

# **Les Modèles Hiérarchiques : Un outil pour gérer les emboîtements d'échelles dans la modélisation intégrée des cycles de vie**

**Etienne RIVOT & Félix MASSIOT-GRANIER, Benoît ARCHAMBAULT,  
Sébastien ROCHETTE, Etienne PREVOST, Olivier LE PAPE**



**UMR Ecologie et Santé des Ecosystèmes  
Ecologie Halieutique**



**AFH - 19-21 Juin 2013 - Bordeaux**

# Integrated life cycle models

- We still need population models!
  - Individuals ↔ Populations ↔ Ecosystem
  - The core of decision analysis
  
- Integrated life cycles models allow considering population models in an ecosystemic framework
  
- Raises challenges ... classics of the old dichotomy between mechanistic and statistical models

Levins, R. 1966. The strategy of model building in population biology. *American Scientist*, 54(4), 421–431.

# Integrated life cycle models - Challenges

## Challenge 1 - Understanding demographics

- **Integrate** various demographic processes and factors that control population dynamics at all life stages
  - with their associated sources of variability and uncertainty
  - within a hierarchy of spatial and temporal scales
    - Spatial      Local (~ site specific)  
                  Global (~ large geographical scale)
    - Temporal    Long term (~ decades)  
                  Short term (~ years)

# Integrated life cycle models - Challenges

## Challenge 2 - Data assimilation

- Assimilate various sources of data and information, with their associated sources of uncertainty, to extract an **integrated signal**

- Survey methods are heterogeneous among life stages, in time/space
- Observations errors are the rule, and must be accounted for
- Data are incomplete

Some key demographic transitions can not be directly observed

- Borrowing strength (« Robin Hood Approach », Punt 2011)
- Informative priors (« Standing on the shoulders of giants », Hilborn and Lierman, 1998)

# Hierarchical Bayesian Models

5

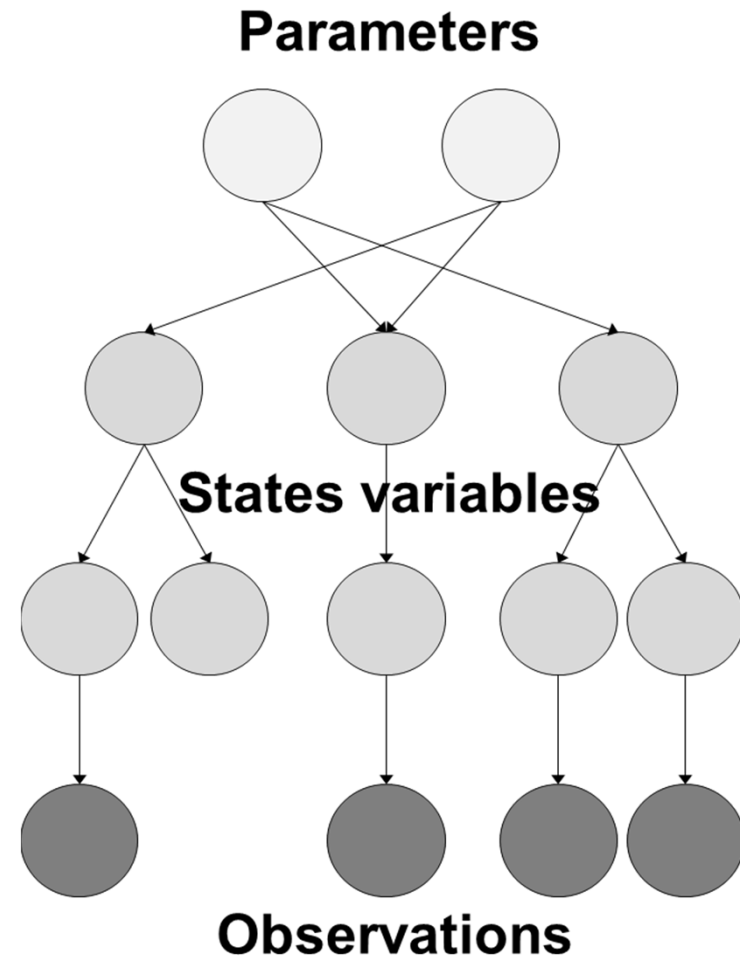
(Clark, 2005 ; Buckland et al., 2007 ; Cressie et al., 2009 ; Kerry and Schaub, 2011 ; Parent and Rivot, 2012)

Conditional probability distributions used for *cause* → *effect* relationships for process and observations

+ (Informative) priors on parameters

- Param. / Demographics / Observations are modeled in a **probabilistic** logic
- Dependencies (Param → States → Obs.) are built in a **mechanistic** logic
- **Synthesis** between mechanistic and probabilistic approaches (~ data assimilation)

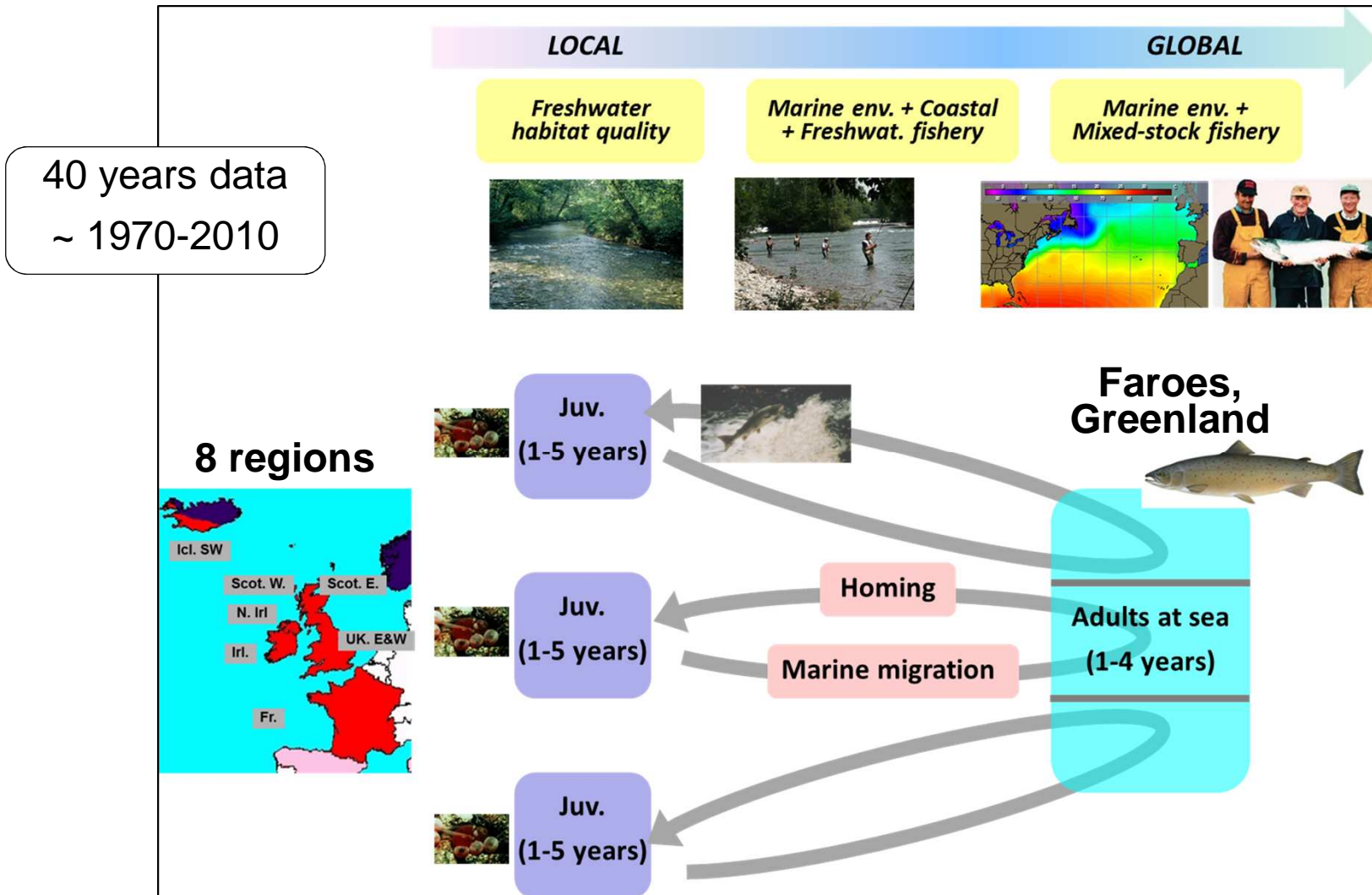
- Allows for building complex network of dependencies, in a hierarchy of time and spatial scales
  1. Demography
  2. Data assimilation (“borrowing strength”)
  3. Informative Priors



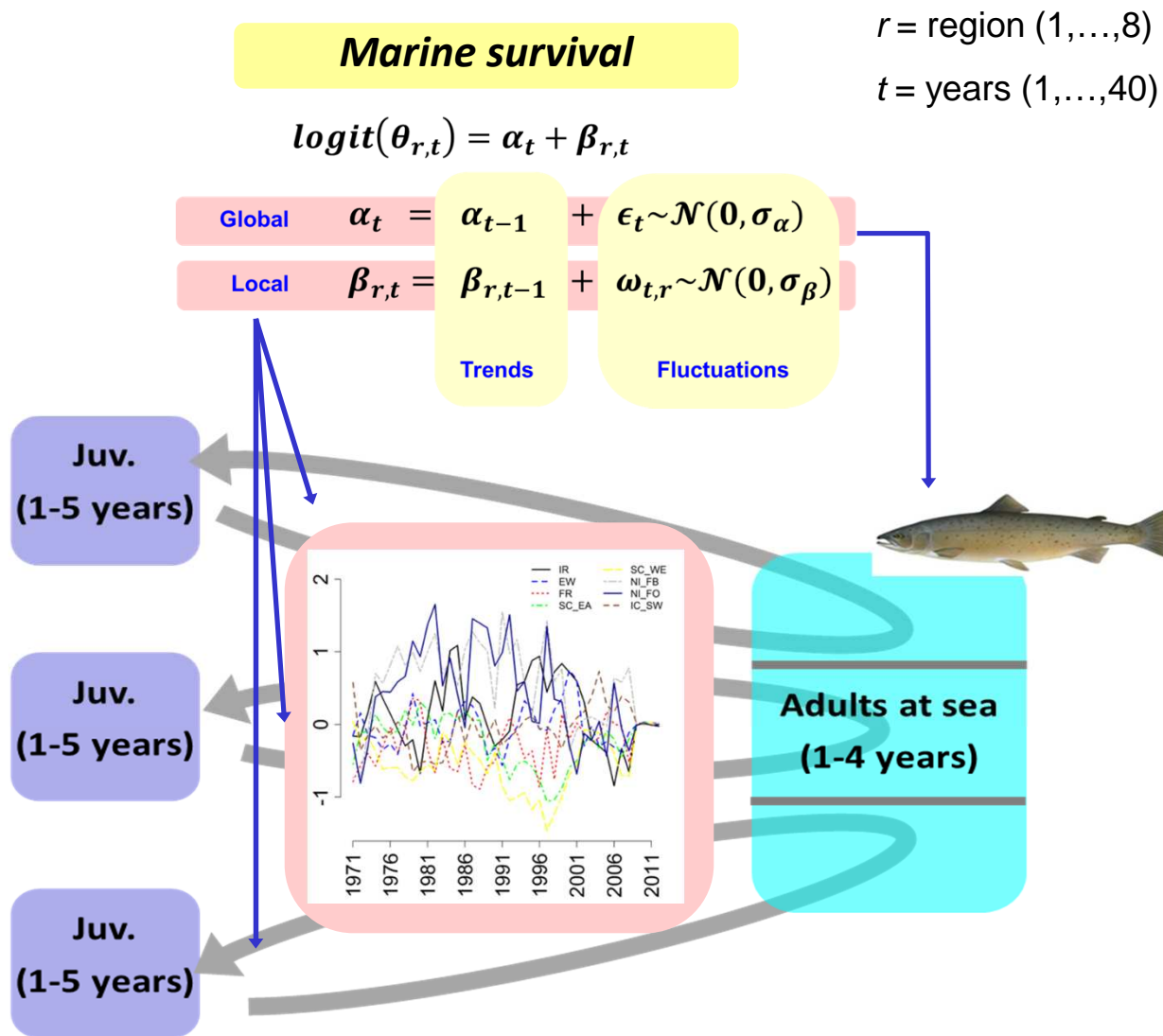
# Atlantic salmon (*Salmo salar*) in the North Atlantic Ocean



F. Massiot-Granier, V. Bret, E. Prévost



# Disentangling the different scales in the A. Salmon freshwater and marine survival



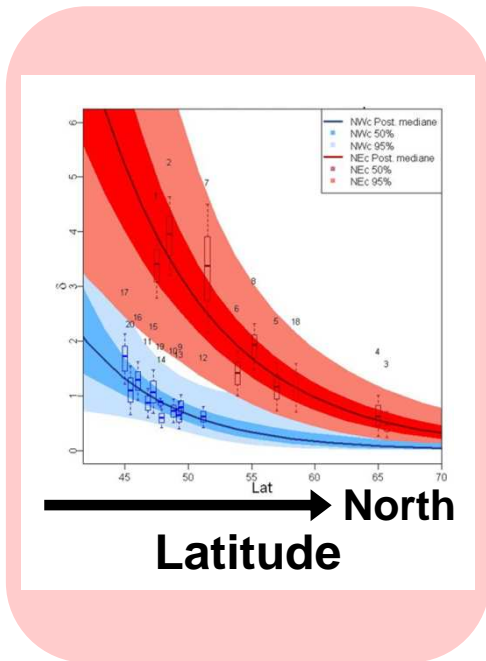
# Disentangling the different scales in the A. Salmon freshwater and marine survival

## Eggs → Smolts survival

21 index rivers

HBM → Borrowing strength  
 "Data rich" → "data poor"

$$Mort_r = e^{\gamma_r + \lambda_r \cdot Latitude_r}$$



## Marine survival

$$logit(\theta_{r,t}) = \alpha_t + \beta_{r,t}$$

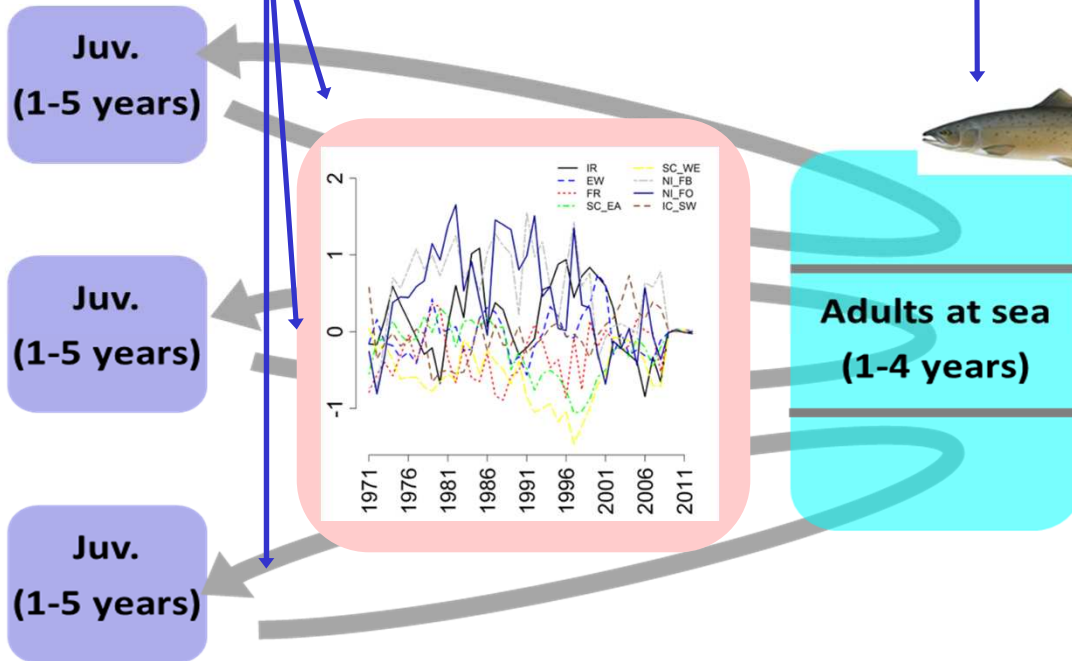
Global  $\alpha_t = \alpha_{t-1} + \epsilon_t \sim \mathcal{N}(0, \sigma_\alpha)$

Local  $\beta_{r,t} = \beta_{r,t-1} + \omega_{t,r} \sim \mathcal{N}(0, \sigma_\beta)$

Trends

Fluctuations

$r = \text{region } (1, \dots, 8)$   
 $t = \text{years } (1, \dots, 40)$

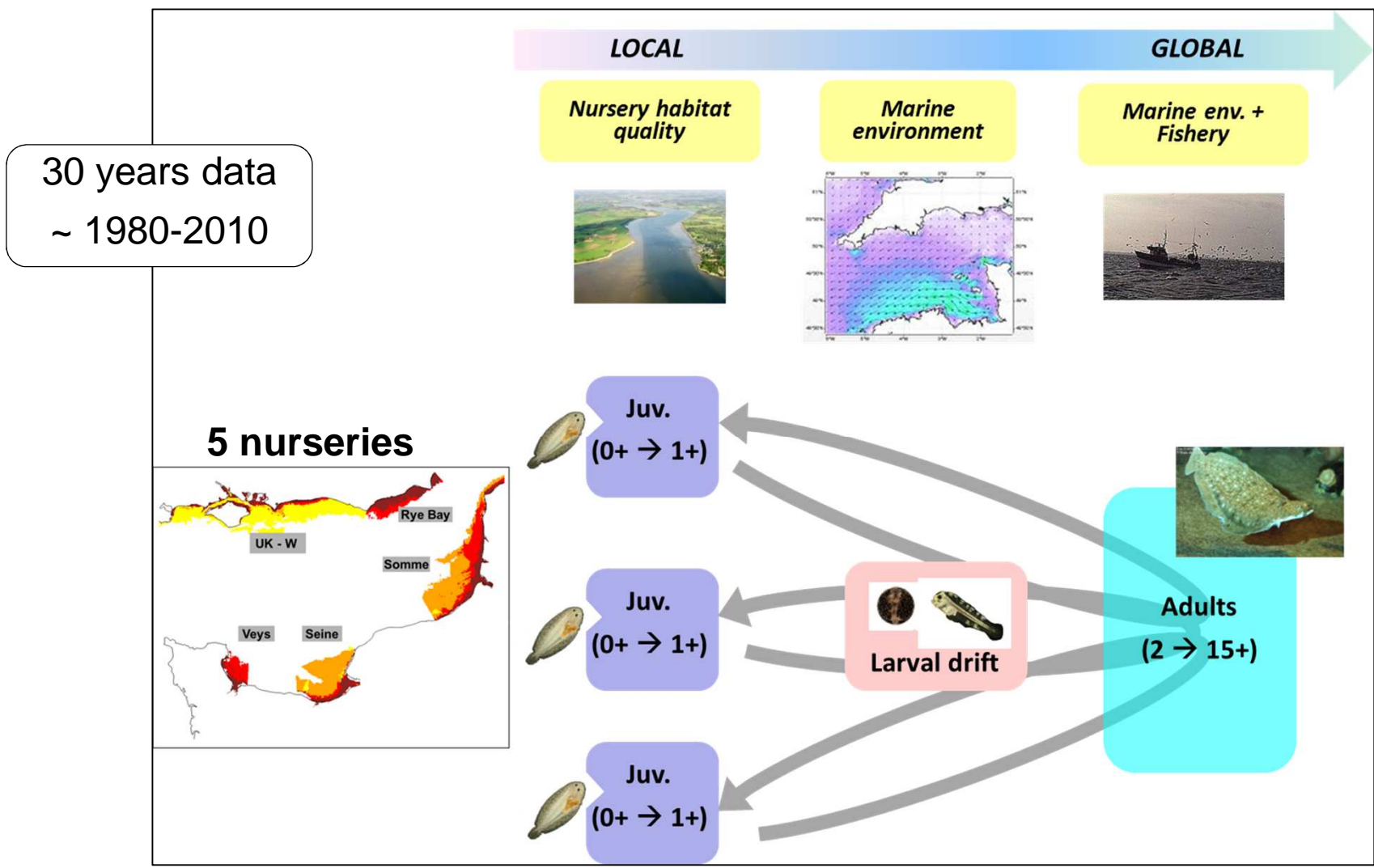




# Common sole (*Solea solea*) in the Eastern Channel



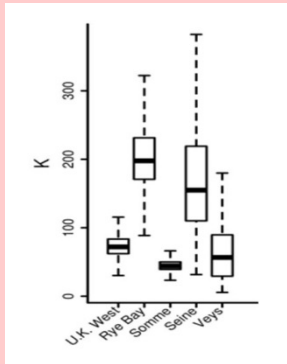
S. Rochette, B. Archambault,  
O. Le Pape, M. Huret



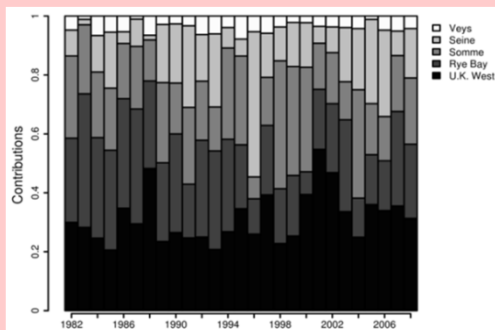
# Merging Lagrangian model for larval drift and survival within a life cycle of the common Sole in the E. Channel

**Nursery-specific density-dependent mortality**

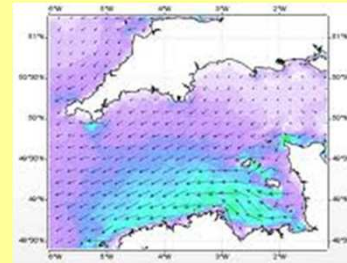
**Nursery-specific carrying capacity**



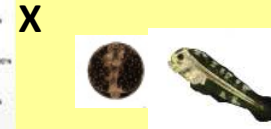
**Contribution of each nursery to the total recruitment**



**Larval drift and survival Lagrangian model (MARS 3D)**



**Biological hyp.**

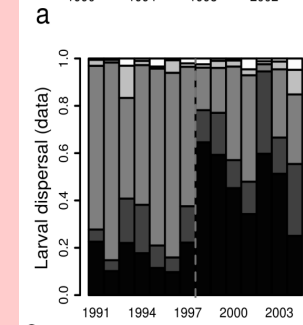
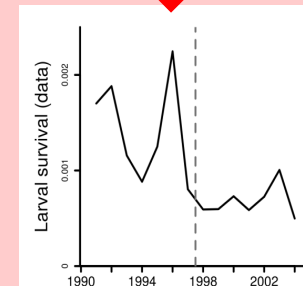


**Informative prior**

Juv. (0+ → 1+)

Juv. (0+ → 1+)

Juv. (0+ → 1+)



**Adults (2 → 15+)**

# Conclusions

- HBMs are useful tools to move from simple population dynamics to integrated life cycle for an ecosystemic approach to population dynamics and demography
- Have the potential for stimulating collaborations and for a synthesis between different fields of science
  - Field ecology (data) and data assimilation
  - Demographics & pop. dynamics
  - Mechanistic models (IBMs ...)
  - Maths. / Statistics / Comput. Sciences
- Provide insights to a better understanding of the factors controlling life cycles in a hierarchy of time and spatial scales

**Thanks for your attention !**



**and welcome in the *Bayesounours World* !**