





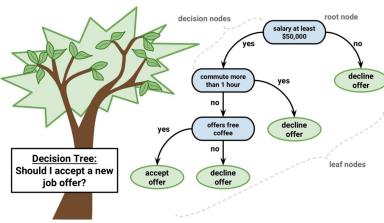
Ensemble Learning

Artificial Intelligence and Machine Learning

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What is a decision tree?

- Machine learning model with a flowchart-like structure
- Each internal nodes represents a "test" on a feature of our dataset
- Final nodes (leaf) are the predicted class or value
- Available both for classification and regression tasks



Example: Iris dataset

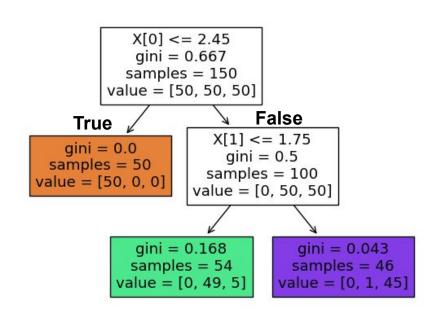
```
from sklearn.datasets import load iris
from sklearn.tree import DecisionTreeClassifier, plot tree
import matplotlib.pyplot as plt
iris=load iris()
X = iris.data[:,2:] #petal length and width
y = iris.target
clf=DecisionTreeClassifier(max depth=2)
clf.fit(X,y)
plot tree(clf, filled=True)
plt.show()
```

Iris decision tree

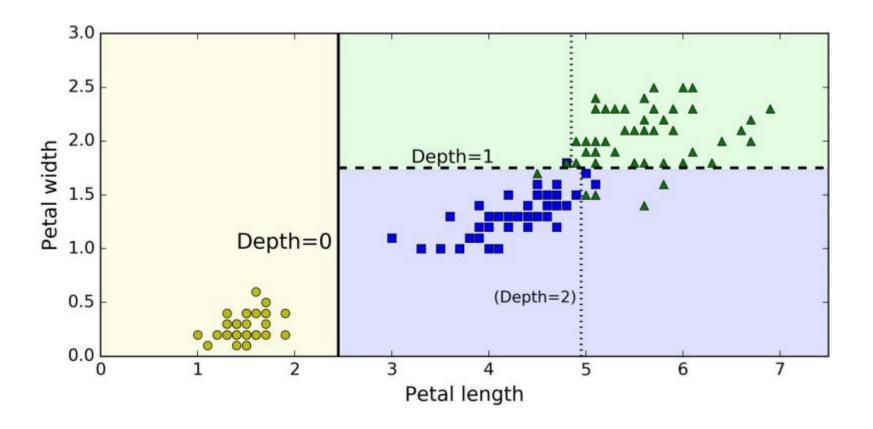
- X[0] : petal length
- X[1]: petal width
- Gini: Impurity metric
 - A node is "pure" if Gini=0
 - If all training instances it applies to belong to the same class

$$G_i = 1 - \sum_{k=1}^{n} p_{i,k}^2$$

- $p_{i,k}$ is the ratio of class k instances among the training instances in the i^{th} node.
- Gini for the depth-2 left node is $1 (0/54)^2 (49/54)^2 (5/54)^2 = 0.168$
- Ratios corresponds also to output probabilities:
 - 0% for Iris-setosa
 - 90.7% for iris-Versicolor
 - 9,3% for iris-Virginica



Decision boundaries



CART training algorithm

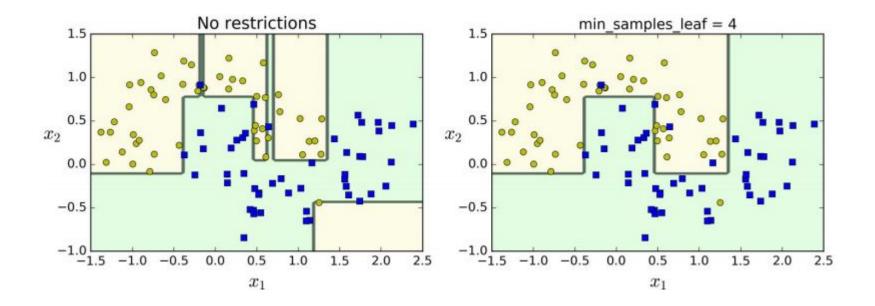
- CART (Classification and Regression Tree algorithm)
- Simple idea:
 - Splits the training set in two subsets using a single feature k and a threshold t_k (e.g., "petal length <= 2.45")
 - b. To choose k and t_k it looks for the purest subsets (weighted by their size) minimizing a cost function
 - c. It stops once it reaches the maximum depth selected as a parameter

$$J(k, t_k) = \frac{m_{\text{left}}}{m} G_{\text{left}} + \frac{m_{\text{right}}}{m} G_{\text{right}}$$
 where
$$\begin{cases} G_{\text{left/right}} \text{ measures the impurity of the left/right subset,} \\ m_{\text{left/right}} \text{ is the number of instances in the left/right subset.} \end{cases}$$

Problems: Overfitting

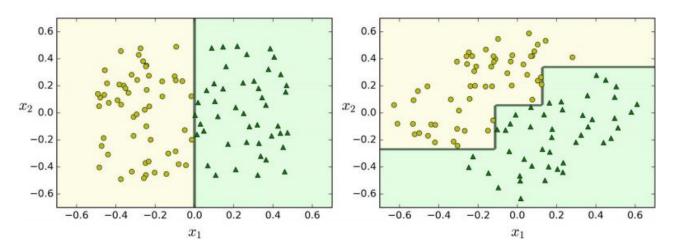
- Decision trees make less assumptions about the training data with respect to linear models
- If left unconstrained the tree structure will adapt itself to the training-set very well!
 - Actually, most of the time it overfits the training-set!
- One solution can be reduce the degree of freedom of the model during the training phase (Regularization)
 - Select a max depth
 - Pruning
 - Select a a minimum number of samples a leaf node must have
 - Select a max feature number that are evaluated for splitting each node

Regularization with min sample leaf



Instability

- A decision tree often finds orthogonal decision boundaries
- Orthogonality makes them sensitive to training set rotation
- A decision tree can fits perfectly a training set but probably will not generalize well in some cases
- An idea to solve this problem is to apply the Principal Component Analysis (PCA) to data in order to have a better orientation of the training data

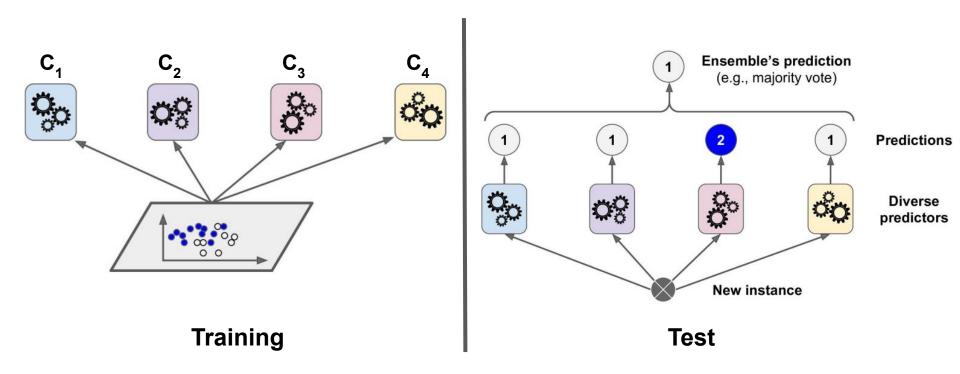


Ensemble learning

- When you are looking for an answer to a complex problem it is better to ask to different experts and then aggregate their answer (Wisdom of the crowd)
- Key idea: aggregate the predictions of a group of predictors to get better predictions than the best single predictor



Ensemble learning



Why the aggregation is better?

- Suppose you have a slightly biased coin: 51% chance of coming up heads
 - If you toss the coin 1000 times: you will have more or less 510 heads and 490 tails
 - What is the probability of having a majority of heads after 1000 tosses?
 - It is around 73%
 - If 10'000 tosses: the probability is around 95%
 - The more you toss the coin, the higher the probability is

- This is true, when each toss is independent
 - The result of each toss does not depend on the others!



Why the aggregation is better?

- N = 1000
- k = n° of heads that in N tosses has to be at least higher than (N/2)+1
- The probability that after N tosses the majority are heads is the sum of the probability of k successes with k in range of (N/2)+1 to N

$$\sum_{k=(N/2)+1)}^{N} \binom{N}{k} p^k (1-p)^{n-k}$$

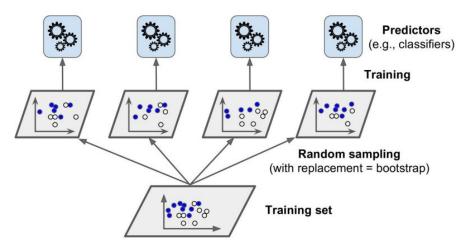


Independent predictors

- Similarly, we can build 1000 classifiers that are individually correct only 51% of the time (slightly better than the random guessing -> weak classifier)
- If you predict the majority voted class, you can hope to 73% accuracy
- This is true if all classifiers are perfectly independent!
 - They should make uncorrelated errors, otherwise they are likely to make the same type of errors and the majority could be wrong!
 - If all classifiers are trained on the same dataset, are they independent?

Bootstrap aggregating (Bagging)

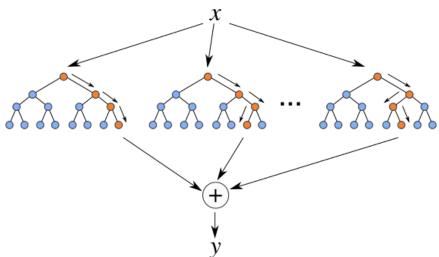
- We have only one training-set and we want to use only one algorithm
- We can train the classifiers on different random subsets of the training-set
- When sampling is performed with replacement, this method is called bootstrap
- Each individual predictor has a higher bias than if it were trained on the original dataset, but aggregation reduces both bias and variance



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Random forest

- Random forest is a bagging-based model where we make an aggregation of decision trees
- It also sub-samples a fraction of the features when fitting a decision tree to each bootstrap sample



Hard voting classifier

- Another solution to get independent classifiers is to use different algorithms
- Then we can make hard voting

```
from sklearn.linear_model import LogisticRegression
from sklearn.ensemble import RandomForestClassifier
from sklearn.svm import SVC
from sklearn import datasets
from sklearn.metrics import accuracy_score
from sklearn.model_selection import train_test_split
from sklearn.ensemble import VotingClassifier
```

```
# import data
X, y = datasets.load_breast_cancer(return_X_y=True)
```

Hard voting classifier

```
log clf = LogisticRegression()
rf clf= RandomForestClassifier(n estimators = 10)
svm clf = SVC()
X train, X test, y train, y test = train test split(X, y,
test size=0.33, stratify=y)
voting clf=VotingClassifier(estimators=[('lr',log clf),('rf',rf
clf), ('svc', svm clf)], voting='hard')
voting clf.fit(X train, y train)
y pred= voting clf.predict(X test)
print("Voting "+str(accuracy score(y test, y pred)))
```

Hard voting classifier

```
log clf.fit(X train, y train)
y pred= log clf.predict(X test)
print("Logistic "+str(accuracy score(y test, y pred)))
rf clf.fit(X train, y train)
y pred = rf clf.predict(X test)
print("Random forest "+str(accuracy score(y test, y pred)))
svm clf.fit(X train, y train)
y pred = svm clf.predict(X test)
print("Support Vector Machine
"+str(accuracy score(y test, y pred)))
```

Bagging with sklearn

from sklearn.ensemble import BaggingClassifier

```
from sklearn.ensemble import BaggingClassifier
model= BaggingClassifier(n_estimators = 200)
model.fit(X_train,y_train)
```

```
from sklearn.ensemble import RandomForestClassifier
model= RandomForestClassifier(n_estimators = 1000)
model.fit(X train,y train)
```

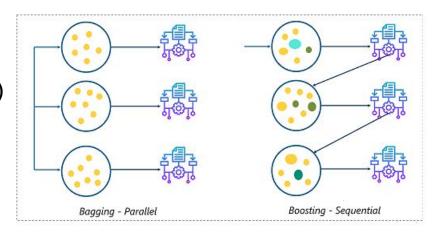
Exercise Breast cancer

- X, y = datasets.load_breast_cancer(return_X_y=True)
- Split data (70% training and 30% test)
- Train a single decision tree, a bagging classifier with 200 trees and a random forest
- Repeat each training 10 times and consider the average accuracy
- What is the best classifier?
- ADDITIONAL for home: Try to use regularization parameters:
 - https://scikit-learn.org/stable/modules/generated/sklearn.ensemble.Baggi ngClassifier.html
 - https://scikit-learn.org/stable/modules/generated/sklearn.ensemble.Rand omForestClassifier.html
 - https://scikit-learn.org/stable/modules/generated/sklearn.tree.DecisionTre eClassifier.html

BREAK

Boosting

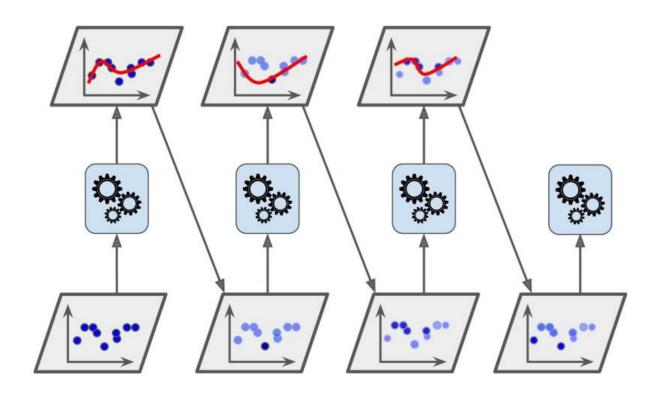
- Similar to bagging, but now the idea is to train predictors sequentially
- Each predictor tries to correct its predecessor
- Many boosting methods available:
 - Adaptive boosting (AdaBoost)
 - Gradient boosting
 - XGboost (Extreme gradient boosting)



AdaBoost

- To correct a predecessor one way is to pay attention to the training instances that the predecessor under-fitted
- The new predictor will focus more on the hard cases (adaptive)
 - A first base predictor is trained and used to make predictions on the training-set
 - b. The relative weight of misclassified training instances is then increased
 - c. A second classifier is trained using the updated weights and so on
 - d. To each classifier a weight is assigned
 - Once all predictors are trained, the ensemble makes predictions like bagging, except that predictors have different weights depending on their accuracy on the training-set

AdaBoost



AdaBoost algorithm

- Each instance weight w⁽ⁱ⁾ is initially set to 1/m where m is the number of observations
- For each predictor j is computed a weighted error-rate r_j

$$r_{j} = \frac{\sum_{i=1}^{m} w^{(i)}}{\sum_{i=1}^{m} w^{(i)}}$$

where y-hat is the jth predictors' prediction for the ith instance

AdaBoost algorithm

- A predictor's weight α_i is then computed using a learning-rate parameter η
- The more accurate the predictor is, the higher its weight will be.

$$\alpha_j = \eta \log \frac{1 - r_j}{r_j}$$

- After that, the instance weights are updated using the equation below
- All the instance weights are then normalized dividing by the sum of all weights

for
$$i = 1, 2, \dots, m$$

$$w^{(i)} \leftarrow \begin{cases} w^{(i)} & \text{if } \hat{y}_j^{(i)} = y^{(i)} \\ w^{(i)} \exp(\alpha_i) & \text{if } \hat{y}_i^{(i)} \neq y^{(i)} \end{cases}$$

AdaBoost predictions

- \bullet Finally, AdaBoost computes the predictions of all predictors and weights them using α_{j}
- The predicted class is the one that gets the majority of weighted votes

$$\hat{y}(\mathbf{x}) = \underset{k}{\operatorname{argmax}} \sum_{j=1}^{N} \alpha_j \qquad \text{where N is the number of predictors}$$

$$\hat{y}_j(\mathbf{x}) = k$$

Gradient Boosting

- The key idea of GB is additive modelling
 - We can have a final learner $F_L = f_0(x) + f_1(x) + f_2(x) + ... + f_{L-1}(x)$
 - Where L is the number of learners we want to use (for example n°of trees)
 - \circ $f_0(x)$ is the first predictor we train on the training-set
 - Each epoch of training we add a new learner that tries to increase the accuracy of the previous one by minimizing a Loss function using gradient descent and not by increasing instances' weight like AdaBoost

- Let's play a game: You are given a dataset in the form of $(x_1, y_1), ..., (x_n, y_n)$.
- The task is to fit a model F(x) to minimize the square loss (sum of square difference between y_true and y_pred)
- Suppose your friend wants to helè you and gives you his model F
- You check his model and find the model is good but not perfect because it makes some mistakes:
 - For example: $F(x_1) = 0.5$ while $y_1 = 0.6$
 - \circ F(x₂) = 1.8 while y₂=1.7
- You cannot change the model F or change its parameters but you want to improve it

- We cann an additional model (regression tree for example) h to F, so the new prediction will be F(x)+h(x)
- You wish to improve the model such that

$$F(x_1) + h(x_1) = y_1$$

$$F(x_2) + h(x_2) = y_2$$

$$F(x_n) + h(x_n) = y_n$$

$$h(x_1) = y_1 - F(x_1)$$

$$h(x_2) = y_2 - F(x_2)$$

Can any regression tree h achieve this goal perfectly? Probably not! But some regression might be able to do this approximately

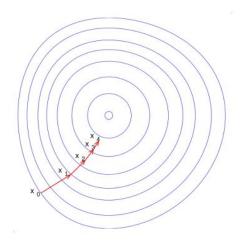
- To get this predictor h we can fit a regression tree with a different formulation of the original dataset where the target value to predict will be y F(x)
- $\{x_1, y_1 F(x_1), ..., (x_n, y_n F(x_n))\}$
- y_i $F(x_i)$ are usually called residuals
- We want to minimize the entire loss for all training examples (1,...,n)
- To minimize this loss we can use the gradient descent

Gradient descent

Gradient Descent

Minimize a function by moving in the opposite direction of the gradient.

$$\theta_i := \theta_i - \rho \frac{\partial J}{\partial \theta_i}$$



- Loss function L(y, F(x)) = $(y F(x))^2 / 2$
- We want to minimize $J = \sum_{i} L(y_i, F(x_i))$ by adjusting $F(x_1), F(x_2) \dots F(x_n)$

$$\frac{\partial J}{\partial F(x_i)} = \frac{\partial \sum_i L(y_i, F(x_i))}{\partial F(x_i)} = \frac{\partial L(y_i, F(x_i))}{\partial F(x_i)} = F(x_i) - y_i$$

So we can interpret residuals as negative gradients

$$y_i - F(x_i) = -\frac{\partial J}{\partial F(x_i)}$$

 So at the end we can go back to see the predictors at stage m+1 and define it as

$$F_{m+1}(x) = F_m(x) + h(x) = y$$

$$F_{m+1}(x) = F_m(x) - \nabla_m L(y, F_m(x)) = F_m(x) - \left[\frac{\partial L(y, F_m(x))}{\partial F_m(x)}\right]$$

With a learning-rate equal to 1

Boosting with sklearn

```
from sklearn.ensemble import AdaBoostClassifier
model= AdaBoostClassifier(n estimators = 200)
model.fit(X train, y train)
from sklearn.ensemble import GradientBoostingClassifier
model= GradientBoostingClassifier(n estimators = 200)
model.fit(X train, y train)
from xqboost import XGBClassifier
model = XGBClassifier(n estimators = 200)
model.fit(X train, y train)
```

Exercise - Classification

- X, y = datasets.load_breast_cancer(return_X_y=True)
- Train an ensemble model with 200,1000,2000 trees with AdaBoost, GradientBoosting and XGBoost
- Compute the accuracy and the confusion matrix
- Which model performs better?
- How results change if you increase the test-set size percentage?
- Try to use regularization parameters

Exercise 2 - Regression

- X, y = datasets.load_boston(return_X_y=True)
- Train an ensemble model with 200,1000,2000 trees with AdaBoost, GradientBoosting and XGBoost
- Compute the rmse
- Which model performs better?
- How results change if you increase the test-set size percentage?
- Try to use regularization parameters