A comparison of three learning methods to predict N₂O fluxes and N leaching



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> June 20*th*, 2010 MASHS 2010, Lille





2 Presentation of three learning methods







Presentation of three learning methods



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- To that aim, a platform has been developed in the EC-project CCAT (Cross-Compliance Assessment Tool) that calculated the impact of cross-compliance legislations on various indicators (GHG, soil pollution, economic indicators, ...);
- But platform needs simplified models which make fast simulations because, in this case, the biogeochemical model DNDC could not be used directly.



Framework of this study

Selection of HSMU related to **corn culture** (about 20 000 spatial units).

Fertilization is done by adding N which produces N_2O (greenhouse gazes) and N leaching (water pollution).

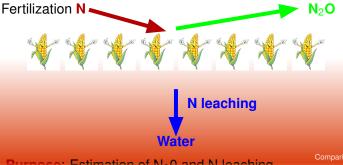




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Purpose: Estimation of N₂0 and N leaching.



5 data sets corresponding to different scenarios coming from the biogeochemical simulator DNDC-EUROPE.

- S1: Baseline scenario (conventional corn cropping system): 18 794 HSMU
- S2: like S1 without tillage: 18 830 HSMU
- S3: like S1 with a limit in N input through manure spreading at 170 kg/ha y: 18 800 HSMU
- S4: rotation between corn (2y) and catch crop (3y): 40 536 HSMU
- S5: like S1 with an additional application of fertilizer in winter: 18 658 HSMU



Data and problem

11 input variables

- N FR (N input through fertilization; kg/ha y);
- N MR (N input through manure spreading; kg/ha y);
- Nfix (N input from biological fixation; kg/ha y);
- Nres (N input from root residue; kg/ha y);
- BD (Bulk Density; g/cm³);
- SOC (Soil organic carbon in topsoil; mass fraction);
- PH (Soil PH);
- Clay (Ratio of soil clay content);
- Rain (Annual precipitation; mm/y);
- Tmean (Annual mean temperature; C);
- Nr (Concentration of N in rain; ppm).



2 outputs to estimate from the inputs:

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Comparison of 3 methods:

- **multi-layer perceptrons** (neural networks): [Bishop, 1995]
- O Support Vector Machines (SVM): [Boser et al., 1992]
- random forests: [Breiman, 2001]





Presentation of three learning methods

3 Methodology and results



Multilayer perceptrons (MLP)

A "one-hidden-layer perceptron" is a function of the form:

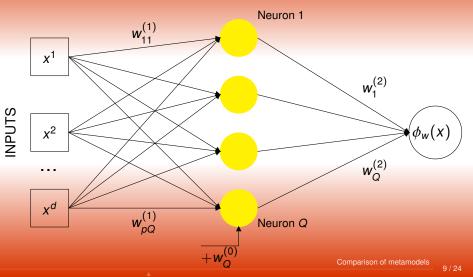
$$\Phi_{w}: x \in \mathbb{R}^{d} \to \sum_{i=1}^{Q} w_{i}^{(2)} G\left(x^{T} w_{i}^{(1)} + w_{i}^{(0)}\right) + w_{0}^{(2)}$$

where:

- the *w* are the **weights** of the MLP that have to be learned from the learning set;
- G is a given activation function: typically, $G(z) = \frac{1-e^{-z}}{1+e^{-z}}$;
- *Q* is the number of neurons on the hidden layer. It controls the flexibility of the machine. *Q* is a **hyper-parameter** that is usually tuned during the learning process.



Symbolic representation of MLP





• Learning the weights: w are learned by a mean squared error minimization scheme :

$$w^* = \arg\min_w \sum_{i=1}^N L(y_i, \Phi_w(x_i)).$$



• Learning the weights: w are learned by a mean squared error minimization scheme penalized by a weight decay to avoid overfitting (ensure a better generalization ability):

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• Tuning of the hyper-parameters, *C* and *Q*: simple validation has been used to tune first *C* and *Q*.



Basic linear SVM for regression: $\Phi_{(w,b)}$ is of the form $x \to w^T x + b$ with (w, b) solution of

$$\arg\min\sum_{i=1}^{N}L_{\epsilon}(y_i,\Phi_{(w,b)}(x_i))+\lambda||w||^2$$

where

- λ is a regularization (hyper) parameter (to be tuned during the learning process);
- L_ϵ(y, ŷ) = max{|y − ŷ| − ϵ, 0} is an ϵ-insensitive loss function
 See ϵ-insensitive loss function



Basic linear SVM for regression

SVM

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 Kernel trick: in fact, φ is never explicit but used through a kernel,

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Common kernel: Gaussian kernel

$$K_{\gamma}(u,v)=e^{-\gamma \|u-v\|^2}$$

is known to have good theoretical properties both for accuracy and generalization.



Learning SVM

• Learning (w, b): $w = \sum_{i=1}^{N} \alpha_i K(x_i, .)$ and *b* are calculated by an exact optimization scheme (quadratic programming). The only step that can be time consumming is the calculation of the kernel matrix:

 $K(x_i, x_j)$ for i, j = 1, ..., N.

12/24



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The resulting $\hat{\Phi}^N$ is known to be of the form:

$$\hat{\Phi}^{N}(x) = \sum_{i=1}^{N} \alpha_{i} K(x_{i}, x) + b$$

where only a few α_i are non zero. The corresponding x_i are called **support vectors**.



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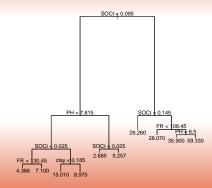
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• Tuning of the hyper-parameters, $C = 1/\lambda$, ϵ and γ : simple validation has been used. To limit waste of time, ϵ has not been tuned in our experiments but set to the default value (1) which ensured 0.5N support vectors at most.



From regression tree to random forest

Example of a regression tree



Each **split** is made such that the two induced subsets have the greatest homogeneity possible.

The **prediction of a final node** is the mean of the Y value of the observations belonging to this node.

Comparison of metamodel



Random forest

Basic principle: combination of a large number of under-efficient regression trees (the prediction is the mean prediction of all trees).



Random forest

Basic principle: combination of a large number of under-efficient regression trees (the prediction is the mean prediction of all trees). For each tree, **two simplifications** of the original method are performed:

- A given number of observations are randomly chosen among the training set: this subset of the training data set is called in-bag sample whereas the other observations are called out-of-bag and are used to control the error of the tree;
- For each node of the tree, a given number of variables are randomly chosen among all possible explanatory variables.

The best split is then calculated on the basis of these variables and of the chosen observations. The chosen observations are the same for a given tree whereas the variables taken into account change for each split.



Learning a random forest

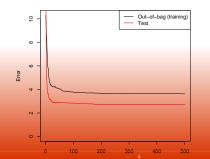
Random forest are **not very sensitive to hyper-parameters** (number of observations for each tree, number of variables for each split): the default values have been used.



Learning a random forest

Random forest are **not very sensitive to hyper-parameters** (number of observations for each tree, number of variables for each split): the default values have been used.

The number of trees should be large enough for the mean squared error based on out-of-sample observations to stabilize:





Methodology and results



Data and problem

Presentation of three learning methods









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The data set has been split into a training set and a test set (on a 80%/20% basis);





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- O The machine has been learned from the training set (with a full validation process for the hyperparameter tuning);
- The performances have been calculated on the basis of the test set: for the test set, predictions have been made from the inputs and compared to the true outputs.



Numerical results

		MLP	RF	SVM	LM
N ₂ O	Scenario 1	90.7%	92.3%	91.0%	77.1%
	Scenario 2	80.3%	85.0%	82.3%	62.2%
	Scenario 3	85.1%	88.0%	86.7%	74.6%
	Scenario 4	88.6%	90.5%	86.3%	78.3%
	Scenario 5	80.6%	84.9%	82.7%	42.5%
N leaching	Scenario 1	89.7%	93.5%	96.6%	70.3%
	Scenario 2	91.4%	93.1%	97.0%	67.7%
	Scenario 3	90.6%	90.4%	96.0%	68.6%
	Scenario 4	80.5%	86.9%	90.6%	42.5%
	Scenario 5	86.5%	92.9%	94.5%	71.1%

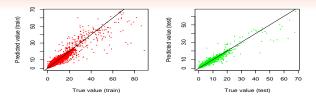
Comparison of metamodels

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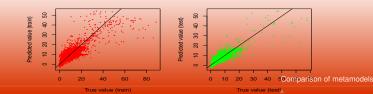


Graphical results

Random forest (S1, N₂O)



Linear model (S1, N₂O)





Further comparisons

	Time for training	Time for using once the model is trained		
RF	+ (~ 15 minutes)	= (~ 5 seconds)		
SVM	=	- (~ 20 seconds)		
		sensitive to N		
MLP	-	+ (~ 1 second)		
		sensitive to d		

Times correspond to the prediction for about 20 000 HSMU on a Dell Latitude D830, Intel Core 2DUO 2.2GHz, 2GO RAM, OS Ubuntu Linux 9.10 (and, for RF, to the training of about 15 000 HSMU) with R.

Time for DNDC: around 200 hours by using a desktop computer and around 2 days by using a cluster of computers.



Understanding which inputs are important

Importance: A measure to estimate the importance of the input variables can be defined by:

- for a given input variable randomly permute the input values and calculate the prediction from this new randomly permutated inputs;
- compare the accuracy of these predictions to accuracy of the predictions obtained with the true inputs: the increase of mean squared error is called the importance.



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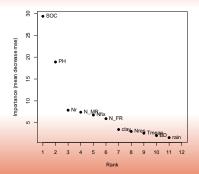
This comparison is made on the basis of data that are not used to define the machine, either the validation set or the out-of-bag observations.



Methodology and results

Understanding which inputs are impor-

Example (S1, N₂O, RF):



The variables SOC and PH are the most important for accurate predictions.

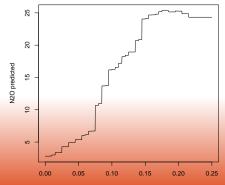
21 / 24



Methodology and results

Understanding the relation between a given input and the output

Evolution of N₂O in function of the value of SOC



Comparison of metamode



Conclusion and further work

- Learning approaches can provide accurate estimations of the variable of interest much faster than the biogeochemical simulator;
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- Learning approaches can provide accurate estimations of the variable of interest much faster than the biogeochemical simulator;
- The random forest approach is now implemented in DNDC to allow fast test of policies impacts.
- Other statistical nonparametric approaches (based on splines) are currently under test.



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> Thank you for your attention. Questions? Comments?

> > Comparison of metamodel





$\epsilon\text{-insensitive loss function}$

