





# Agenda

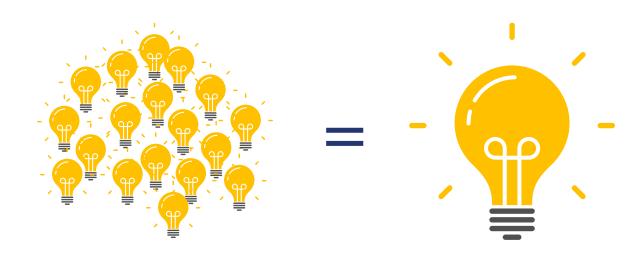
- Lecture 2: Generalization error, bias and variance, data splitting, Cross-validation.
- Lecture 3: Model evaluation
- Lecture 4: Decision Trees Grow, Splitting and Stopping criteria, Prune, Final Evaluation
- EL: SVM (datacamp)

#### **Today**

- Introduction to ensemble models
- Random Forest, ADAboost and XGBoost
- New Assignment
- Coding session

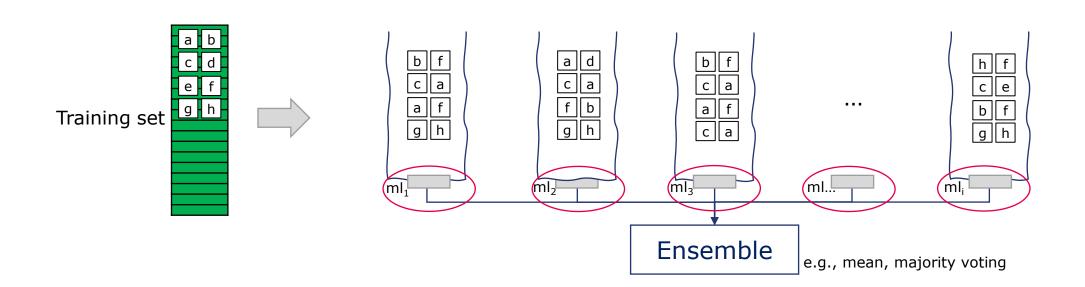


• A highly accurate model can be obtained by a collection of multiple individual weak learners



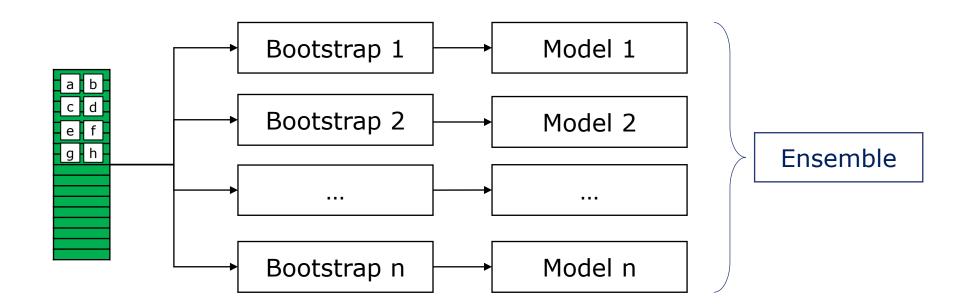


• Remember bagging?





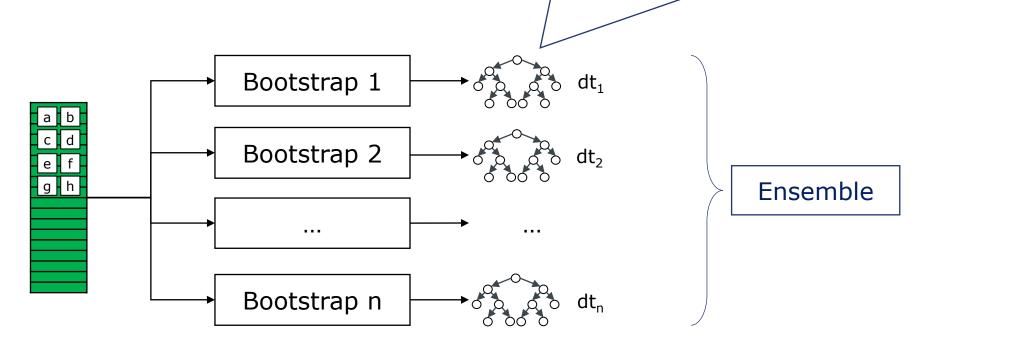
#### Bagging





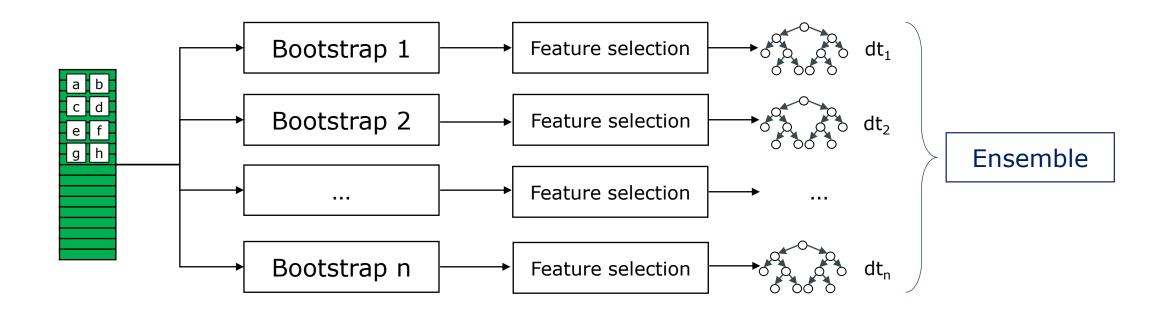
Bagging with trees

PROBLEM: If one feature is dominant the trees will always split based on that feature





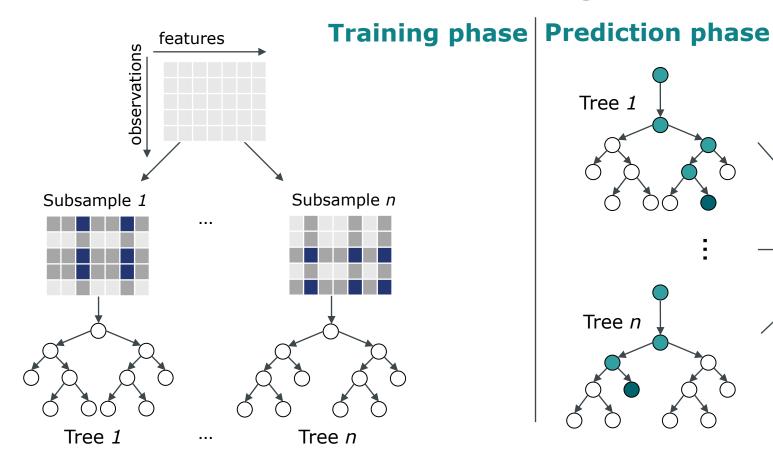
Bagging + feature selection + trees = RANDOM FOREST





# Random Forest – A fast glance!

Bagging, works in paralel



# Tree 1 Predicted distribution Regression problems Majority voting Classification problems



#### Example

	Gender	Age	Smoke	ВМІ	Heart Disease
1	М	28	0	46	0
2	М	39	1	116	1
3	F	55	0	66	0
4	М	18	0	46	0
5	F	87	1	88	0
6	F	58	1	135	1
7	F	77	0	65	0
8	М	60	0	116	1
9	М	42	0	74	1



#### Example

	Gender	Age	Smoke	ВМІ	Heart Disease	
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Subsetting

- A percentage of samples (rows)
- A number of features (columns)
- Create a decision tree



#### Example

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- Repeat a number of times



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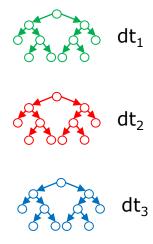


- Subsetting
  - A percentage of samples (rows)
  - A number of features (columns)
- Create a decision tree
- Repeat a number of times
- We now have three trees, each with their own understanding of the problem, i.e., each with their own prediction
- Mean or majority voting



Main parameters - for R: ranger, for Py: sklearn)

	Gender	Age	Smoke	ВМІ	Heart Disease
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R: sample.fraction (0.63) **Py**: max\_samples (None)

**R:** mtry (sqrt)

**Py:** max\_features (sqrt)

- Subsett
  - min.node.size (1) A percent ge of samp max.depth (unlim)
  - A number of features
- Create a decision tree
- Repeat a number of times

**R:** num.trees (500)

e trees, each with their **Python:** n\_estimators (100) own understanding of the problem, i.e., each with their own prediction

Mean or majority voting

(see decision trees)



#### Two more things

OOB Score

		Gender	Age	Smoke	ВМІ	Heart Disease	
$\subseteq$	1	M	28	0	46	0	$\Box$
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These observation not picked. Should we waste them? No! Out-Of-Bag (OOB) samples. Can be used to evaluate the model! For free!

This is an oversimplification. What really happens is:

- For every tree, keep track of which samples were OOB.
  - In our example: 1 is OOB for red and blue, 2 is OOB for green, 3 is OOB for all trees, etc.
- At the end, a prediction is done by passing each sample through all trees for which they were OOB.
  - In our example: 1 predicted by using only red and blue, 2 predicted by using only green, 3 predicted by using the entire forest
- The OOB score is the number of correctly predicted rows from the OOB sample



#### Two more things

Feature importance

		Gender	Age	Smoke	ВМІ	Heart Disease
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Mean Decrease impurity (specific to Random Forest)

We created trees from subset of features!
We could see how much each feature contributed to the decrease in purity when they were selected! For free!

Feature Permutation importance (generic)

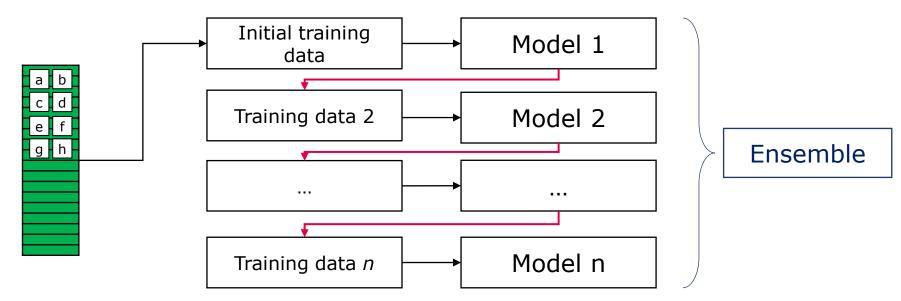
If no change feature is useless

We take <u>one</u> feature and we randomly <u>reshuffle</u> it. Then we see <u>how</u> this <u>reschuffling impacted</u> the prediction <u>error</u> (the larger the impact the more important the feature). Not for free, but more robust.



#### Boosting

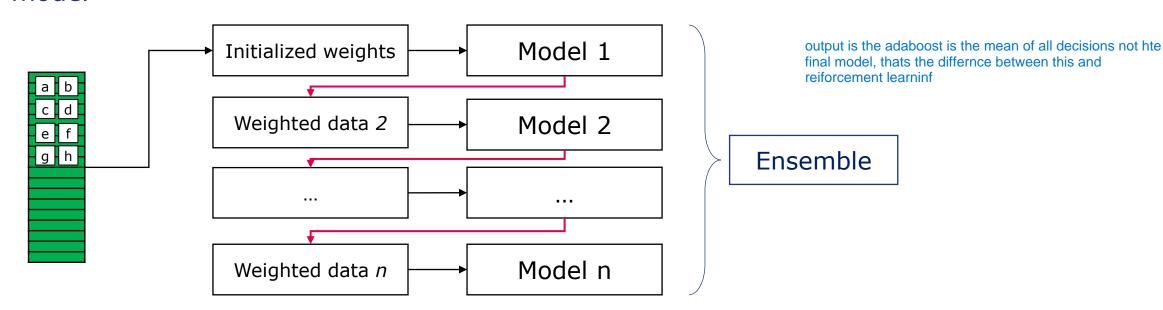
• Differently from bagging, boosting is not parallel. The models are built in sequence, each one considering the output of the previous one.





#### Adaboost

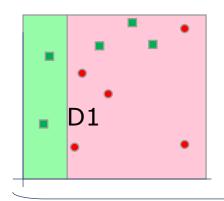
 In AdaBoost, the output of one model is used to weight the input data of the next model

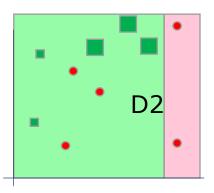


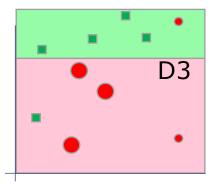


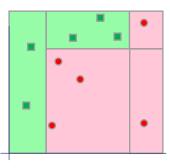
#### Adaboost Example

slower then RF





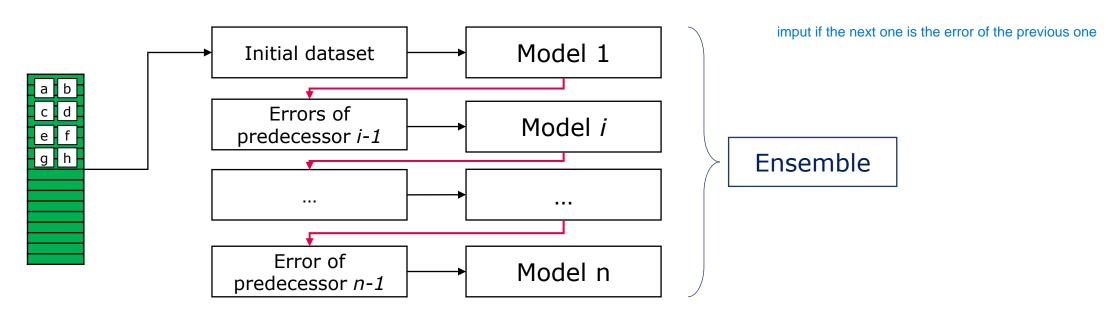






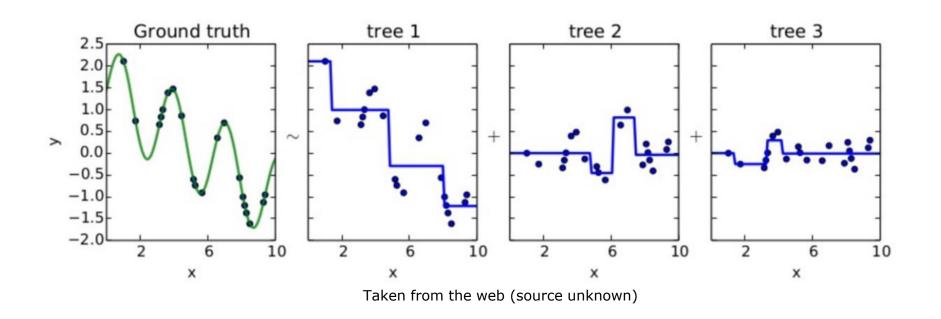
#### **Gradient Boosting**

In Gradient Boosting, each model fits the residual error of the previous model





#### **Gradient Boosting Example**





# **Gradient Boosting**

#### Example of parameters

- Shrinkage:
  - **Iterations (Py:** *num\_round,* **R:** *nrounds*): num of boosting iterations to perform (i.e., n. of trees). The more trees the better but at higher computational costs. It is the main parameter to control model performance. Typical value: Iterations > 100.
  - **Learning rate (Py: eta, R: eta)**: Scales the contribution of each iteration. Small learning rates lead to higher number of iterations. Typical value: [0.01,...0.1,...,0.5], default = 0.3.
  - Min\_split\_loss (Py: gamma, R: gamma): Minimum loss reduction required to make a further partition on a leaf node of the tree. The larger gamma, the more conservative the algorithm. Like purity gain in decision trees.
- + all the familiar ones from decision trees (maxdepth, min node size, etc.)
- Super documentation!!!
  - https://xgboost.readthedocs.io/en/latest/R-package/index.html
  - https://xgboost.readthedocs.io/en/latest/python/index.html



### Ensemble models from decision trees

#### Summary

- Ensemble
  - Combining multiple (weak) learners
  - Random Forest:
    - Bagging + feature selection + Decision Tree
  - AdaBoost
    - Boosting (each model is trained on a dataset weighted depending on previous models) +
       Decision Tree
  - Gradient Boosting:
    - Boosting (each model is trained based on the residuals of the previous model) + Decision Tree
    - XGBoost is an implementation of Gradient Boosting



## Exercise 2 - Due on 04.04.2021 23.59 CET

Import the EEG data from the files section in Teams







- 16 EEG numerical features
- 1 label (eyes closed or open)
- Train and test are already split (eeg\_training.csv, eeg\_test.csv)
- Create a classifier that can detect if the patient has open or closed eyelid
- Try Decision Trees, Random Forest, AdaBoost, and XGBoost
- Attention: This is not the same data that can be found online. Do not copy!
- Hints: Are there NAs? Is it balanced? Are there useless features? Are there outliers?



# Appendix I

#### Strategies for preprocessing

- Missing values
  - Omit the observations containing NAs -> If there are only fews
  - Omit the features containing NAs -> If a feature contains mostly NAs
  - Impute the NAs -> Manual (e.g. median), Manual by group of similars, model-based (eg. kNN)
- Imbalanced data
  - Undersampling -> Drop samples from the majority class
  - Oversampling -> Duplicate samples from the minority class
  - Synthetic generation of new data points (e.g. ROSE)



# Appendix II

