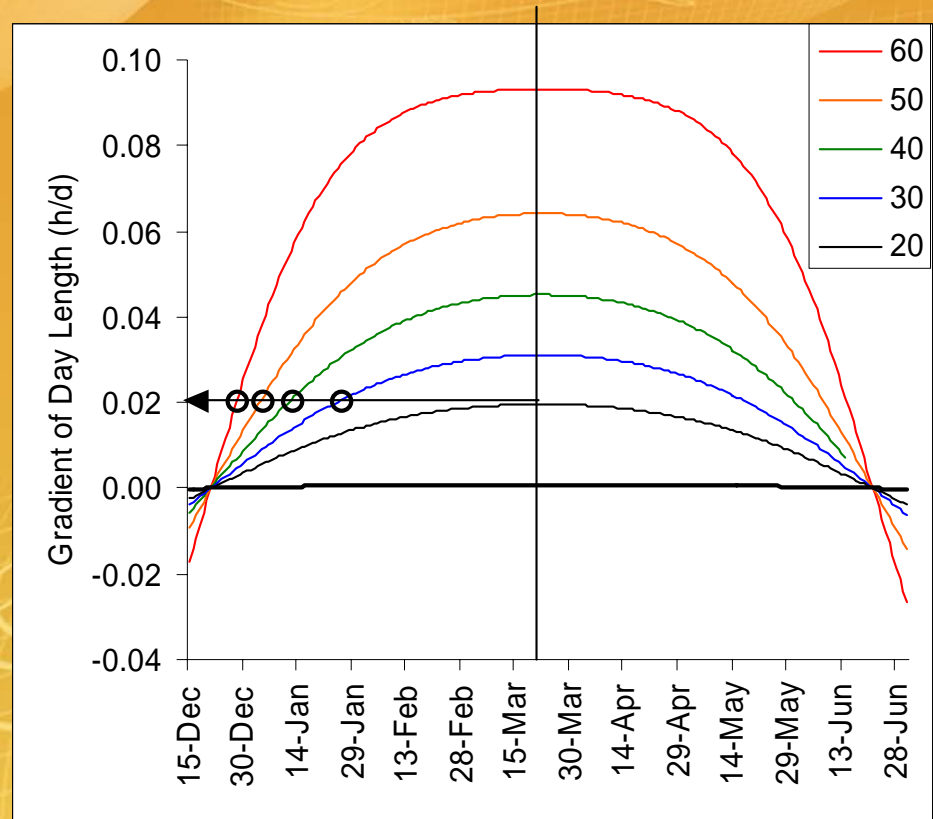


D decreases when gradient increases

With FL the size (Fork Length) in m, MSS the Maximum Sustainable Speed (in Body length.s⁻¹)

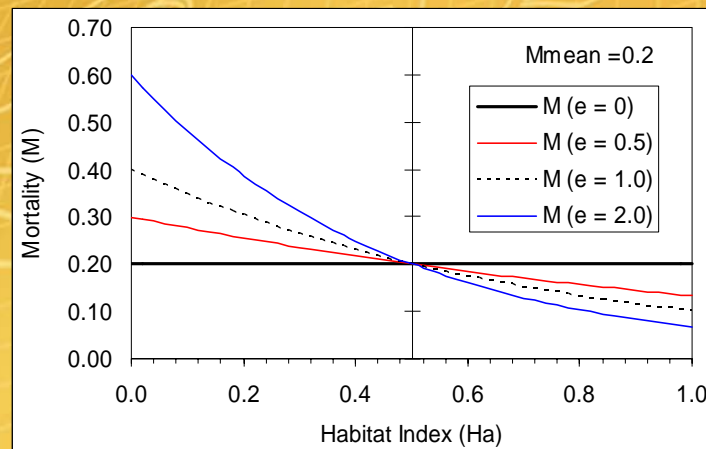
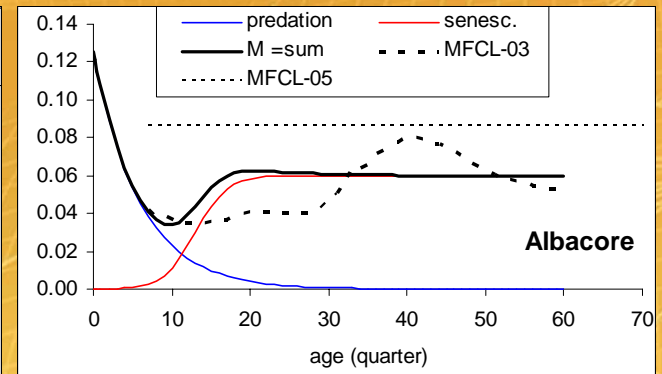
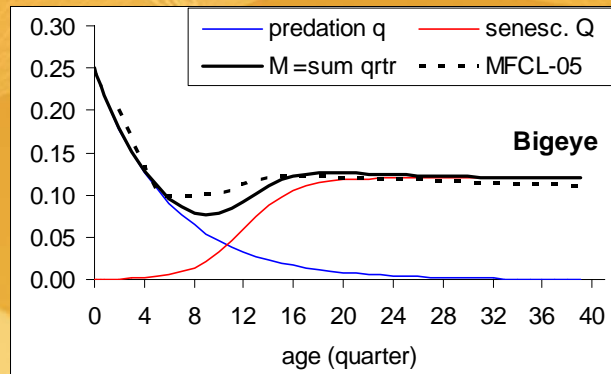
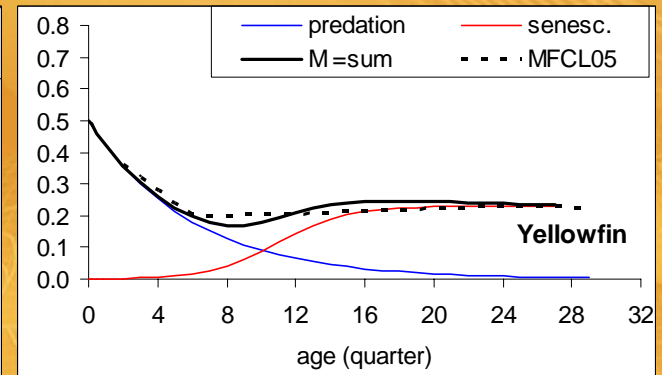
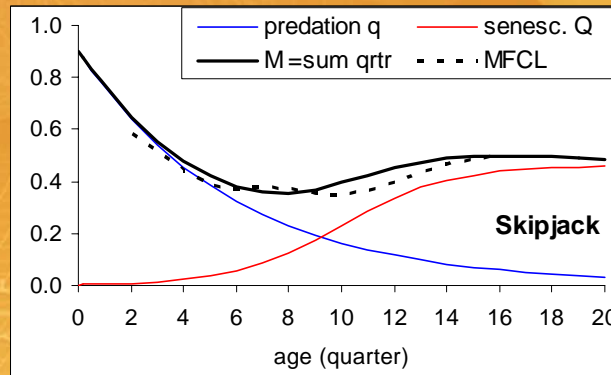
Spawning seasonality

Adult movement based on feeding habitat but switch to the spawning habitat with change in day length. The switch occurs based on a threshold value (>0.03 h/d).



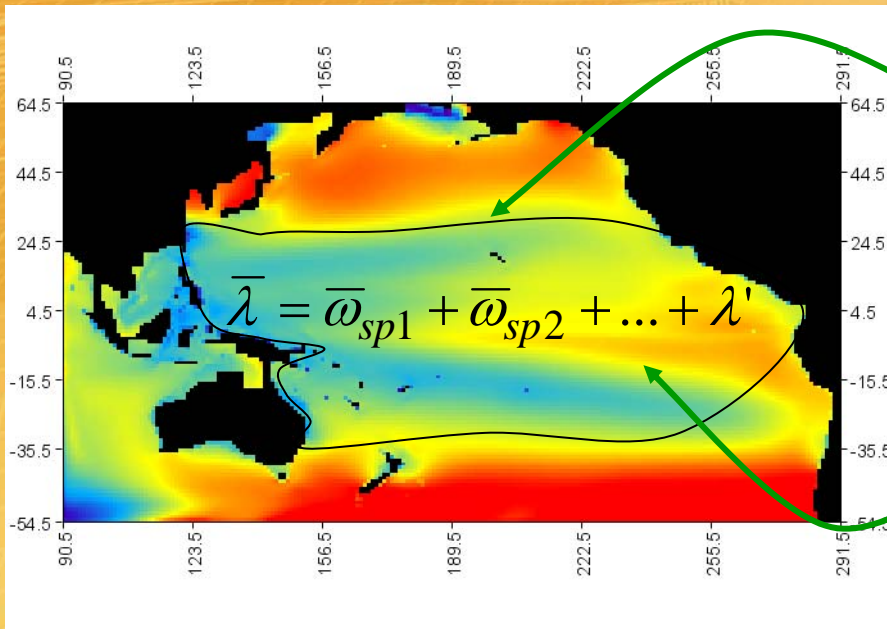
Natural mortality

M is represented by two functions (predation and senescence), and coefficient-at-age can vary in time and space based on habitat value.



Coupling prey (forage) and predators (tuna)

- it is possible to have from zero to N potential predators species explicitly described in the model.
- As a counterpart, this is relying on the assumption that the predators present an ‘ideal free distribution’, such that the total forage mortality by these species would be equal to $\lambda = f(\theta)$
- *can be considered as an ~ equilibrium state.*



Over the “specific predator area”, the mean forage mortality (for a given component) is the sum of the mortalities due to the predator species described in the model + a residual mortality λ' due to all other predators

Locally, in each cell, the forage mortality due to food requirements of described predators, $\omega_{i,j}$ is calculated according to physical accessibility of the predator species (age) to the forage component considered and to their daily ration (% of body mass)

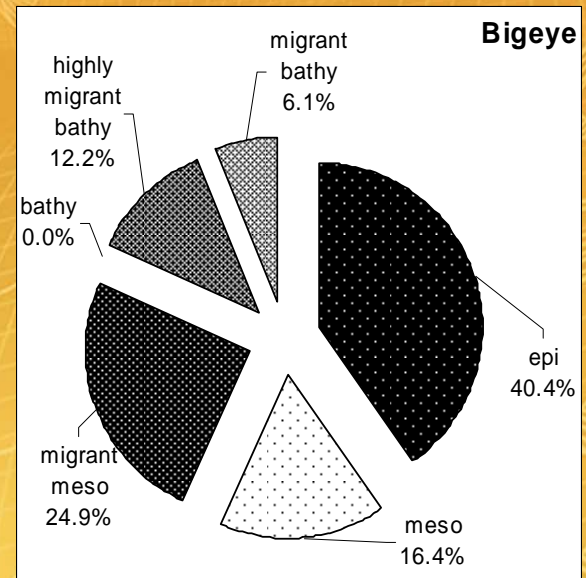
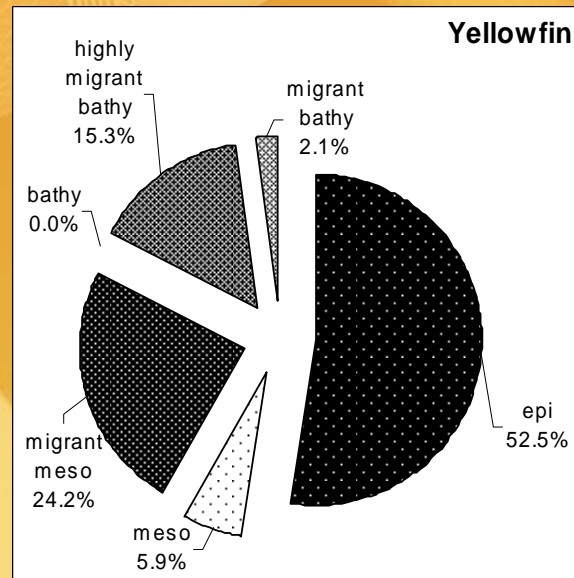
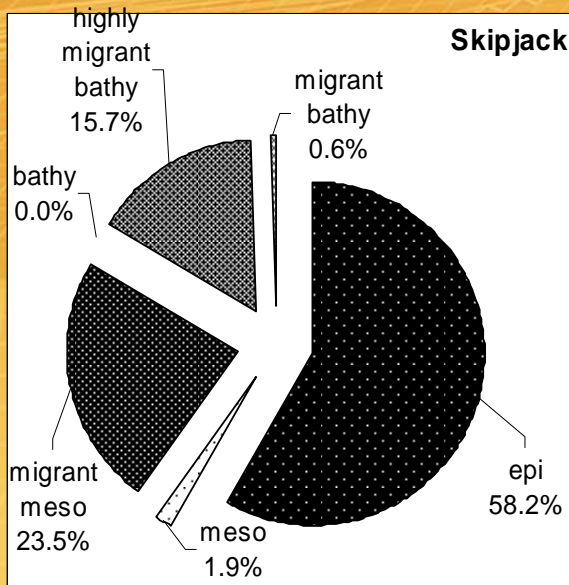
If sum of $\bar{\omega}_{sp}$ above $\bar{\lambda}$ -> ERROR:

biomass of predators cannot be sustained by the forage component

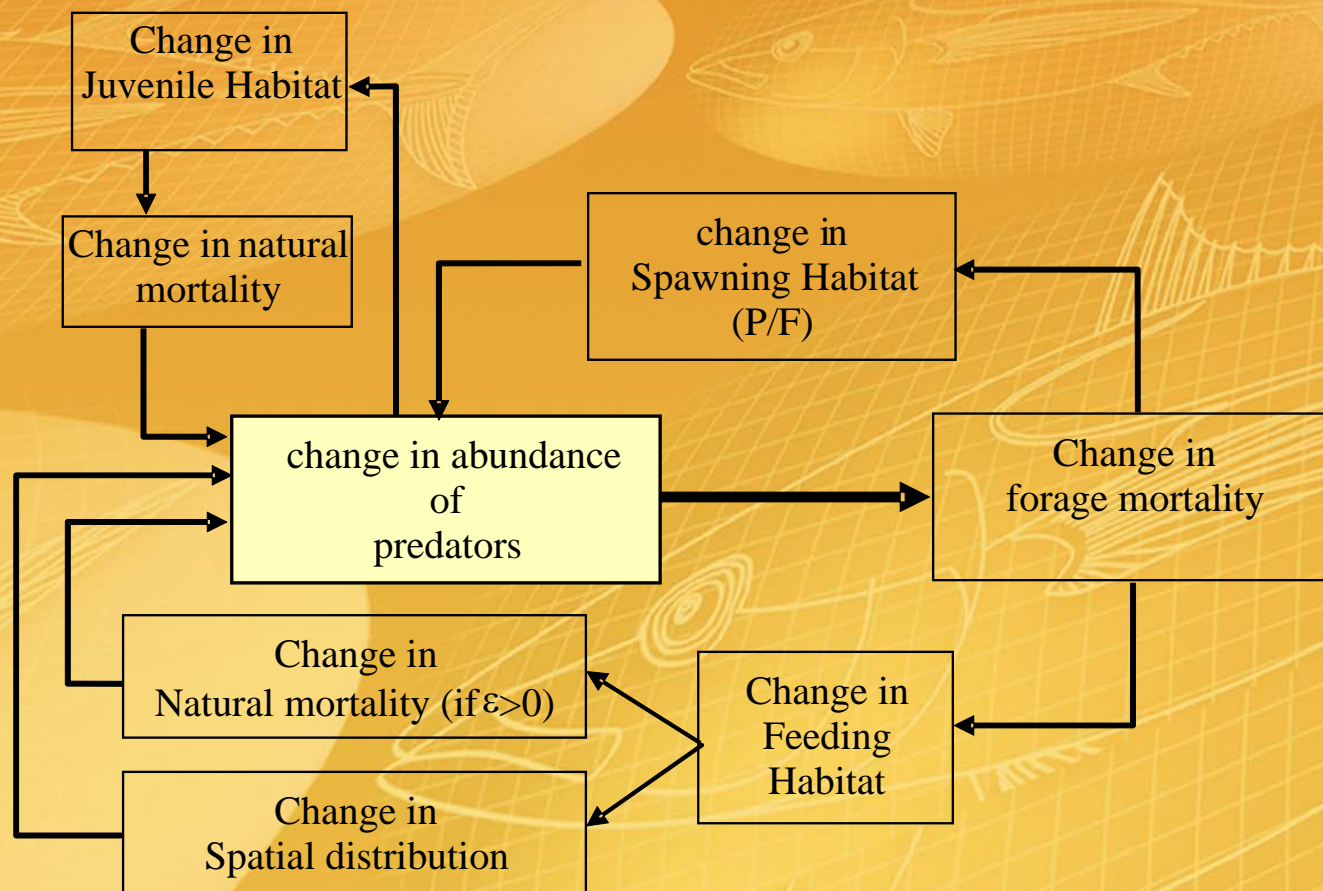
Outside specific predator area $m = \lambda$

Inside specific predator area $m = \omega_{i,j} + \lambda'$

Average forage consumption by species (all age classes) based on accessibility to forage components

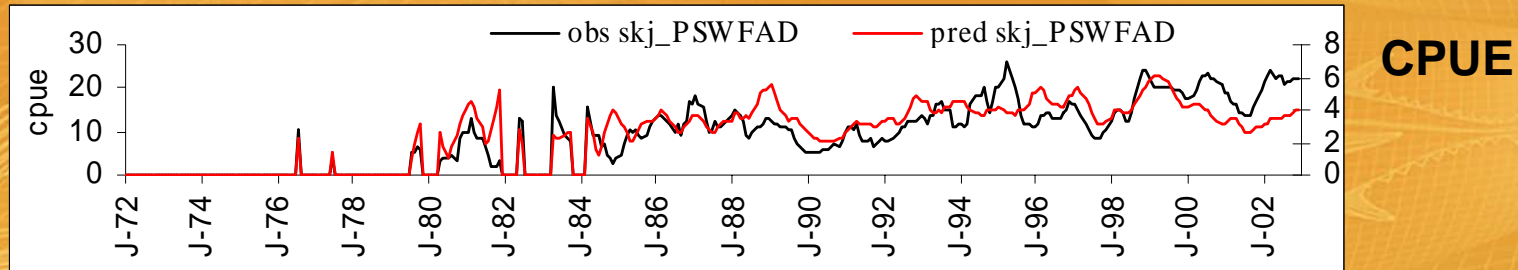


Running single vs multi-species simulations with SEAPODYM: What are the effect of interaction between top predator species like tuna?

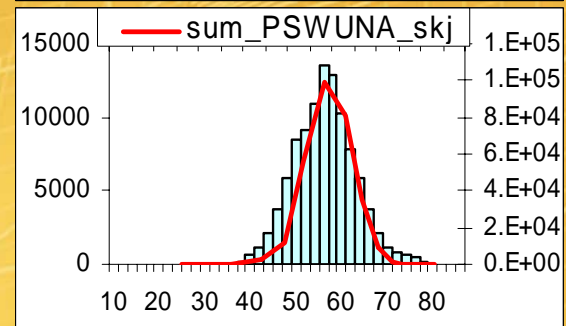
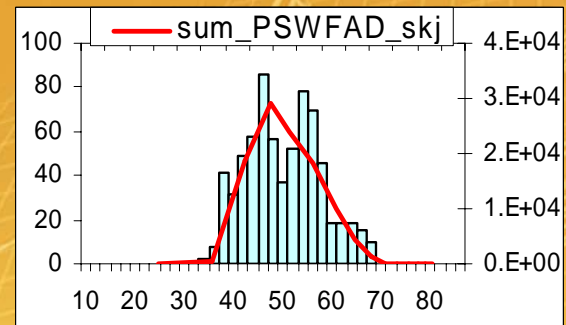
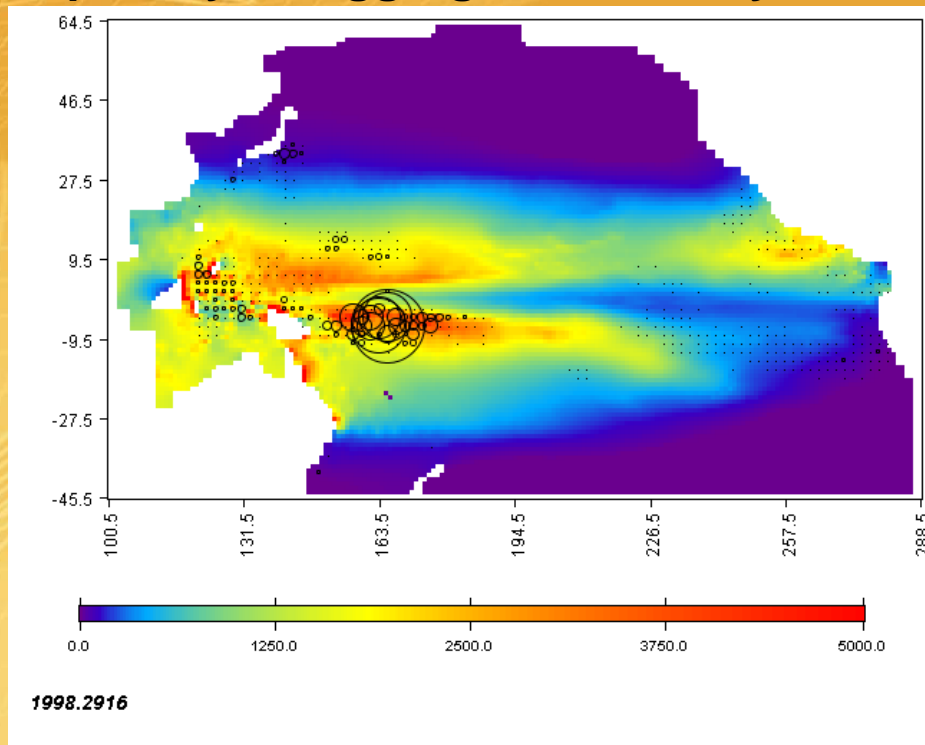


Model outputs and evaluation

Multiple Fisheries → compare Prediction vs Observation



Spatially-disaggregated monthly catch



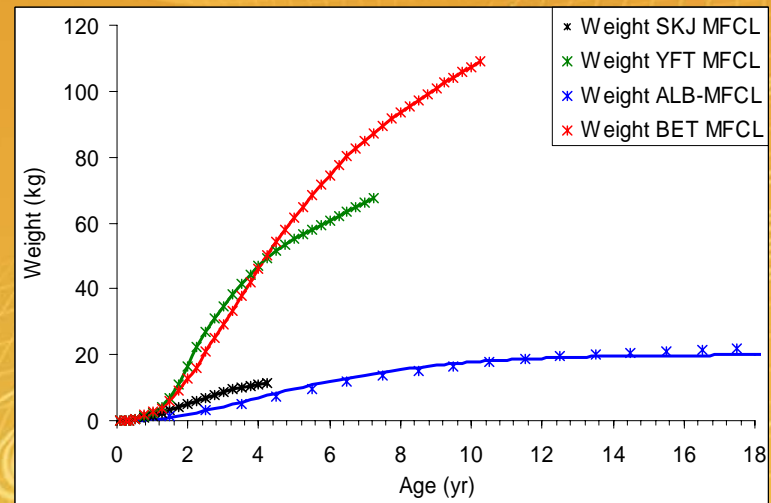
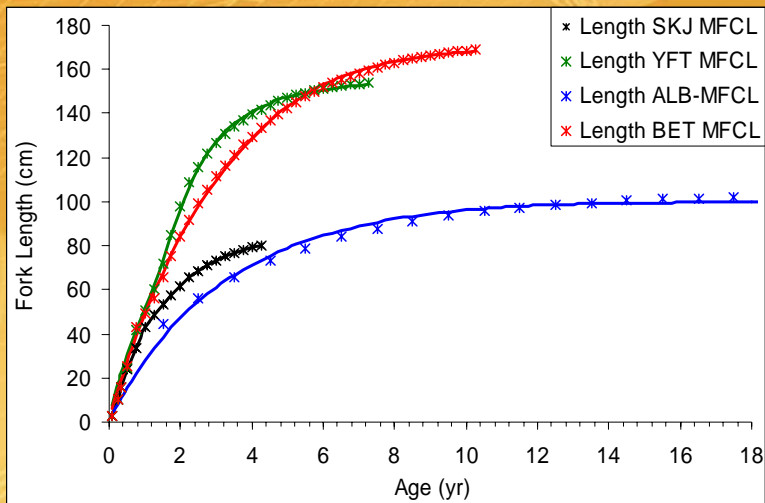
Length-frequency distribution
(by fishery, time and space)

The background is a warm, golden-yellow color. It features several faint, stylized illustrations of fish, including skipjack, yellowfin, and bigeye tuna, rendered in a light, sketchy style. A subtle grid pattern is overlaid on the background, creating a textured effect.

**Application to skipjack,
yellowfin and bigeye tuna**

Table 2. Parameterisation of the populations structure in SEAPODYM

	skipjack	yellowfin	Bigeye	Albacore
Number of age classes (quarter) after juvenile phase	16	28	40	74
Age at first maturity (quarter)	4	7	11	17
Age (quarter) at recruitment	3	3	3	7



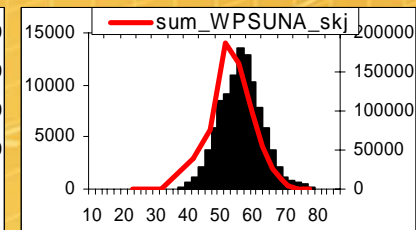
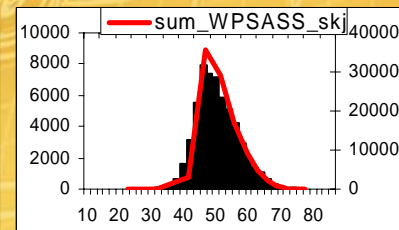
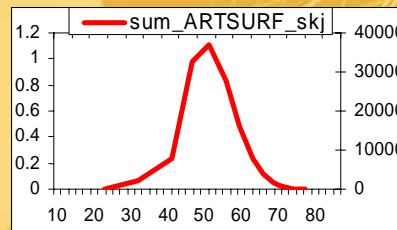
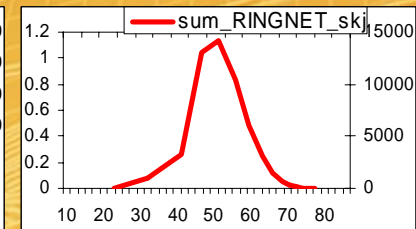
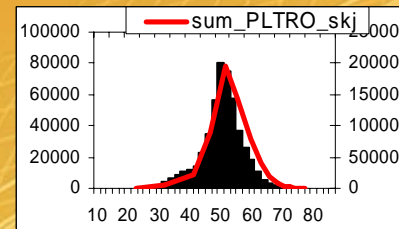
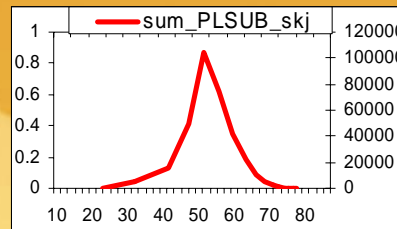
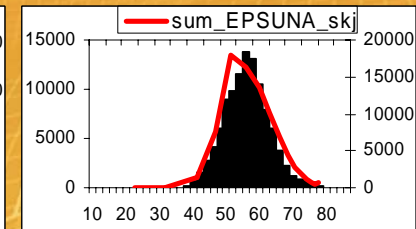
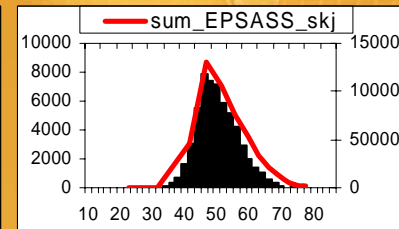
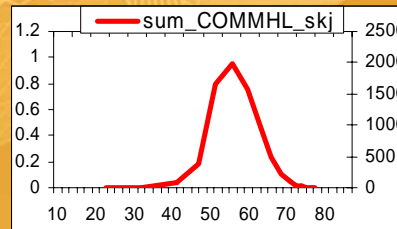
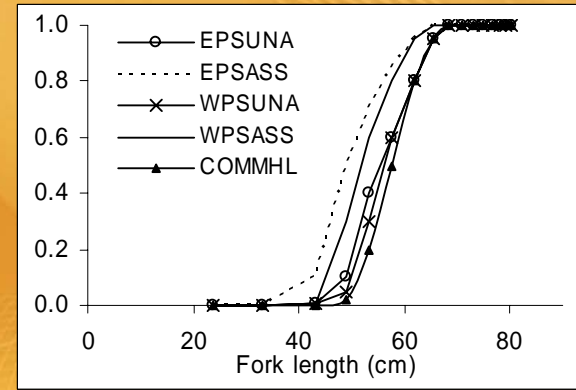
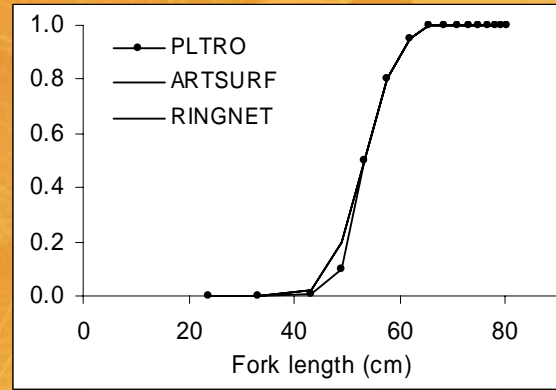
Length-at-age and weight-at-age coefficients estimated from MFCL analyses (crosses) and functions (curves) used to define the coefficient used in SEAPODYM simulations

Fisheries

Category code	Description / source / resolution
PURSE SEINE	
WPSASS	Aggregated data of purse seine fisheries in the WCPO Sets associated to animals, log or FAD
WPSUNA	Aggregated data of purse seine fisheries in the WCPO Unassociated sets (i.e. free schools)
EPSASS	Aggregated data of purse seine fisheries in the EPO Sets associated to animals, log or FAD
EPSUNA	Aggregated data of purse seine fisheries in the EPO Unassociated sets (i.e. free schools)
POLE-AND-LINE	
PLTRO	Aggregated data of tropical (25°N-25°S) pole-and-line fisheries data
PLSUB	Aggregated data of sub-tropical pole-and-line fisheries (mostly Japanese domestic fleets)
LONGLINE	
LLP80	Aggregated data of longline fisheries before 1980 (The pre-1980/post-1980 categories was to (very roughly) define the change from targetting yellowfin to targetting bigeye)
LLSHW	Aggregated data of longline shallow after 1980 (mainly TW and mainland Chinese LL offshore fleets)
LLDEEP	Aggregated data of deep longline fisheries after 1980
LLMIX	Aggregated data of “mixed” longline fisheries after 1980
DIVERSE	
RINGNET	Aggregated data of ringnet fisheries (mainly Philippines, Indonesia)
ARTSURF	Aggregated data of artisanal surface fisheries (including ringnet, mainly Philippines, Indonesia)
COMMHL	Aggregated data of commercial handline fisheries (Philippines, Indonesia, PNG, US)
GILLNET	Aggregated data of gillnet fisheries
TROLL	Aggregated data of troll fisheries

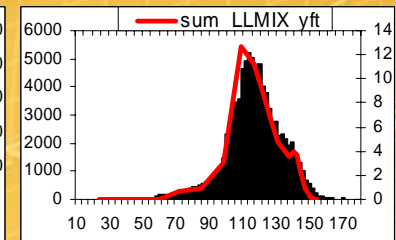
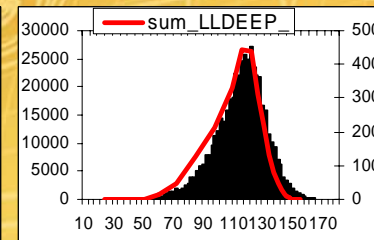
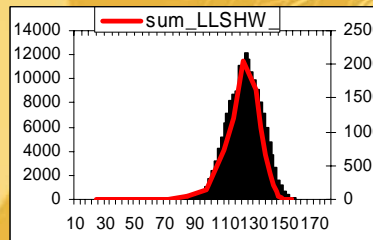
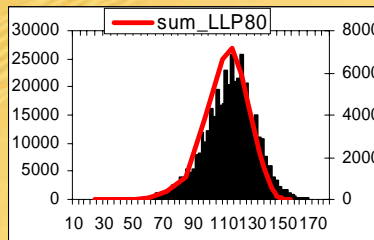
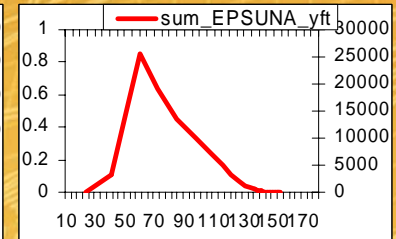
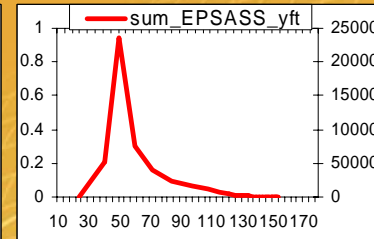
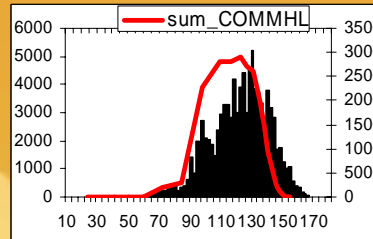
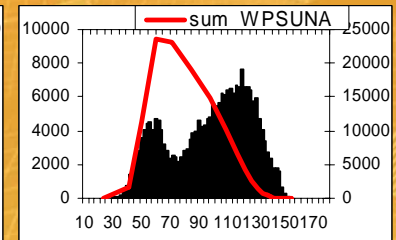
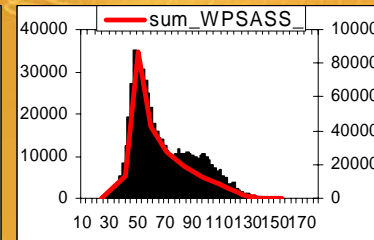
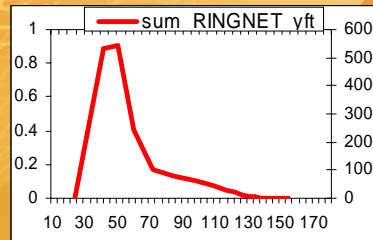
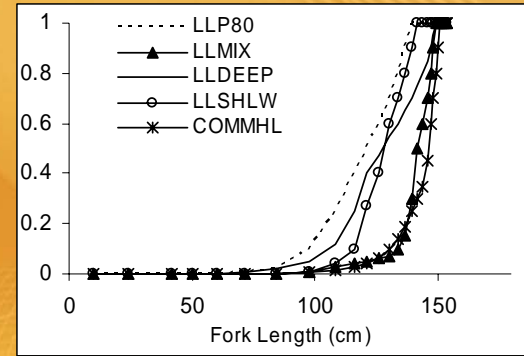
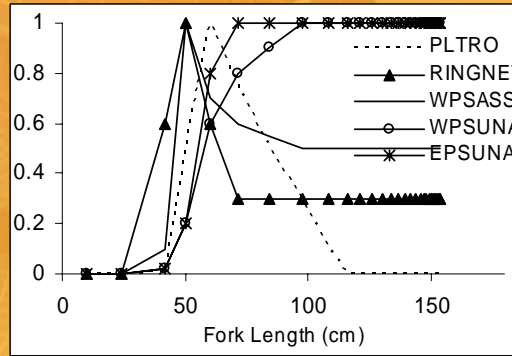
Selectivity

Skipjack



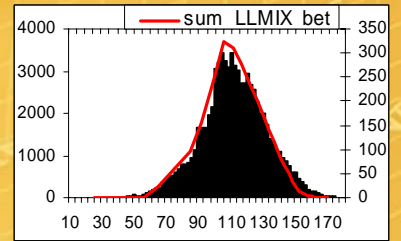
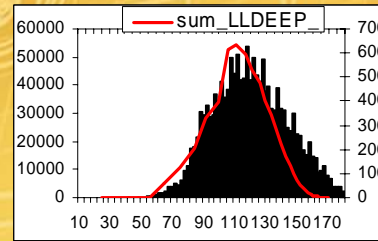
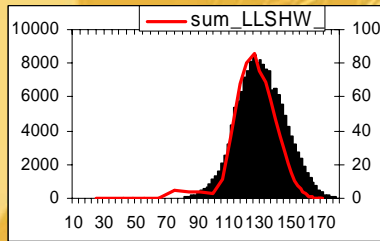
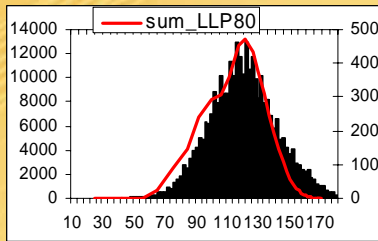
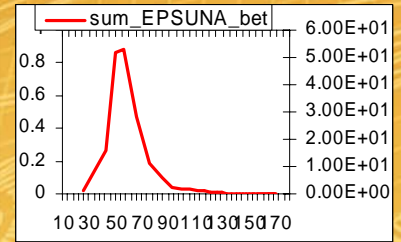
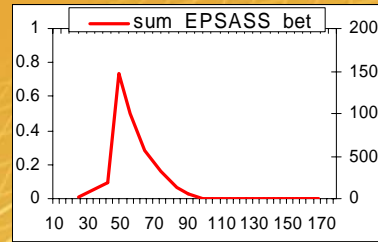
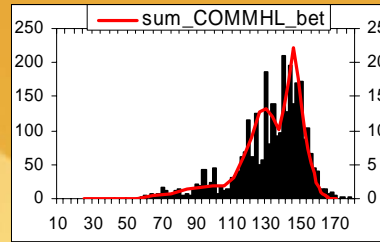
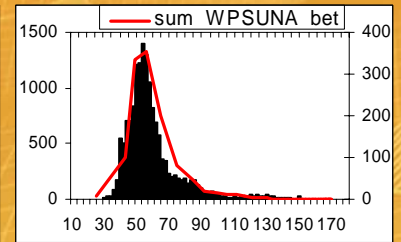
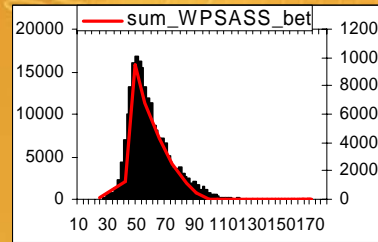
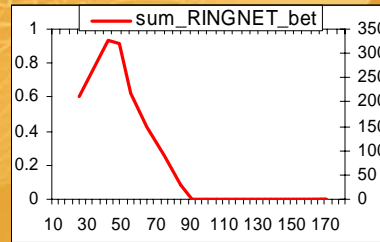
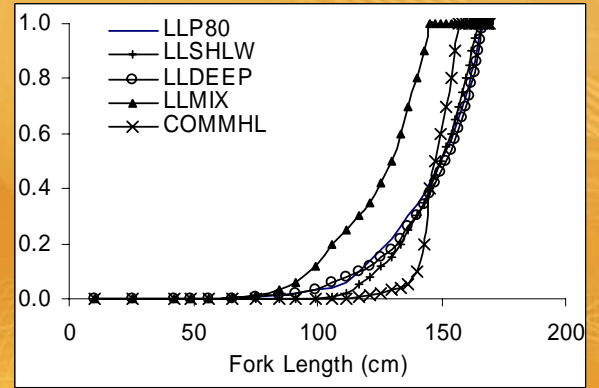
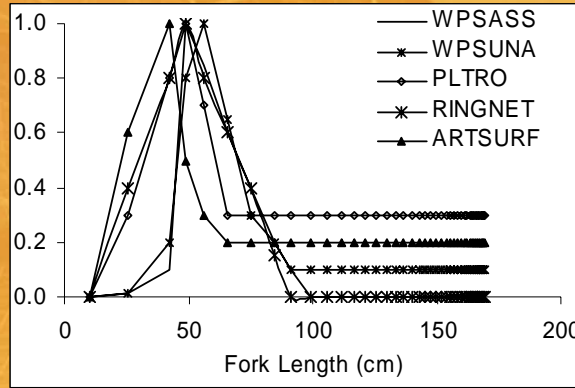
Selectivity

yellowfin



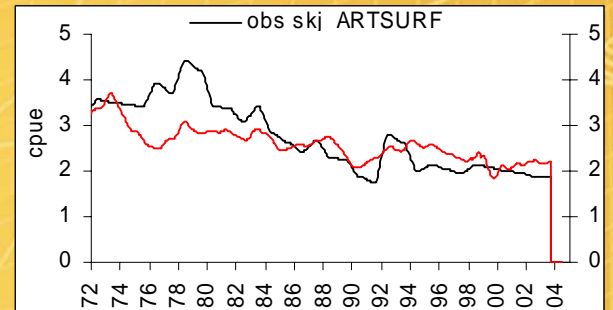
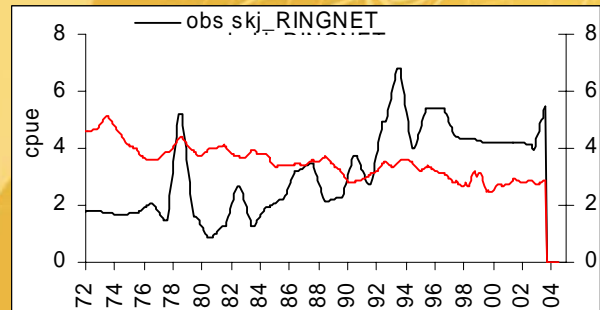
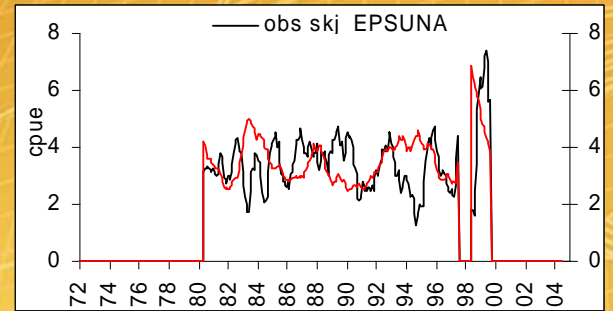
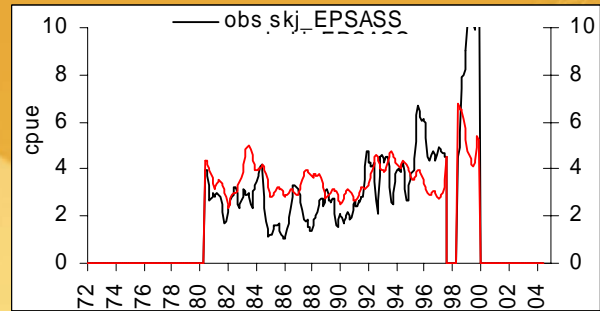
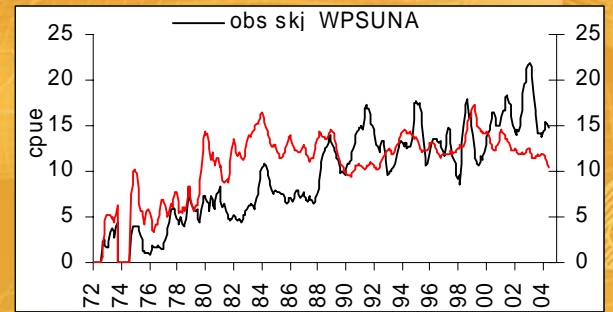
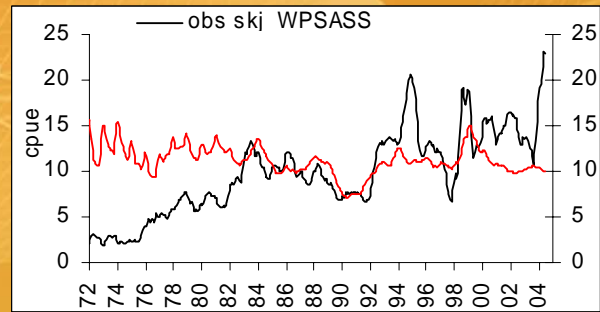
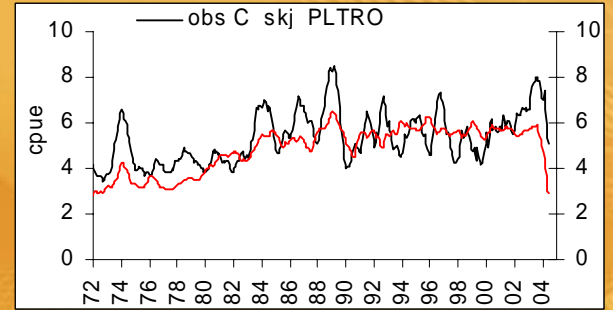
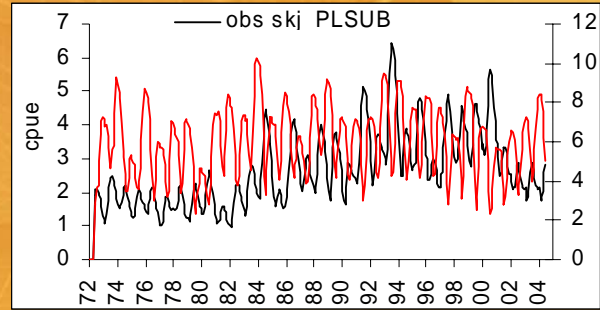
Selectivity

bigeye



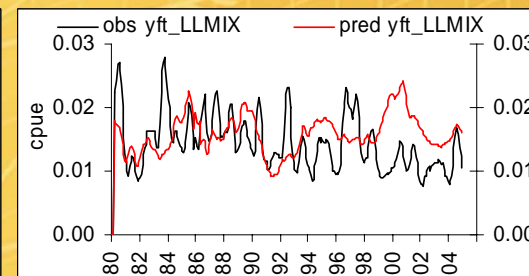
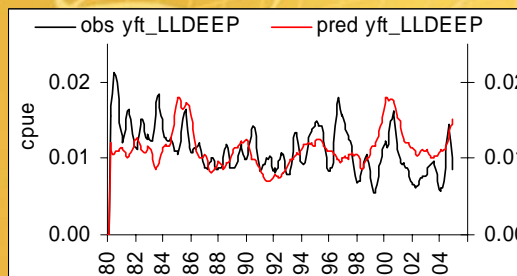
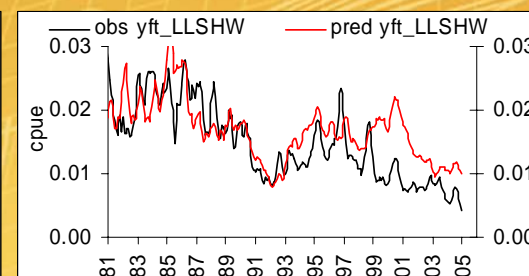
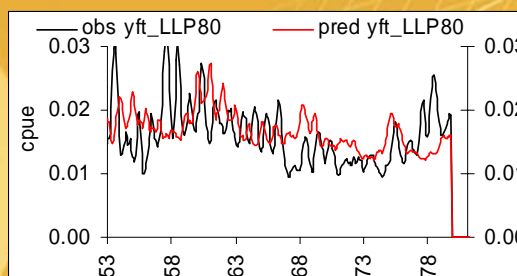
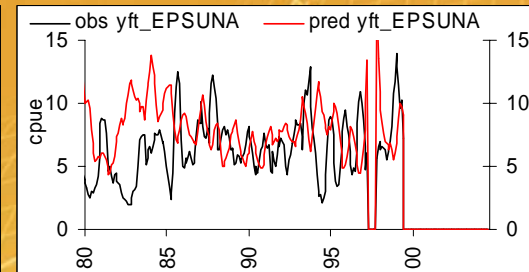
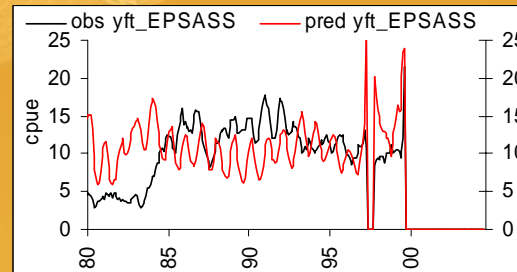
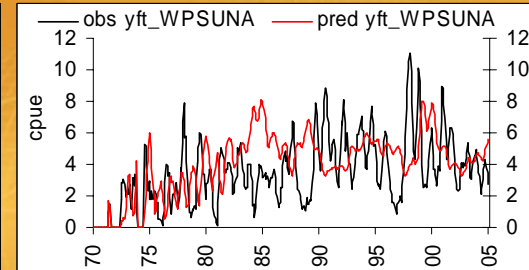
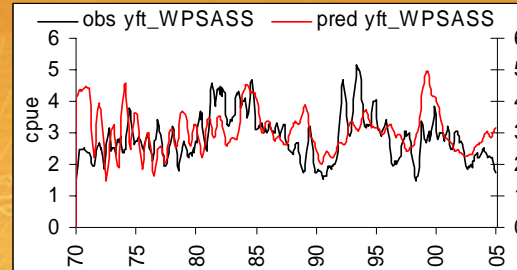
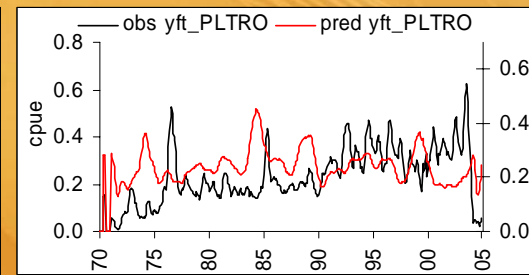
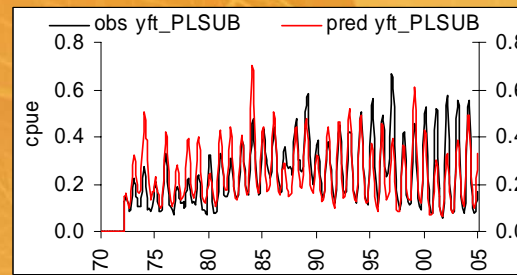
Predicted and observed CPUE

skipjack



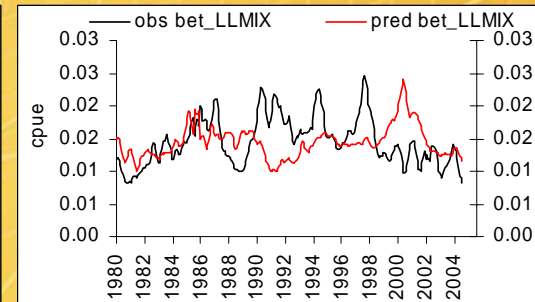
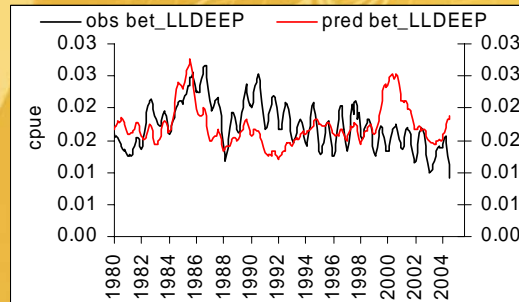
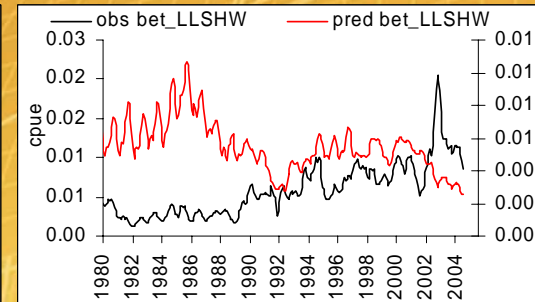
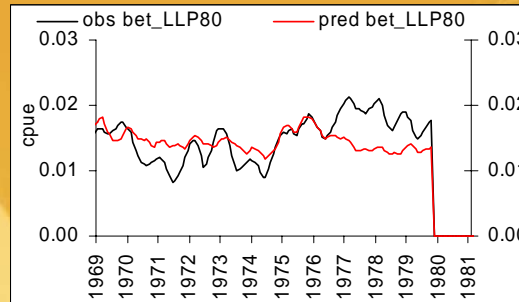
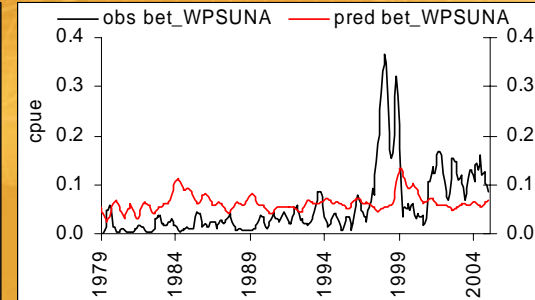
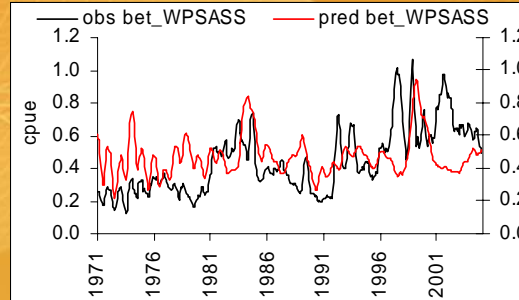
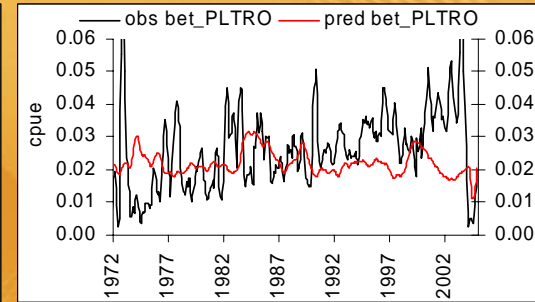
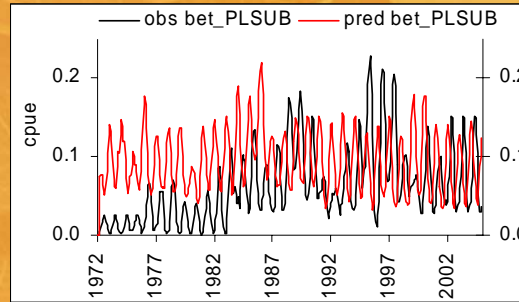
Predicted and observed CPUE

yellowfin



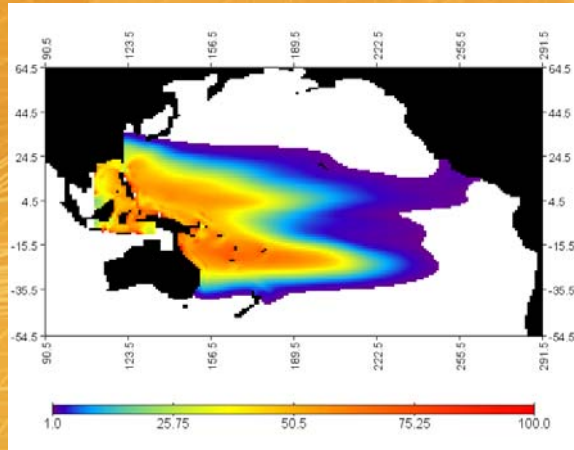
Predicted and observed CPUE

bigeye

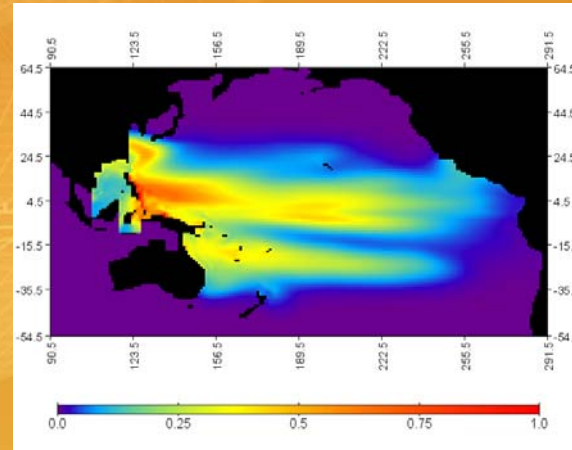


Skipjack

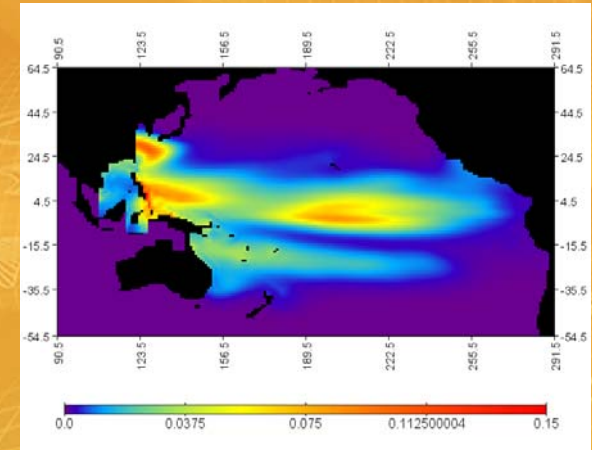
1950-75



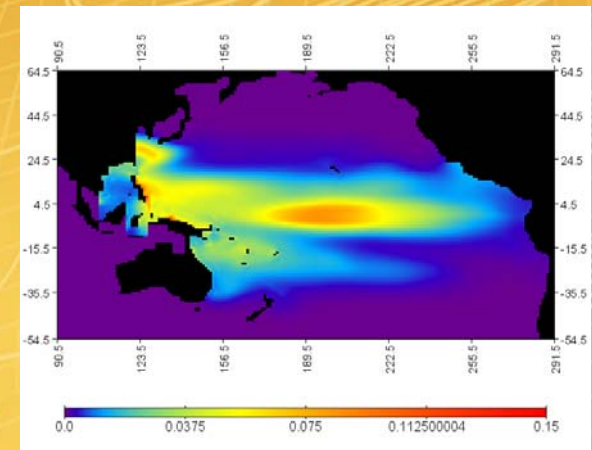
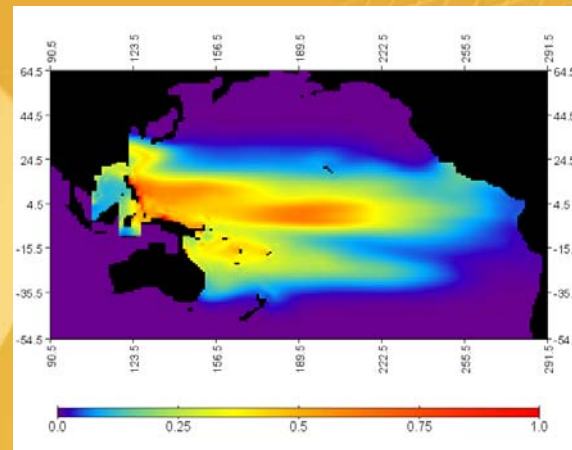
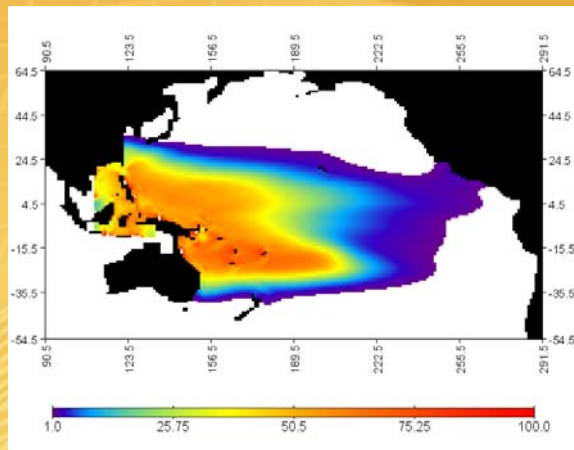
Yellowfin



Bigeye



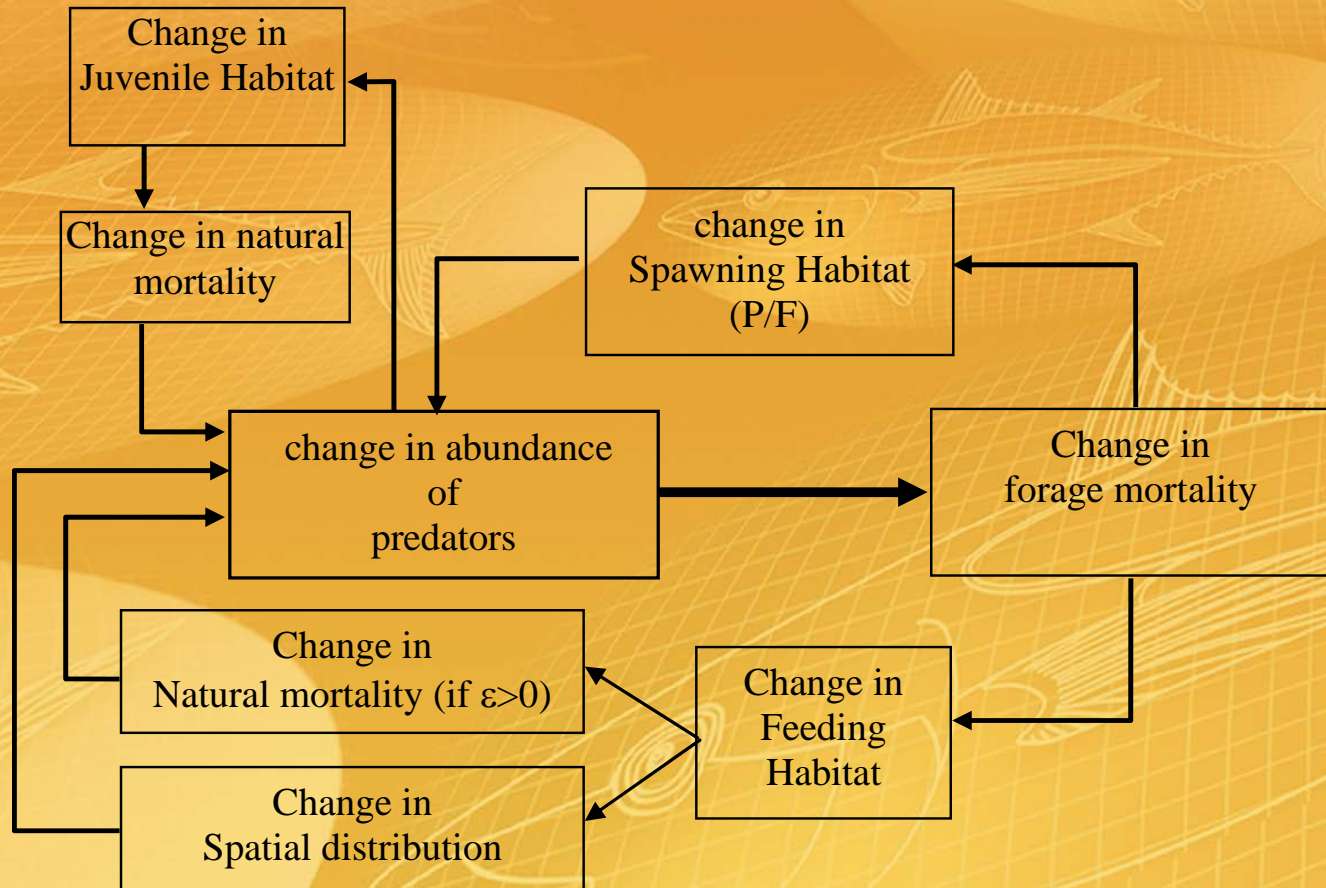
1976-98



Average predicted distribution of juvenile (age 2-3 months) biomass during decadal period 1950-75 and 1976-98

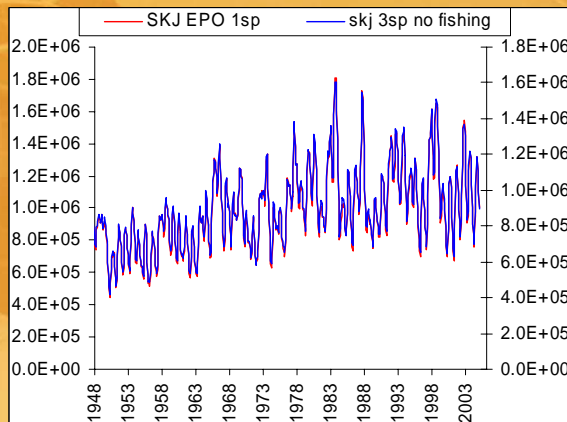
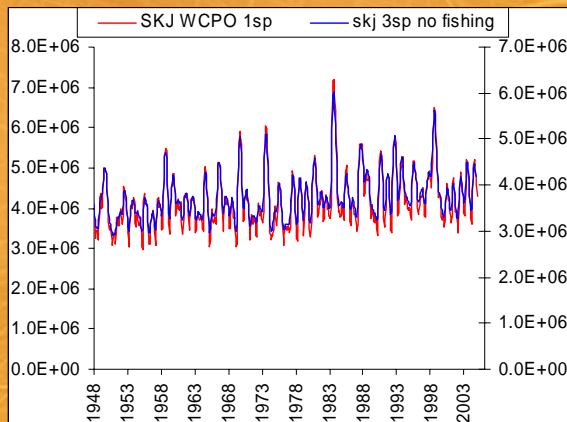
There are large overlaps between spawning and juvenile feeding grounds.

What are the interactions?

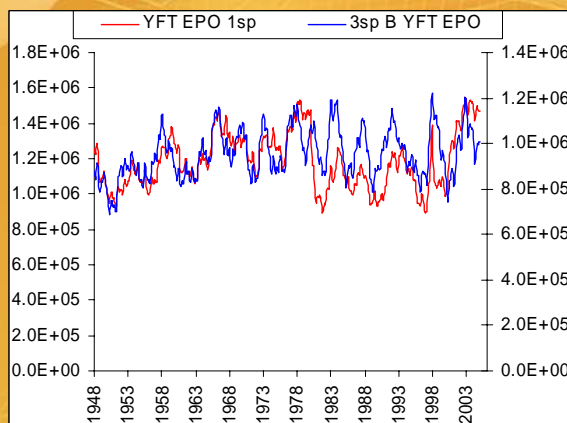
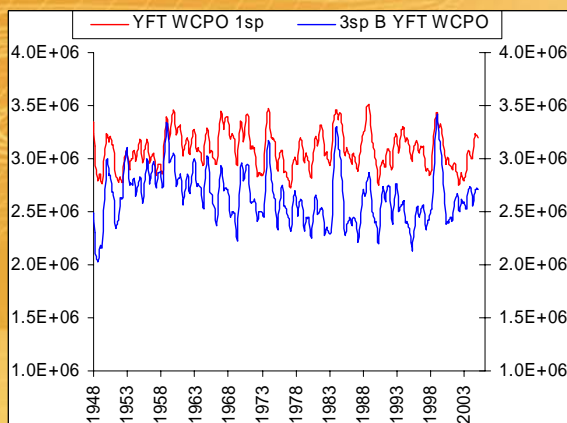


Comparing single vs multi-species simulations

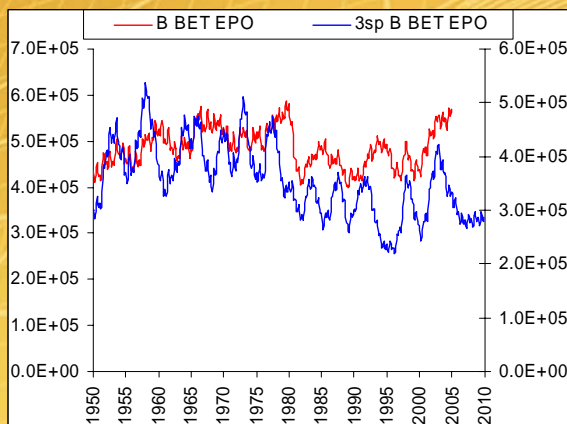
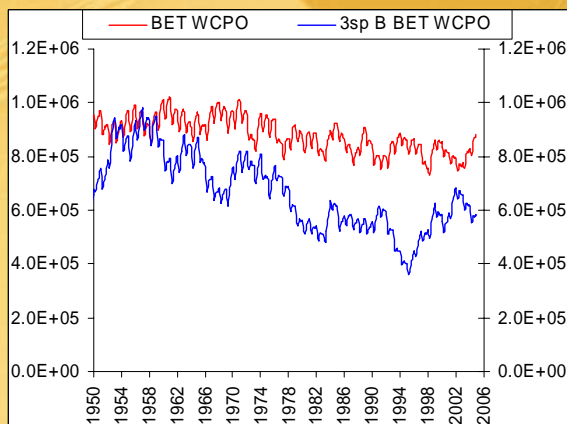
SKJ



YFT



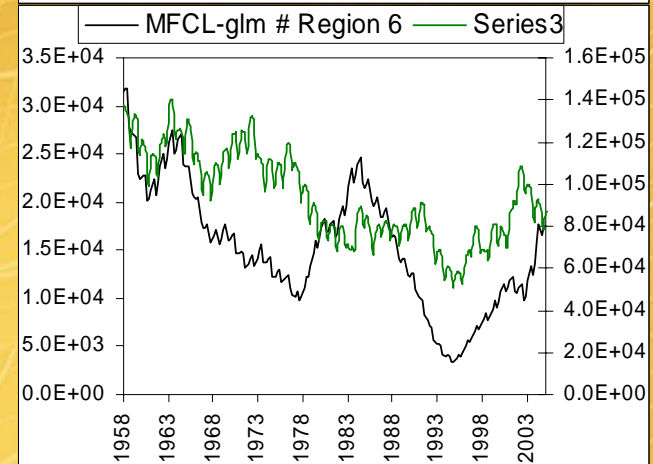
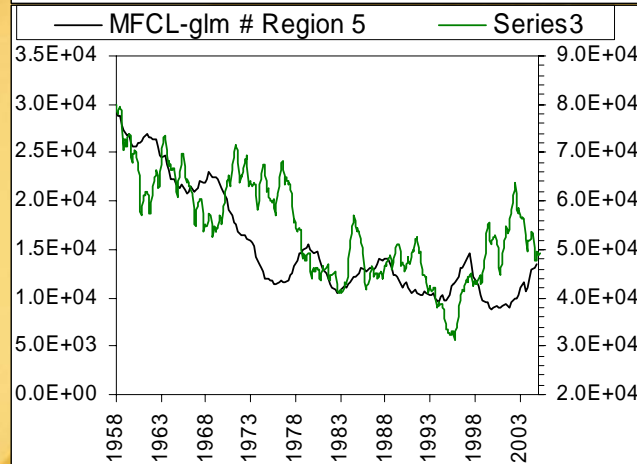
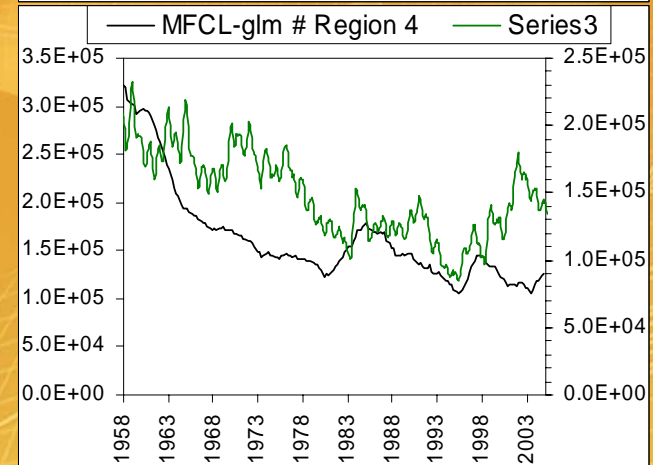
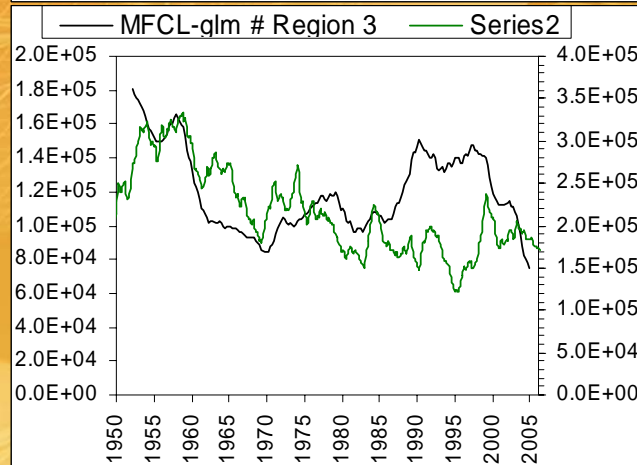
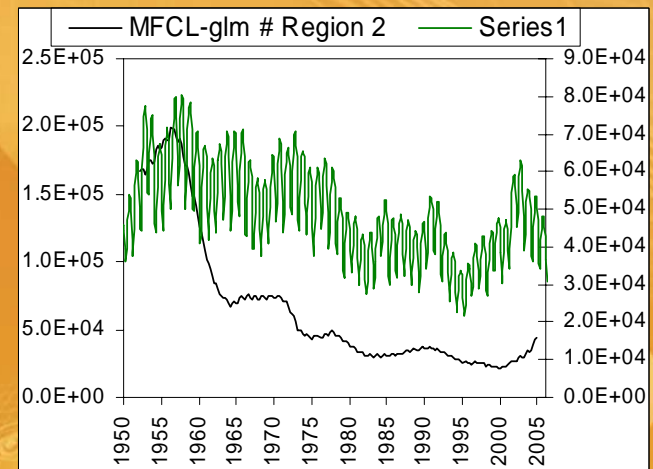
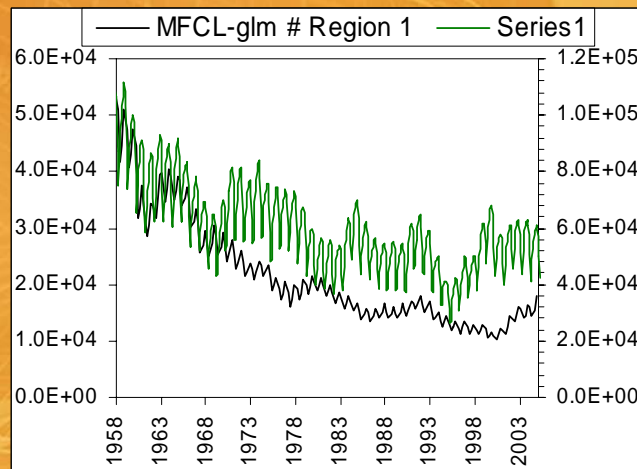
BET



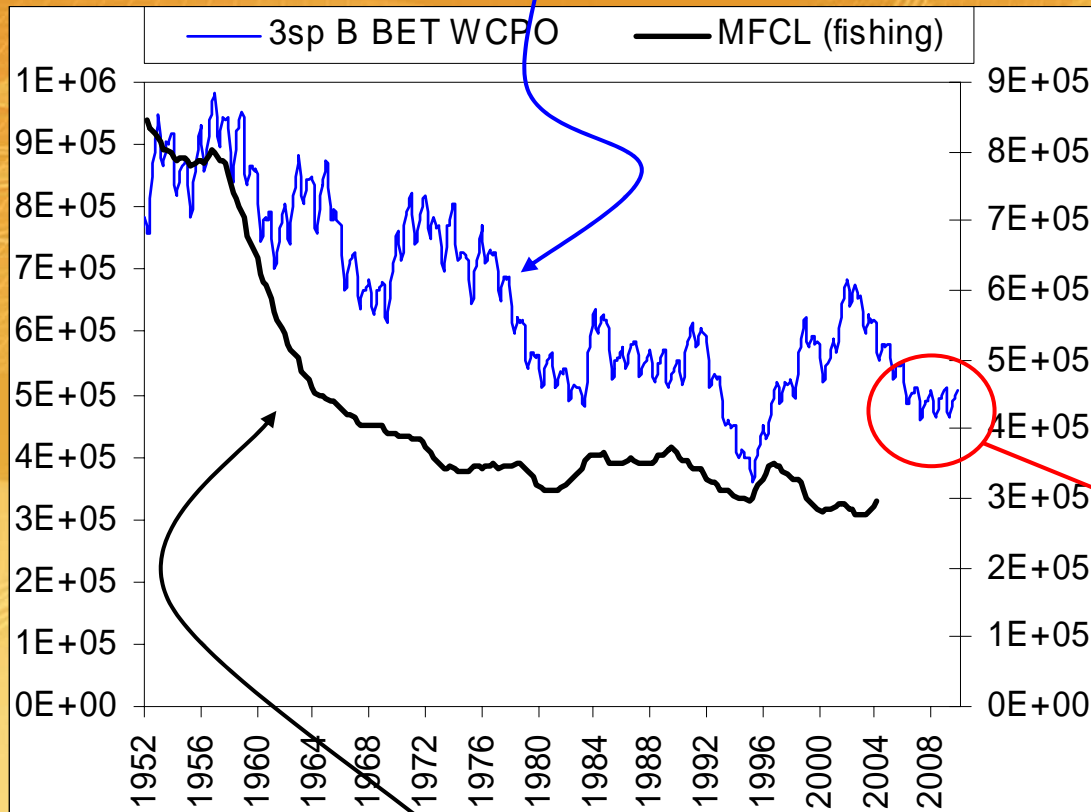
BET
total biomass
Comparison by
region

MFCL (with fisheries):
black curves

Seapodym (3-species,
without fisheries): green
curves



Seapodym total biomass (multi-species simulation – no fishing)



MFCL total biomass with fishing

5-yrs prediction
based on
climatological
environmental data

Conclusions

- In absence of optimization function, a reasonable parameterization for 3 species and their fisheries was obtained.
- The model capture important changes in the population dynamics that explain a large part of time space variability in the catch and CPUE.
- Decline in bigeye stock in the late 1950's and during 1960's is reproduced by the model and due to *natural variability* AND *species interactions*.
- There is no sign of increase in bigeye stock biomass for 5-year projection based on environmental climatology.
- It is now possible to run “what... if” scenarios to test management options in a spatial multi-species and multi-fisheries context.