

# UiO: University of Oslo

Outline of the lecture

# Generalized Additive Models

- Definition
- Fitting algorithm
- Tree-based Methods
  - Background
  - How to grow a regression tree
- Bagging
  - Bootstrap aggregation
  - Bootstrap trees

STK-IN4300 Statistical Learning Methods in Data Science

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Generalized Additive Models: introduction

From the previous lecture:

- linear regression model are easy and effective models;
- often the effect of a predictor on the response is not linear;

local polynomials and splines.

# **Generalized Additive Models:**

- flexible statistical methods to identify and characterize nonlinear regression effects;
- lerger class than the generalized linear models.

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Generalized Additive Models: additive models

Consider the usual framework:

- ullet  $X_1,\ldots,X_p$  are the predictors;
- Y is the response variable;
- ullet  $f_1(\cdot),\ldots,f_p(\cdot)$  are unspecified smooth functions.

Then, an  $\underline{\text{additive model}}$  has the form

$$E[Y|X_1,...,X_p] = \alpha + f_1(X_1) + \cdots + f_p(X_p).$$

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#### Generalized Additive Models: more generally

As linear models are extended to generalized linear models, we can generalized the additive models to the generalized additive models,

$$g(\mu(X_1,\ldots,X_p)) = \alpha + f_1(X_1) + \cdots + f_p(X_p),$$

where:

- $\mu(X_1,\ldots,X_p)=E[Y|X_1,\ldots,X_p]$  is the link function;
- $g(\mu(X_1,\ldots,X_p))$  is the link function;
- classical examples:
  - $g(\mu) = \mu \leftrightarrow \text{identity link} \rightarrow \text{Gaussian models};$
  - $g(\mu) = \log(\mu/(1-\mu)) \leftrightarrow \text{logit link} \rightarrow \text{Binomial models};$
  - $g(\mu) = \Phi^{-1}(\mu) \leftrightarrow \text{probit link} \rightarrow \text{Binomial models};$
  - $g(\mu) = \log(\mu) \leftrightarrow \text{logarithmic link} \rightarrow \text{Poisson models};$

· ...

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Fitting algorithm: difference with splines

When implementing splines:

- each function is modelled by a basis expansion;
- the resulting model can be fitted with least squares.

Here the approach is different:

- each function is modelled with a smoother (smoothing splines, kernel smoothers, . . . )
- ullet all p functions are simultaneously fitted via an algorithm.



Generalized Additive Models: semiparametric models

Generalized additive models are very flexible:

• not all functions  $f_i(\cdot)$  must be nonlinear;

$$g(\mu) = X^T \beta + f(Z)$$

in which case we talk about semiparametric models.

nonlinear effect can be combined with qualitative inputs,

$$g(\mu) = f(X) + g_k(Z) = f(X) + g(V, Z)$$

where k indexes the level of a qualitative variable V.

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Fitting algorithm: ingredients

Consider an additive model

$$Y = \alpha + \sum_{j=1}^{p} f_j(X_j) + \epsilon.$$

We can define a loss function,

$$\sum_{i=1}^{N} \left( y_i - \alpha - \sum_{j=1}^{p} f_j(x_{ij}) \right)^2 + \sum_{j=1}^{p} \lambda_j \int \{f_j''(t_j)\}^2 dt_j$$

- $\lambda_i$  are tuning parameters;
- the minimizer is an additive cubic spline model,
  - each  $f_i(X_i)$  is a cubic spline with knots at the (unique)  $x_{ij}$ 's.



Fitting algorithm: constrains

The parameter  $\alpha$  is in general not identifiable:

- same result if adding a constant to each  $f_j(X_j)$  and subtracting it from  $\alpha$ ;
- by convention,  $\sum_{j=1}^{p} f_j(X_j) = 0$ :
  - the functions average 0 over the data;
  - $\bullet$   $\alpha$  is therefore identifiable;
  - in particular,  $\hat{\alpha} = \bar{y}$ .

If this is true and the matrix of inputs X has full rank:

- the loss function is convex;
- the minimizer is unique.

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Fitting algorithm: remarks

#### Note:

- the smoother S can be (when applied only at the training points) represented by the  $N \times N$  smoothing matrix S,
  - the degrees of freedom for the j-th terms are trace(S);
- for the generalized additive model, the loss function is the penalized log-likelihood;
- the backfitting algorithm fits all predictors,
  - not feasible when p >> N.



Fitting algorithm: backfitting algorithm

# The backfitting algorithm:

- 1. Initialization:  $\hat{\alpha} = N^{-1} \sum_{i=1}^{N} y_i$  and  $\hat{f}_j \equiv 0 \ \forall j$
- 2. In cycle, j = 1, ..., p, 1, ..., p, ...

$$\hat{f}_j \leftarrow \mathcal{S}_j \left[ \{ y_i - \alpha - \sum_{k \neq j} \hat{f}_k(x_{ik}) \}_1^N \right]$$

$$\hat{f}_j \leftarrow \hat{f}_j - \frac{1}{N} \sum_{i=1}^N \hat{f}_j(x_{ij}) \}$$

until  $\hat{f}_i$  change less than a pre-specified threshold.

 $S_j$  is usually a cubic smoothing spline, but other smoothing operators can be used.

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#### Tree-based Methods: introduction

Consider a regression problem, Y the response, X the input matrix.

A tree is a recursive binary partition of the feature space:

- each time a region is divide in two or more regions;
  - until a stopping criterion applies;
- at the end, the input space is split in M regions  $R_m$ ;
- a constant  $c_m$  is fitted to each  $R_m$ .

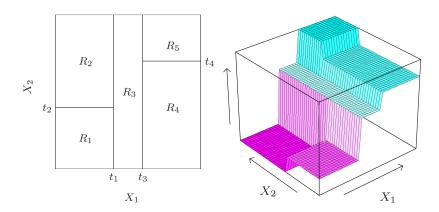
The final prediction is

$$\hat{f}(X) = \sum_{m=1}^{M} \hat{c}_m \mathbb{1}(X \in R_m),$$

where  $\hat{c}_m$  is an estimate for the region  $R_m$  (e.g., ave $(y_i|x_i \in R_m)$ ).

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# Tree-based Methods: introduction

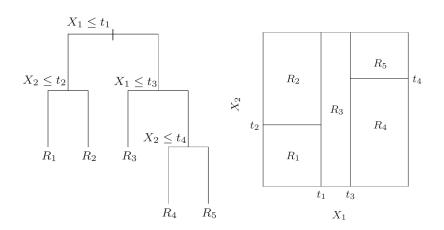


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#### Tree-based Methods: introduction





Tree-based Methods: introduction

#### Note:

- the split can be represented as a junction of a tree;
- this representation works for p > 2;
- each observation is assigned to a branch at each junction;
- the model is easy to interpret.

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#### How to grow a regression tree: split

How to grow a regression tree:

- we need to automatically decide the splitting variables . . .
- ...and the splitting points;
- we need to decide the shape (topology) of the tree.

Using a sum of squares criterion,  $\sum_{i=1}^{N}(y_i-f(x_i))^2$ ,

- the best  $\hat{c}_m = \operatorname{ave}(y_i|x_i \in R_m)$ ;
- finding the best partition in terms of minimum sum of squares is generally computationally infeasible

↓ go greedy

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#### How to grow a regression tree: greedy algorithm

# Starting with all data:

- for each  $X_i$ , find the best split point s
  - define the two half-hyperplanes,
    - $R_1(j,s) = \{X | X_j \le s\};$
    - $R_2(j,s) = \{X|X_j > s\};$
  - the choice of s can be done really quickly;
- for each j and s, solve

$$\min_{j, s} \left[ \min_{c_1} \sum_{x_i \in R_1(j, s)} (y_i - c_1)^2 + \min_{c_2} \sum_{x_i \in R_2(j, s)} (y_i - c_2)^2 \right]$$

- the inner minimization is solved by
  - $\hat{c}_1 = \operatorname{ave}(y_i | x_i \in R_1(j, s));$
  - $\hat{c}_2 = \operatorname{ave}(y_i | x_i \in R_2(j, s)).$
- the identification of the best (j, s) is feasible.

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### How to grow a regression tree: cost-complexity pruning

Consider a tree  $T \subset T_0$ , computed by pruning  $T_0$  and define:

- $R_m$  the region defined by the node m;
- |T| the number of terminal nodes in T;
- $N_m$  the number of observations in  $R_m$ ,  $N_m = \#\{x_i \in R_m\}$ ;
- $\hat{c}_m$  the estimate in  $R_m$ ,  $\hat{c}_m = N_m^{-1} \sum_{r:\in R_m} y_i$ ;
- $Q_m(T)$  the loss in  $R_m$ ,  $Q_m(T) = N_m^{-1} \sum_{x_i \in R_m} (y_i \hat{c}_m)^2$ .

# Then, the **cost complexity criterion** is

$$C_{\alpha}(T) = \sum_{m=1}^{|T|} N_m Q_m(T) + \alpha |T|.$$



#### How to grow a regression tree: when to stop

#### The tree size:

- is a tuning parameter;
- it controls the model complexity;
- its optimal values should be chosen from the data.

# Naive approach:

- split the tree nodes only if there is a <u>sufficient decrease</u> in the sum-of-squares (e.g., larger than a pre-specified threshold);
  - intuitive:
  - short-sighted (a split can be preparatory for a split below).

# Preferred strategy:

- grow a large (pre-specified # of nodes) or complete tree  $T_0$ ;
- prune (remove branches) it to find the best tree.

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### How to grow a regression tree: cost-complexity pruning

The idea is to find the subtree  $T_{\hat{\alpha}} \subset T_0$  which minimizes  $C_{\alpha}(T)$ :

- $\forall \alpha$ , find the unique subtree  $T_{\alpha}$  which minimizes  $C_{\alpha}(T)$ ;
- through weakest link pruning:
  - successively collapse the internal node that produce the smallest increase in  $\sum_{m=1}^{|T|} N_m Q_m(T)$ ;
  - until the single node tree;
  - find  $T_{\alpha}$  within the sequence;
- find α̂ via cross-validation.

# Here the tuning parameter $\alpha$ :

- governs the trade-off between tree size and goodness of fit;
- ullet larger values of lpha correspond to smaller trees;
- α = 0 → full tree.

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#### Classification trees: definition

No major differences between regression and classification trees:

• define a class  $k \in \{1, \dots, K\}$  for each region,

$$k_m = \operatorname{argmax}_k \, \hat{p}_{mk} = \operatorname{argmax}_k \, \left\{ N_m^{-1} \sum_{x_i \in R_m} \mathbb{1}(y_i = k) \right\};$$

- change the loss function from  $Q_m(T)$  to:
  - 0-1 loss:  $N_m^{-1} \sum_{x_i \in R_m} \mathbb{1}(y_i \neq k_m);$  Gini index:  $\sum_{k=1}^K \hat{p}_{mk} (1 \hat{p}_{mk});$  deviance:  $\sum_{k=1}^K \hat{p}_{mk} \log \hat{p}_{mk};$

  - ▶ all three can be extended to consider different error weights.

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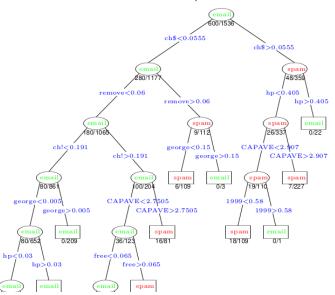
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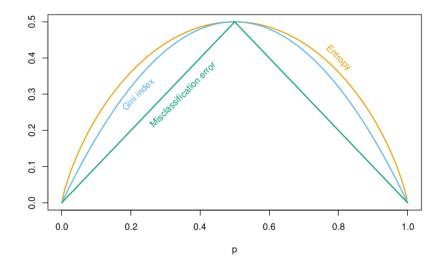
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# Classification trees: example



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#### Classification trees: loss functions



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Tree-based Methods: remarks

#### Tree-based methods:

- fast to construct, interpretable models;
- can incorporate mixtures of numeric and categorical inputs;
- immune to outliers, resistant to irrelevant inputs;
- lack of smoothness;
- difficulty in capturing additive structures;
- highly unstable (high variance).

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#### Bagging: Galton (1907)

NATURE 450

[MARCH 7, 1907

carion to those portroins of southern sew macket or war an altitude less than 4000 feb. William (1960) and the sex of the

tion is similar in all respects to its predecessors; it contains most valuable data relating to the North Sea and

450		ONL		[WIARCH 7, 1907		
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We have also received the official meteorologic books for South Australia (1904) and Mysore (1905) of these works contain valuable means for previous Forty Years of Southern New Mexico Climate.	s). Both	the length of Array o -100°	Estimates in lbs.	Observed deviates from 1207 lbs.	Normal p.e =37	Excess of Observed ove Normal
No. 59 of the New Mexico College of Agriculture the meteorological data recorded at the experimenta	contains	°5	1074	- 133	- 90	+43
from 1892 to 1905 inclusive, together with re	esults of	10	1109	- 98	- 70	+ 28
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'Report on the Climate of New Mexico." The s	station is	30	1174	- 33	- 29	+ 4
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(fourteen years) 76° 8, mean minimum 41° 4,		60	1219	+ 12	+14	- 2
maximum 106° (which occurred several times),		65	1225	+ 18	+21	- 3
minimum 1° (December, 1895). The mean annu	ual rain-	70	1230	+ 23	+ 29	- 6
fall was 8.8 inches; the smallest yearly amo	unt was	93 75	1236	+ 29	+ 37	- 8
3-5 inches, in 1873, the largest 17-1 inches, in 190		80	1243	+ 36	+46	- 10
of the rain falls during July, August, and Se		85	1254	+ 47	+ 57	~ 10
The relative humidity is low, the mean annual amou	unt being	90	1267	+ 52	+70	- 18
about 51 per cent. The bulletin was prepared by	by J. D.	95	1293	+ 86	+90	- 4

 $q_1, q_2$ , the first and third quartiles, stand at 25° and 75° respectively. m, the median or middlemost value, stands at  $g_0$ °. The dressed weight proved to be 1:198 lbs.

According to the democratic principle of "one vote one value," the middlemost estimate expresses the vox populi, every other estimate being condemned as too low or too high by a majority of the voters (for fuller explanation see "One Vote, One Value," NATURE, February 28, 414). Now the middlemost estimate is 1207 lb., if the weight of the dressed ox proved to be 1198 lb.;

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the rainy days being 172 in number

VOX POPULI

I N these democratic days, any investigation into the trustworthiness and peculiarities of popular judgments

was of the usual type, so far that they closely in its neighbourhood and became more sparse as the distance from it is Diagram, from the tabular values.



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# Bagging: Galton (1907)

After having arrayed and analyzed the data, Galton (1907) stated:

It appears then, in this particular instance, that the vox populi is correct to within 1 per cent of the real value, and that the individual estimates are abnormally distributed in such a way that it is an equal chance whether one of them, selected at random, falls within or without the limits of -3.7 per cent and +2.4 per cent of their middlemost value.

Concept of "Wisdom of Crowds" (or, as Schapire & Freund. 2014, "how it is that a committee of blockheads can somehow arrive at a highly reasoned decision, despite the weak judgement of the individual members.")



Bagging: Galton (1907)

In 1907, Sir Francis Galton visited a country fair:

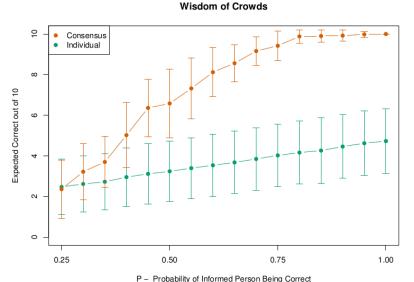
A weight-judging competition was carried on at the annual show of the West of England Fat Stock and Poultry Exhibition recently held at Plymouth. A fat ox having been selected, competitors bought stamped and numbered cards [...] on which to inscribe their respective names, addresses, and estimates of what the ox would weigh after it had been slaughtered and "dressed". Those who guessed most successfully received prizes. About 800 tickets were issued, which were kindly lent me for examination after they had fulfilled their immediate purpose.

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# Bagging: wisdom of crowds



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# Bagging: translate this message into trees

How do can we translate this idea into tree-based methods?

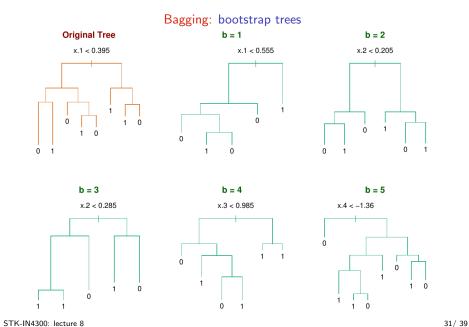
- we can fit several trees, then aggregate their results;
- problems:
  - "individuals" are supposed to be independent;
  - we have only one dataset . . .

How can we mimic different datasets while having only one?

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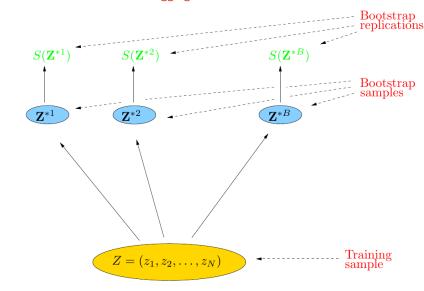


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# Bagging: the solution is . . .

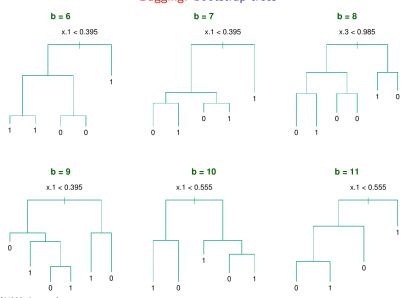


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#### Bagging: bootstrap trees

The procedure so far:

- generate bootstrap samples;
- fit a tree on each bootstrap sample;
- obtain B trees.

At this point, aggregate the results. How?

- consensus:  $\hat{G}(x) = \operatorname{argmax}_k q_k(x), k \in \{1, \dots, K\},$ 
  - where  $q_k(x)$  is the proportion of trees voting for the category k;
- probability:  $\hat{G}(x) = \operatorname{argmax}_k B^{-1} \sum_{b=1}^B p_k^{[b]}(x)$ ,  $k \in \{1, \dots, K\}$ ,
  - where  $p_k^{[b]}(x)$  is the probability assigned by the b-th tree to category k;

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# Bagging: general

In general, consider the training data  $Z = \{(y_1, x_1), \dots, (y_N, x_N)\}$ . The **bagging** (boostrap **agg**regating) estimate is define by

$$\hat{f}_{\mathsf{bag}}(x) = E_{\hat{\mathcal{D}}}[\hat{f}^*(x)],$$

where:

- $\hat{\mathcal{P}}$  is the empirical distribution of the data  $(y_i, x_i)$ ;
- $\hat{f}^*(x)$  is the prediction computed on a bootstrap sample  $Z^*$ ;
- i.e.,  $(y_i^*, x_i^*) \sim \hat{\mathcal{P}}$ .

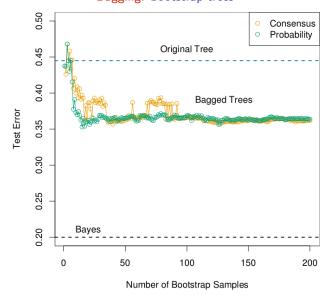
The empirical version of the bagging estimate is

$$\hat{f}_{\text{bag}}(x) = \frac{1}{B} \sum_{b=1}^{B} \hat{f}^*(x),$$

where B is the number of bootstrap samples.



#### Bagging: bootstrap trees



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Bagging: variance

Bagging has smaller prediction error because it reduces the variance component,

$$\begin{split} E_{\mathcal{P}}[(Y - \hat{f}^*(x))^2] &= E_{\mathcal{P}}[(Y - f_{\mathsf{bag}}(x) + f_{\mathsf{bag}}(x) - \hat{f}^*(x))^2] \\ &= E_{\mathcal{P}}[(Y - f_{\mathsf{bag}}(x))^2] + E_{\mathcal{P}}[(f_{\mathsf{bag}}(x) - \hat{f}^*(x))^2] \\ &\geqslant E_{\mathcal{P}}[(Y - f_{\mathsf{bag}}(x))^2], \end{split}$$

where  $\mathcal{P}$  is the data distribution.

Note that this does not work for 0-1 loss:

- due to non-additivity of bias and variance;
- bagging makes better a good classifier, worse a bad one.

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# Bagging: from bagging to random forests

The average of B identically distributed r.v. with variance  $\sigma^2$  and positive pairwise correlation  $\rho$  has variance

$$\rho\sigma^2 + \frac{1-\rho}{B}\sigma^2.$$

- as B increases, the second term goes to 0;
- the bootstrap trees are p. correlated → first term dominates.

construct bootstrap tree as less correlated as possible

# random forests

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GALTON, F. (1907). Vox populi. Nature 75, 450-451.

Schapire, R. E. & Freund, Y. (2014). *Boosting: Foundations and Algorithms.* MIT Press, Cambridge.

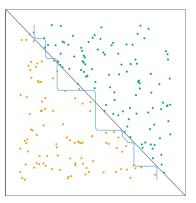
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# Bagging: from bagging to boosting

#### **Bagged Decision Rule**

#### **Boosted Decision Rule**



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