

A walk in random forests

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Background on random forests

Random forests are a class of algorithms used to solve regression and classification problems

- They are often used in applied fields since they handle high-dimensional settings.
- They have good predictive power and can outperform state-of-the-art methods.



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But mathematical properties of random forests remain a bit magical.

1 Construction of random forests

2 Centred Forests

3 Median forests

4 Consistency of Breiman forests

General framework of the presentation

Regression setting

We are given a training set $\mathcal{D}_n = \{(X_1, Y_1), \dots, (X_n, Y_n)\}$ where the pairs $(X_i, Y_i) \in [0, 1]^d \times \mathbb{R}$ are *i.i.d.* distributed as (X, Y) .

We assume that

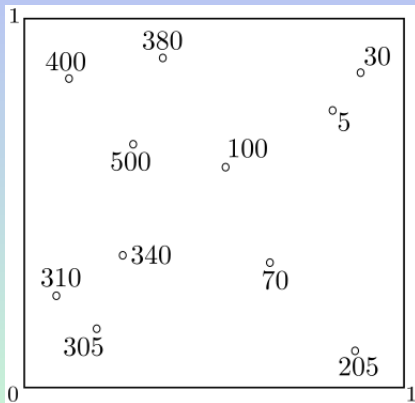
$$Y = m(\mathbf{X}) + \varepsilon.$$

We want to build an estimate of the regression function m using random forest algorithm.



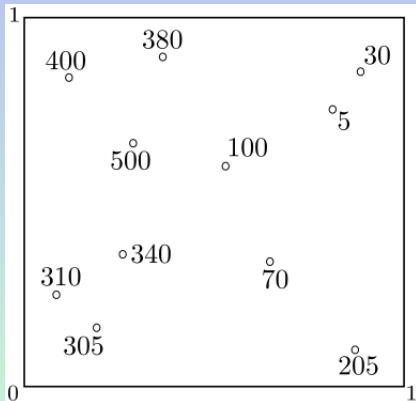
How to build a tree?

- Trees are built recursively by splitting the current cell into two children until some stopping criterion is satisfied.



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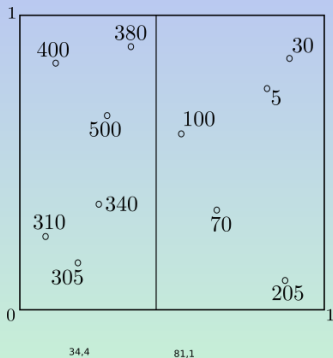


$k = 0$



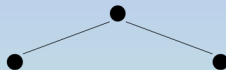
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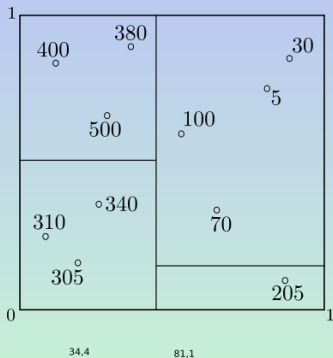
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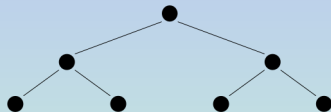
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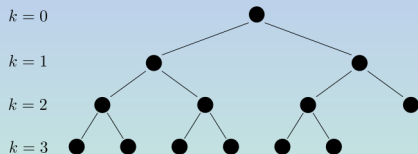
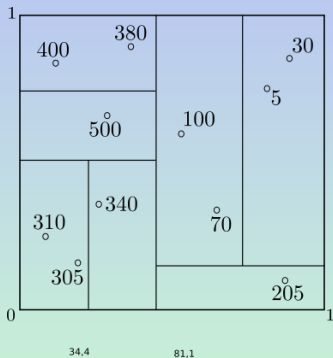
$k = 1$

$k = 2$



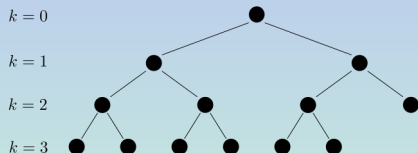
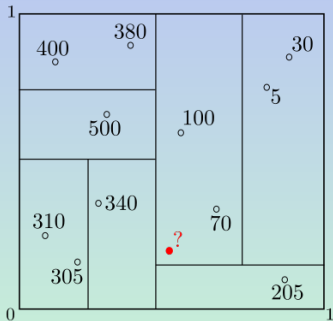
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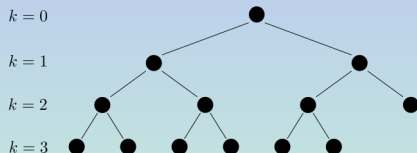
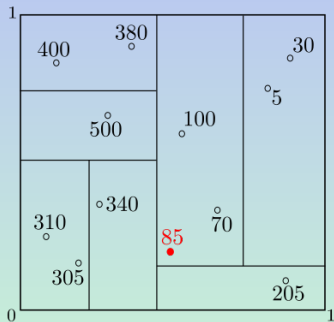
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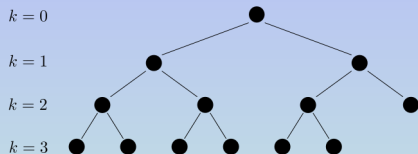
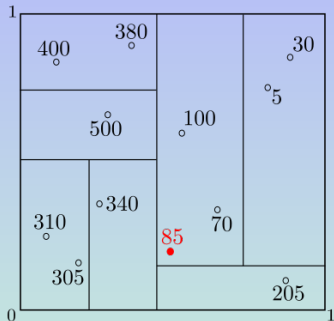


How to build a tree?

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How to build a tree?



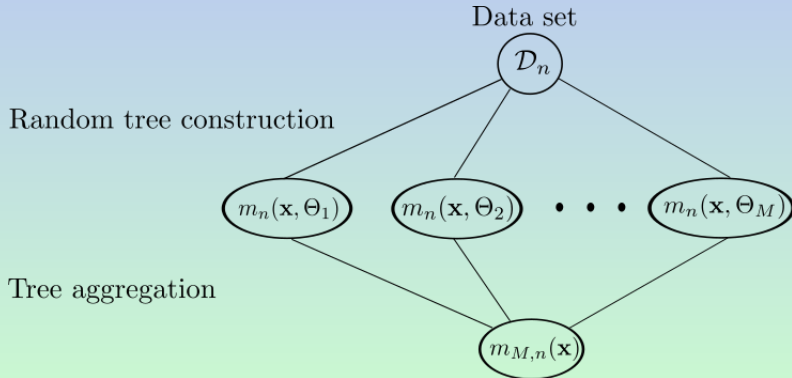
Breiman Random forests are defined by

- 1 A **splitting rule** : minimize the variance within the resulting cells.
- 2 A **stopping rule** : stop when each cell contains less than $\text{nodesize} = 2$ observations.

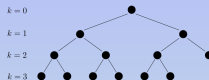
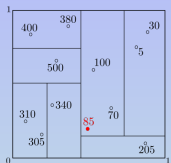
Construction of random forests

Randomness in tree construction

- Resample the data set via bootstrap;
- At each node, preselect a subset of m try variables eligible for splitting.



Construction of Breiman forests



Breiman tree

- Select a_n observations with replacement among the original sample \mathcal{D}_n . Use only these observations to build the tree.
- At each cell, select randomly m_{try} coordinates among $\{1, \dots, d\}$.
- Split at the location that minimizes the square loss.
- Stop when each cell contains less than nodesize observations.

- Random forests were created by Breiman [2001].
- Many theoretical results focus on [simplified version](#) on random forests, whose construction is [independent of the dataset](#). [Biau et al., 2008, Biau, 2012, Genuer, 2012, Zhu et al., 2012, Arlot and Genuer, 2014].
- Analysis of more data-dependent forests:
 - [Asymptotic normality](#) of random forests [Wager, 2014, Mentch and Hooker, 2015].
 - [Variable importance](#) [Louppe et al., 2013].
- Literature review on random forests:
 - [Methodological review](#) [Criminisi et al., 2011, Boulesteix et al., 2012].
 - [Theoretical review](#) [Biau and Scornet, 2016].

Different types of forests

Centred forest		

Different types of forests

Centred forest		
Independent of X_i and Y_i		

Different types of forests

Centred forest

Independent of X_i and Y_i



Different types of forests

Centred forest		Breiman's forests
Independent of X_i and Y_i		
		

Different types of forests

Centred forest		Breiman's forests
Independent of X_i and Y_i		Dependent on X_i and Y_i
		

Different types of forests

Centred forest



Breiman's forests

Independent of X_i and Y_i



Dependent on X_i and Y_i




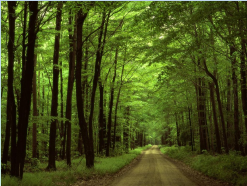

Different types of forests

Centred forest	Median forests	Breiman's forests
Independent of X_i and Y_i		Dependent on X_i and Y_i
		

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Centred forest	Median forests	Breiman's forests
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Tree consistency



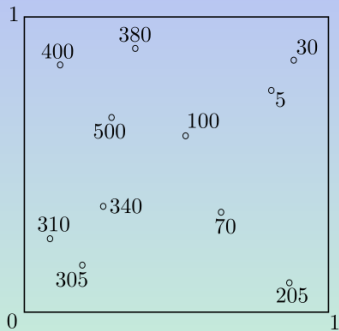
For a tree whose construction is independent of data, if

- 1 $\text{diam}(A_n(\mathbf{X})) \rightarrow 0$, in probability;
- 2 $N_n(A_n(\mathbf{X})) \rightarrow \infty$, in probability;

then the tree is consistent, that is

$$\lim_{n \rightarrow \infty} \mathbb{E} [m_n(\mathbf{X}) - m(\mathbf{X})]^2 = 0.$$

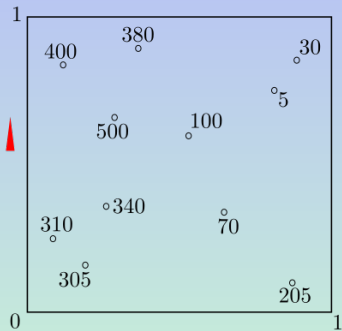
Centered forests



$k = 0$



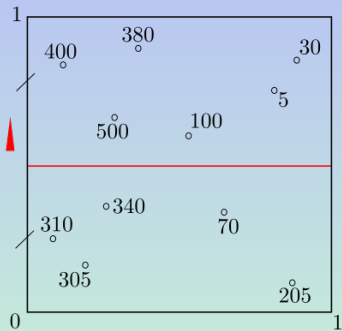
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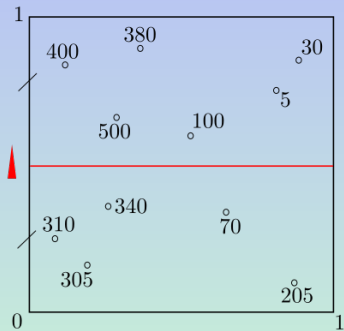
Centered forests



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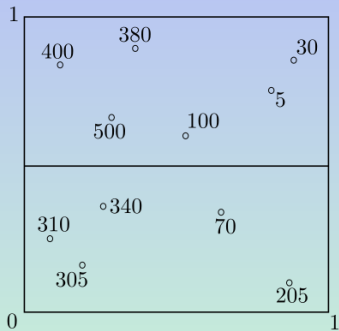
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Centered forests

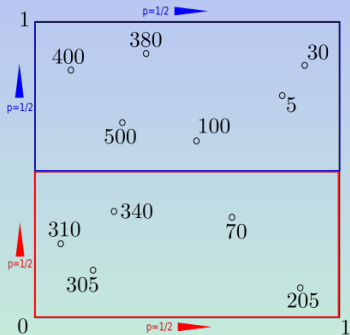


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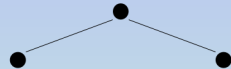


Centered forests

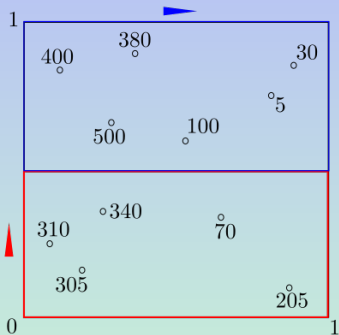


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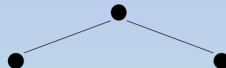


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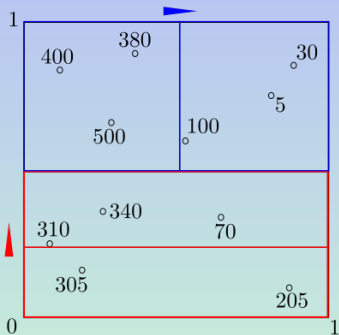


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Centered forests

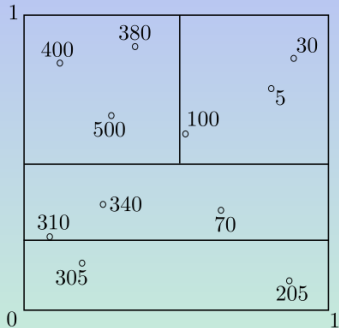


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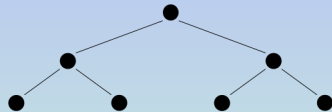
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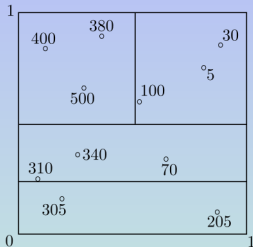
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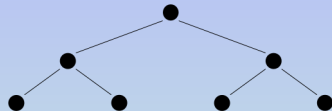
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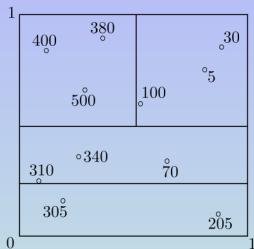
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Theorem (Biau [2012])

Under proper regularity hypothesis, provided $k \rightarrow \infty$ and $n/2^k \rightarrow \infty$, the centred random forest is consistent.

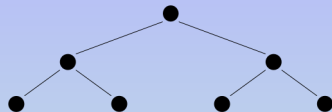
Centered forests



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Theorem (Biau [2012])

Under proper regularity hypothesis, provided $k \rightarrow \infty$ and $n/2^k \rightarrow \infty$, the centred random forest is consistent.

- Forest consistency results from the consistency of each tree.
- Trees are not fully developed.

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Construction of Breiman/Median forests

Breiman tree

- Select a_n observations with replacement among the original sample \mathcal{D}_n . Use only these observations to build the tree.
- At each cell, select randomly m_{try} coordinates among $\{1, \dots, d\}$.
- Split at the location that minimizes the square loss.
- Stop when each cell contains less than $nodesize$ observations.

Construction of Breiman/Median forests

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- Stop when each cell contains less than nodesize observations.

Median tree

- Select a_n observations without replacement among the original sample \mathcal{D}_n . Use only these observations to build the tree.
- At each cell, select randomly $m_{\text{try}} = 1$ coordinate among $\{1, \dots, d\}$.
- Split at the location of the empirical median of X_i .
- Stop when each cell contains exactly $\text{nodesize} = 1$ observation.

Theorem

Assume that **(H1)** is satisfied. Then, provided $a_n \rightarrow \infty$ and $a_n/n \rightarrow 0$, median forests are consistent, i.e.,

$$\lim_{n \rightarrow \infty} \mathbb{E} [m_n(\mathbf{X}) - m(\mathbf{X})]^2 = 0.$$

Remarks

- Good trade-off between simplicity of centred forests and complexity of Breiman's forests.
- First consistency results for fully grown trees.
- Each tree is not consistent but the forest is, because of subsampling.

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Construction of Breiman forests

Breiman tree

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Construction of Breiman forests

Breiman tree

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- Split at the location that minimizes the square loss.
- Stop when each cell contains less than nodesize observations.

Modified Breiman tree

- Select a_n observations without replacement among the original sample \mathcal{D}_n . Use only these observations to build the tree.
- At each cell, select randomly m_{try} coordinates among $\{1, \dots, d\}$.
- Split at the location that minimizes the square loss.
- Stop when the number of cells is exactly t_n .

Assumption (H1)

Additive regression model:

$$Y = \sum_{i=1}^d m_i(\mathbf{X}^{(i)}) + \varepsilon,$$

where

- \mathbf{X} is uniformly distributed on $[0, 1]^d$,
- $\varepsilon \sim \mathcal{N}(0, \sigma^2)$ with ε independent of \mathbf{X} ,
- Each model component m_i is continuous.

Theorem [Scornet et al., 2015]

Assume that **(H1)** is satisfied. Then, provided $a_n \rightarrow \infty$ and $t_n(\log a_n)^9/a_n \rightarrow 0$, random forests are consistent, i.e.,

$$\lim_{n \rightarrow \infty} \mathbb{E} [m_{\infty, n}(\mathbf{X}) - m(\mathbf{X})]^2 = 0.$$

Remarks

- First consistency result for Breiman's original forest.
- Consistency of CART.

Theorem [Scornet et al., 2015]

Assume that **(H1)** and **(H2.1)** are satisfied and let $t_n = a_n$. Then, provided $a_n \rightarrow \infty$ and $a_n \log n/n \rightarrow 0$, random forests are consistent, i.e.,

$$\lim_{n \rightarrow \infty} \mathbb{E} [m_{\infty, n}(\mathbf{X}) - m(\mathbf{X})]^2 = 0.$$

Remarks:

- First result for fully developed forest;
- Importance of subsampling;
- One major drawback: **(H2)** seems impossible to verify.

Sparsity and random forests

- Assume that

$$Y = \sum_{i=1}^S m_i(\mathbf{X}^{(i)}) + \varepsilon,$$

for some $S < d$.

- Denote by $j_{1,n}(\mathbf{X}), \dots, j_{k,n}(\mathbf{X})$ the first k cut directions used to construct the cell containing \mathbf{X} .

Proposition [Scornet et al., 2015]

Let $k \in \mathbb{N}^*$ and $\xi > 0$. Under appropriate assumptions, with probability $1 - \xi$, for all n large enough, we have, for all $1 \leq q \leq k$,

$$j_{q,n}(\mathbf{X}) \in \{1, \dots, S\}.$$

Conclusion

- **Centred forests**: their consistency results from the consistency of each tree.
→ No benefits from using a forest instead of a single tree.
- **Median forests**: the aggregation process can turn inconsistent trees into a consistent forest.
→ Benefits from using a random forest compared to a single tree.
- **Breiman forests**: consistent as well as CART procedure. The splitting criterion asymptotically selects relevant features.
→ Good performance in high-dimensional settings.



Merci pour votre attention !

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How to deal with fully developed Breiman forest?

Let

$$\psi_{i,j}(Y_i, Y_j) = \mathbb{E} \left[\mathbb{1}_{\mathbf{X} \leftrightarrow \mathbf{X}_i} \mathbb{1}_{\mathbf{X} \leftrightarrow \mathbf{X}_j} \mid \mathbf{X}, \Theta, \Theta', \mathbf{X}_1, \dots, \mathbf{X}_n, Y_i, Y_j \right]$$

and $\psi_{i,j} = \mathbb{E} \left[\mathbb{1}_{\mathbf{X} \leftrightarrow \mathbf{X}_i} \mathbb{1}_{\mathbf{X} \leftrightarrow \mathbf{X}_j} \mid \mathbf{X}, \Theta, \Theta', \mathbf{X}_1, \dots, \mathbf{X}_n \right]$.

One assumption **(H2.1)**:

$$\lim_{n \rightarrow \infty} (\log a_n)^{2p-2} (\log n)^2 \mathbb{E} \left[\max_{\substack{i,j \\ i \neq j}} |\psi_{i,j}(Y_i, Y_j) - \psi_{i,j}| \right]^2 = 0.$$