

Borrowing strength from similar stock-recruitment analyses

HBM for Ecological Data, chapter 9.2

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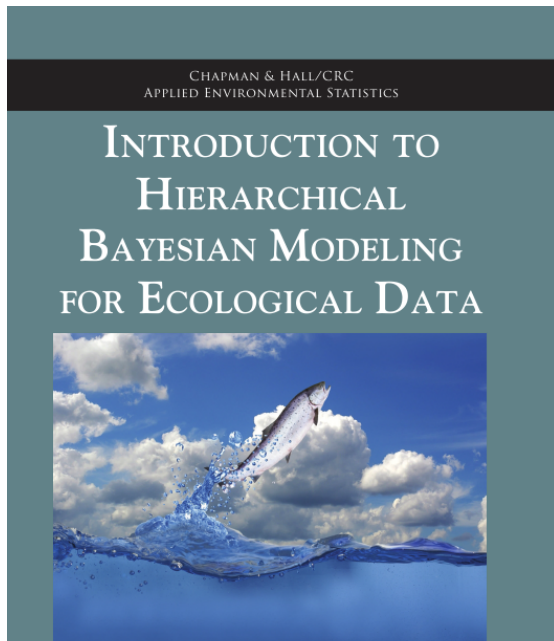
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Hierarchical (*multilevel* or *random effect*) models assume that the dataset being analyzed consists of a *hierarchy* of different *groups* within which records look more alike than between groups.

- SR hierarchical analysis for 13 rivers in Europe : from data-rich to sparse situations.
- Latitude and riverine wetted area accessible to salmon as covariates.
- Hierarchical structure to organize the transfer of information between different units.

In renewable resources management models :

- *Cohort effects* in correlated or familial survival data as opposed to individual behaviors within a group ;
- *Site effects* in meta-analyzes or in *spatially* structured phenomena.

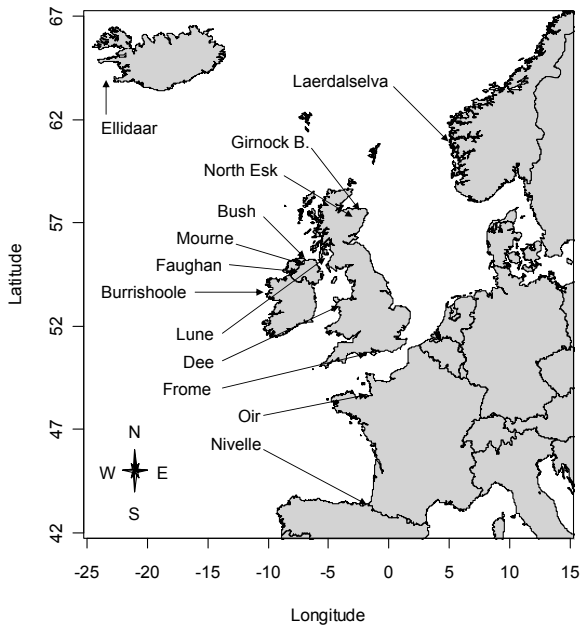


Data for Hierarchical stock-recruitment analysis

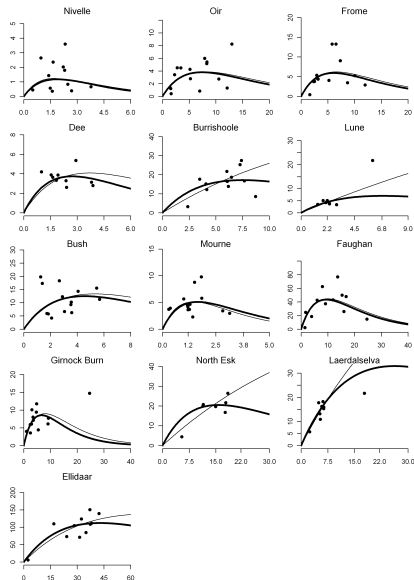
index rivers : a representative sample from the salmon rivers located in western Europe and under the influence of the Gulf Stream.

Biological Reference Points : spawning target S^* , maximum sustainable exploitation rate h^* .

River	Country	Latitude (°N)	Riverine wetted area accessible to salmon (m^2)	Number of SR observations (years)
Nivelle	France	43	320995	12
Oir	France	48.5	48000	14
Frome	England	50.5	876420	12
Dee	England	53	6170000	9
Burrischoole	Ireland	54	155000	12
Lune	England	54.5	4230000	7
Bush	N. Ireland	55	845500	13
Mourne	N. Ireland	55	10360560	13
Faughan	N. Ireland	55	882380	11
Girnock Burn	Scotland	57	58764	12
North Esk	Scotland	57	2100000	6
Laerdalselva	Norway	61	704000	8
Ellidaar	Iceland	64	199711	10



Independent vs hierarchical SR analyses



The Atlantic salmon SR series on the 13 index rivers and fitted SR relationships. S (x -axis) and R (y -axis) are the stock and recruitment variables after standardization for river size expressed in eggs per m^2 of riverine wetted area accessible to salmon. The SR Ricker curves are graphed for two model configurations, the model assuming independence between rivers (thin line) and the hierarchical model (bold line). SR curves are graphed with parameters (S^* , h^*) set at the median of their marginal posterior distributions.

- How is the SR information transferred from the monitored data-rich rivers to set Biological Reference Points for other sparse-data salmon rivers, while accounting for the major sources of uncertainty?
- How can the joint analysis of the SR relationship for the 13 index rivers be used to forecast biological reference points for a new river without any SR data but for which relevant covariates are available?

For river $k = 1, \dots, 13$, one relates the recruitment $R_{k,t}$ rescaled in eggs/m^2 of the cohort born in year t to the associated spawning stock $S_{k,t}$:

$$\begin{cases} \log(R_{k,t}) = h_k^* + \log\left(\frac{S_{k,t}}{1 - h_k^*}\right) - \frac{h_k^*}{S_k^*} S_{k,t} + \varepsilon_{k,t} \\ \varepsilon_{k,t} \stackrel{iid}{\sim} \text{Normal}(0, \sigma_k^2) \end{cases} \quad (1)$$

where σ_k is the standard deviation of the Normal distribution of $\log(R_{k,t})$, S_k^* and h_k^* are respectively the stock which are necessary to guarantee an optimal sustainable exploitation and the associated optimal exploitation rate for the river k .

Poor informative priors, independently for each river k :

$$\begin{cases} h_k^* \sim \text{Beta}(1, 1) \\ S_k^* \sim \text{Uniform}(0, 200) \end{cases}$$

More refined elicitation of S^* : $\mu_{S^*} = 40$ eggs per m^2 with $\sigma_{S^*} = 40$ so that $CV_{S^*} = \sigma_{S^*} / \mu_{S^*} = 1$
Gamma pdf with shape parameter a and scale parameter b such that $\mu_{S^*} = \frac{a}{b}$, $CV_{S^*} = \frac{1}{\sqrt{a}}$ and constrained to the range $]0, 200]$:

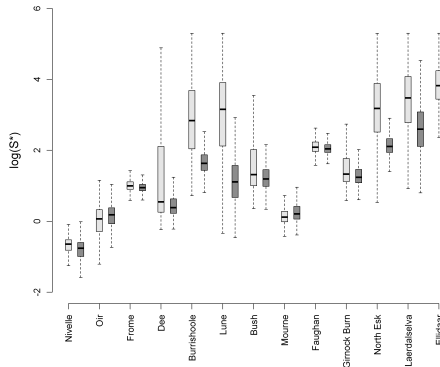
$$\begin{cases} \mu_{S^*} = 40 \text{ eggs}/m^2; CV_{S^*} = 1 \\ a = \frac{1}{CV_{S^*}^2}; b = \frac{1}{\mu_{S^*} \times CV_{S^*}^2} \\ S_k^* \sim \text{Gamma}(a, b) 1_{S_k^* < 200} \end{cases}$$

σ_k^2 constant across all the rivers in the study, and a Gamma prior distribution was assigned on the precision :

$$\begin{cases} \forall k, \sigma_k = \sigma \\ \sigma^{-2} \sim \text{Gamma}(p, q) \end{cases}$$

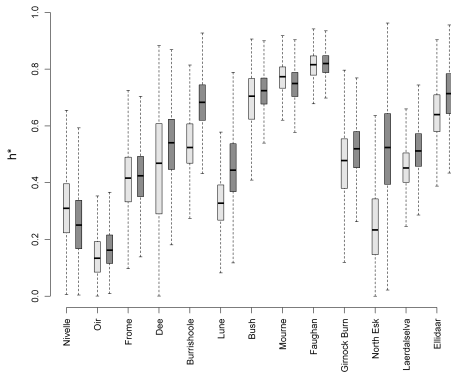
Diffuse $\text{Gamma}(p, q)$ prior for the precision σ^{-2} by letting p and q being very small.

Posterior pdf of Log(Sustainable stock)



Marginal posterior distribution of $\log(S^*)$ (in eggs per m^2) for the 13 index rivers obtained under two model configurations : the model assuming independence between rivers (light gray) and the hierarchical model (dark gray).

Exploitation rate h posterior pdf

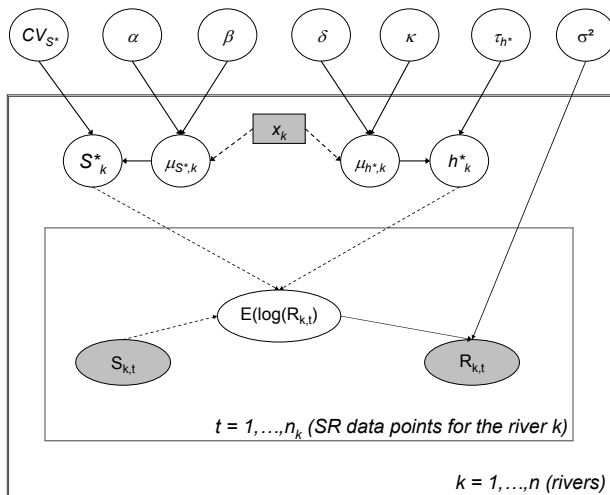


Marginal posterior distribution of h^* for the 13 index rivers obtained under two model configurations : the model assuming independence between rivers (light gray) and the hierarchical model (dark gray).

Posterior inferences on the model assuming independence between rivers

- All distributions exhibit quite heavy tails.
- Depending on the river, the number of observations, and on the contrast between the S values in the observation sample, uncertainty range in the posterior inferences may differ from several orders of magnitude.
- The boxplots of the parameters' posterior pdfs reveal an increasing latitudinal gradient in the S_k^* 's.

Posterior inferences on the hierarchical model with partial exchangeability between rivers (Latitude as a covariate)



Posterior inferences on the hierarchical model with partial exchangeability between rivers (Latitude as a covariate)

S_k is related to latitude of river k denoted x_k :

$$\begin{cases} \log(\mu_{S_k^*}) = \alpha \times x_k + \beta \\ \alpha \sim \text{Uniform}(-5, 5) \\ \beta \sim \text{Uniform}(-50, 50) \end{cases}$$

Parameters a_k and b_k may depend upon the latitude x_k :

$$\begin{cases} CV_{S^*} \sim \text{Uniform}(0, 20) \\ a_k = \frac{1}{CV_{S^*}^2} \\ b_k = \frac{1}{\mu_{S_k^*} \times CV_{S^*}^2} \\ S_k^* \sim \text{Gamma}(a_k, b_k) 1_{S_k^* < 200} \end{cases}$$

Posterior inferences on the hierarchical model with partial exchangeability between rivers (Latitude as a covariate)

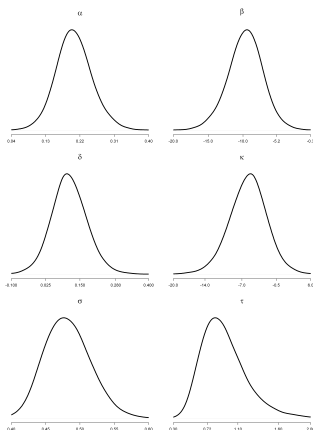
Residual degree of similarity between rivers once the latitude gradient is accounted for :

$$\begin{cases} \text{logit}(\mu_{h_k^*}) = \delta \times x_k + \kappa \\ \delta \sim \text{Uniform}(-5, 5) \\ \kappa \sim \text{Uniform}(-50, 50) \end{cases}$$

Diffuse prior set on precision :

$$\begin{cases} \text{logit}(h_k^*) \sim \text{Normal}(\text{logit}(\mu_{h_k^*}), \tau^2) \\ \tau_{h^*}^{-2} \sim \text{Gamma}(0.001, 0.001) \end{cases}$$

Posterior inferences on the hierarchical model with partial exchangeability between rivers



Marginal posterior probability shapes of the parameters α , β , δ , κ , σ and τ from the hierarchical model.

Posterior inferences on the hierarchical model with partial exchangeability between rivers

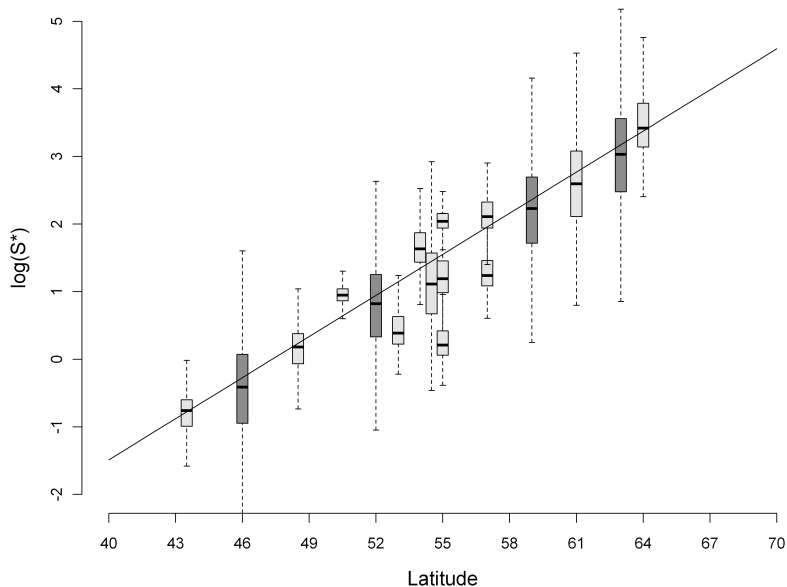
The covariate latitude offers a good statistical explanation of positive variations between rivers in both h^* and S^* . Posterior pdf of S^* and h^* for the monitored 13 rivers reveal :

- Considerable within-river uncertainty in some cases despite SR data being available (*e.g.*, the Lune R. and the Laerdalselva R.) ;
- Significant variations among rivers, even within a relatively narrow latitudinal range (*e.g.*, the Bush R., the Mourne R. and the Faughan R., all located in Northern Ireland) ;
- An increasing trend with latitude.

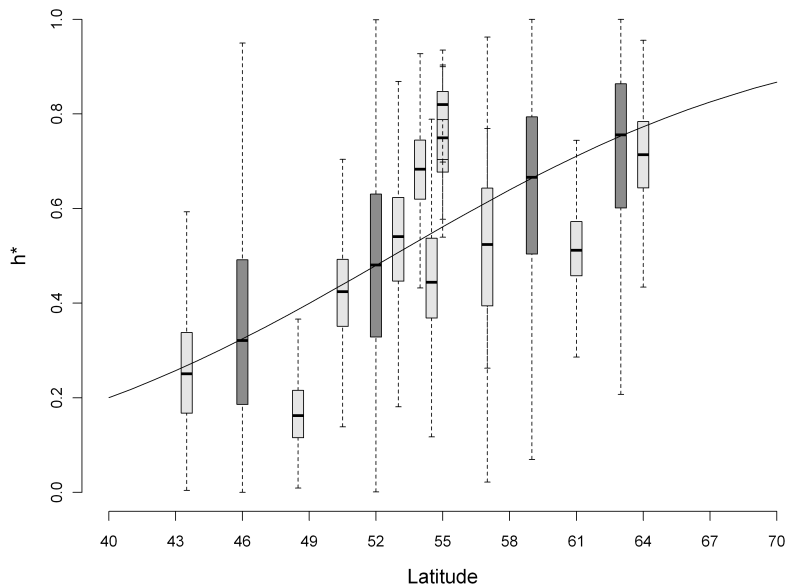
h_{new}^* and S_{new}^* at various latitudes covering the salmon distribution range in the northeast Atlantic area (46° , 52° , 59° and 63° north) .

- Moving north, salmon stocks can sustain higher exploitation rates h^* , can produce higher recruitment at MSY ($R^* = \frac{S^*}{1-h^*}$), but at the same time should be set at higher conservation limits S^* .
- Great remaining uncertainty in the spawning stock, the recruitment and the exploitation rates at MSY for a sparse data river
- important residual variations stem from other explanatory covariates than the riverine wetted area and the latitude.

Posterior inferences on the hierarchical model



Posterior inferences on the hierarchical model





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