

Variable Selection in Additive Models with an Application to Logbook Data on Blue Sharks

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Collaborators & Funding

- Eva Cantoni
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- Important issue in any statistical analysis
- Determine strongest effects that explain the response variable
- Reduces model complexity by admitting a small amount of bias

Motivating Example

- US National Marine Fisheries Service Pelagic Observer Program
- Catches of the blue shark, *Prionace glauca*
- Northeast Coastal and Distant Atlantic

Goals:

- *Statistical:*
 - propose an additive model
 - accommodate covariates which are potentially nonlinearly related to some function of the response (counts)
 - simultaneously fit a model and perform variable selection
- *Ecological:*
 - Are blue shark counts decreasing?

- *Subset selection*: quickly becomes infeasible when the covariate dimension is too large
- *Stepwise procedures*: suffer from dependence on the path chosen through the variable space and may be inconsistent
- *Shrinkage methods*: have emerged and gained popularity in recent years
- **Methods that simultaneously address estimation and variable selection now exist**: modified LASSO, COSSO

Nonnegative Garrote

- Simple approach to variable selection for additive models
- Based on nonnegative garrote idea of Breiman (1995)
- Simultaneously has properties of subset selection, shrinkage and stability
- Computationally reasonable

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Additive Model

$$Y_i = \alpha + \sum_{k=1}^p f_k(\mathbf{x}_{ki}) + \epsilon_i$$

Solves

$$\min_{c_k} \sum_{i=1}^n (y_i - \alpha - \sum_{k=1}^p c_k \hat{g}_k^{h_k}(\mathbf{x}_{ki}))^2$$

under the constraints $c_k \geq 0$ and $\sum_{k=1}^p c_k \leq s$. The final estimate of $f_k(\mathbf{x}_{ki})$ is $\hat{f}_k(\mathbf{x}_{ki}) = c_k \hat{g}_k^{h_k}(\mathbf{x}_{ki})$.

- h_1, \dots, h_p are smoothing parameters of the initial function estimates $\hat{g}_1^{h_1}, \dots, \hat{g}_p^{h_p}$.
- c_k depends on s and s is regarded as an additional parameter.
- Decreasing s has the effect of increasing the shrinkage of the non-zeroed functions and making more of the c_k become zero.
- *Given an initial estimate of all the additive functions in the model and a value for s our method will automatically give a set of coefficients c_1, \dots, c_p that will provide information on the importance of each variable in the model.*

- Smoothing parameters of initial fits must be selected in a reasonable manner
- \Rightarrow We select to use an automatically data driven approach

- Best value of s will be that which minimizes the PE
- \Rightarrow Estimate the PE by V-fold cross-validation

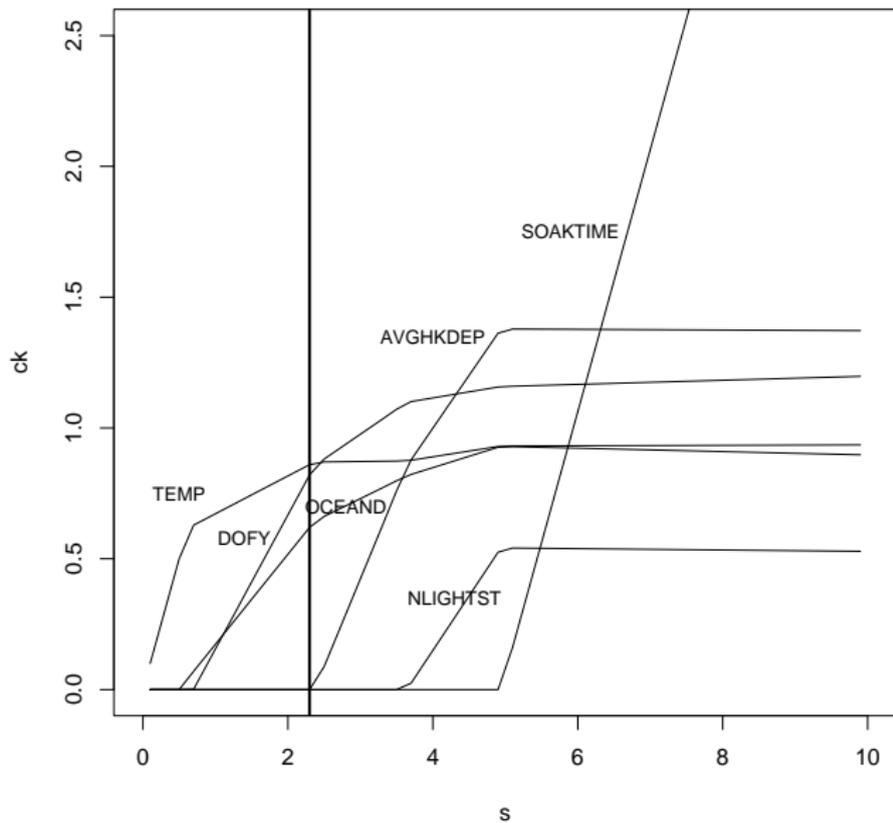
Implementation

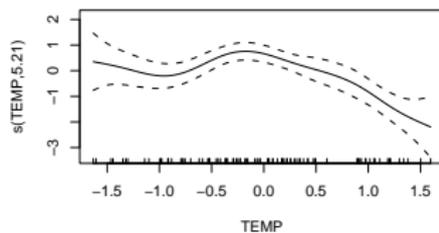
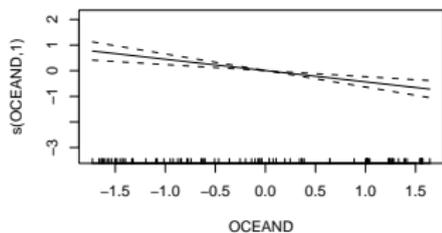
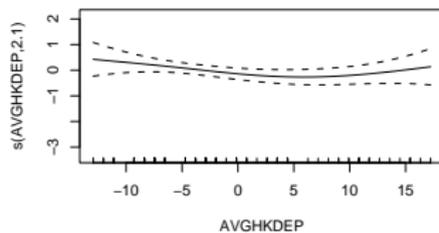
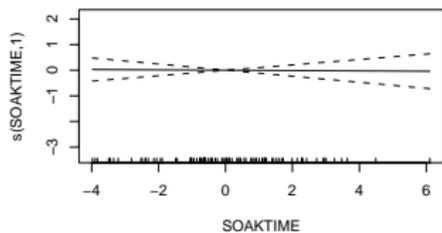
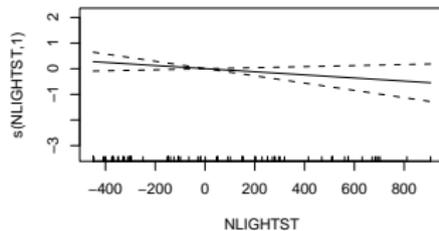
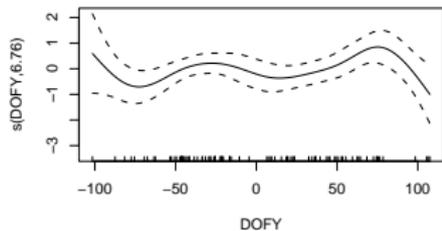
- Two parts:
- \Rightarrow *gam* from the *mgcv* library in R
- \Rightarrow Modified fortran code of Breiman and linked with R

Model

$$\log(\text{bluesharks}+1) = \alpha + f_1(\text{DOFY}) + f_2(\text{NLIGHTST}) + f_3(\text{SOAKTIME}) + f_4(\text{AVGHKDEP}) + f_5(\text{OCEAND}) + f_6(\text{TEMP}) + \log(\text{TOTHOOKS})$$

- Sample size is 91
- Strongest effects are TEMP, OCEAND and DOFY
- SOAKTIME and NLIGHTST can be removed
- AVGHKDEP borderline
- DOFY complicated functional form, TEMP approximately quadratic



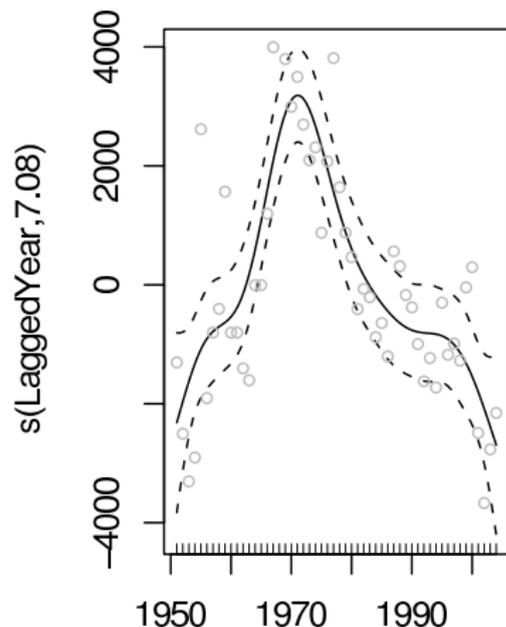


- In terms of predictive ability, as well or better than competitors
- Code readily available and user-friendly

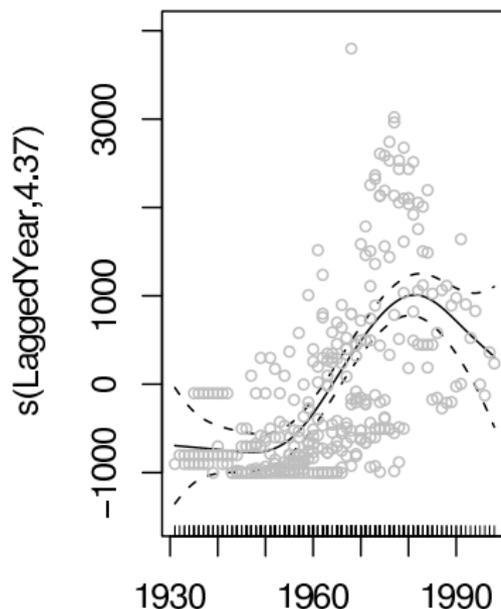
Other Applications

- Patterns of expansion and depletion of invertebrate fisheries on a global scale

Japan



Other countries



- Extension to GAMs
- Robustness aspects