

# CZ 4041 - Machine Learning

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AY 16/17, Semester 2

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# Chapter 1

## Introduction

### 1.1 Definition

Machine learning is a type of artificial intelligence that provides computers with the ability to learn from examples and/or experience without being explicitly programmed.

### 1.2 Different Paradigms

#### 1.2.1 Supervised Learning

- The examples presented to the computer are pairs of inputs (i.e. features) and the corresponding outputs (i.e. labels). Labelled training data.
- If the labels are discrete, it is a **classification** problem.
- If the labels are continuous, it is a **regression** problem.
- The goal is to *learn* a **model** that maps inputs to the correct labels.

$$f : \text{label} = f(\text{input})$$

#### 1.2.2 Unsupervised Learning

- The examples presented to the computer are a set of inputs without any outputs. Unlabelled training data.
- The goal is to *learn* an **intrinsic structure** of the examples. For example, the goal in such problems may be to discover groups of similar examples within the data (i.e. clustering) or to determine the distribution of data within the input space (i.e. density estimation).

#### 1.2.3 Semi-Supervised Learning

- The examples presented to the computer include both labelled data and unlabelled data.
- The goal is to utilize the unlabelled data to help supervised learning.
- More precise f.

#### 1.2.4 Active Learning

- The examples presented to the computer are unlabelled.
- An active learner (i.e. computer) may pose queries in the form of unlabelled examples to be labelled by an oracle (e.g. human annotator).

#### 1.2.5 Transfer Learning

- The goal is to adapt the same learning to different tasks or environments.

#### 1.2.6 Reinforcement Learning

- Learning by **interacting** with an environment to achieve a goal.
- The goal is to learn an optimal policy mapping states to actions.

# Chapter 2

## Overview of Supervised Learning

- In supervised learning, the examples presented to a computer are pairs of inputs and the corresponding outputs (i.e. labelled training data).
- The goal is to *learn* a **model** that maps inputs to the correct labels.
- Given a set of  $\{\mathbf{x}_i, y_i\}$  (i.e. labelled training data) for  $i = 1, \dots, N$  where  $\mathbf{x}_i = [x_{i1}, x_{i2}, \dots, x_{im}]$  (i.e. a vector of features) and  $y_i$  (i.e. output) is a scalar, the goal is to learn a mapping  $f : x \rightarrow y$  (i.e. classifier) by requiring  $f(\mathbf{x}_i) = y_i$ .
- The learned mapping  $f$  is expected to be able to make precise predictions on any unseen  $\mathbf{x}^*$  as  $f(\mathbf{x}^*)$ .

### 2.1 Classification vs. Regression

- In classification problems,  $y$  is discrete.
  - If  $y$  is binary, then it is a binary classification problem.
  - If  $y$  is nominal but not binary, then it is a multi-class classification problem.
- In regression problems,  $y$  is continuous.

### 2.2 Typical Learning Procedure

**Inductive Learning** consists of two phases:

1. Training Phase
  - Given labelled training data, apply supervised learning algorithms to learn a model  $f$  such that  $f(\mathbf{x}_i) = y_i$ .
2. Testing or Prediction Phase
  - Given unseen test data  $\mathbf{x}_i^*$ , use the trained model  $f$  to make predictions for  $f(\mathbf{x}_i^*)$ .

**Lazy Learning** involves taking an unseen test data  $\mathbf{x}_i^*$  and using an algorithm to find the appropriate  $\{\mathbf{x}_i, y_i\}$  pair from the training data to output a predicted label  $y^*$ .

### 2.3 Evaluation of Supervised Learning

- The learned predictive model should be able to make predictions on previously unseen data as accurately as possible.
- The whole training data is usually divided into two sets for training and testing.
- The training set is used to learn the predictive model and the test set is used to validate the performance of the learned model.

#### Common Evaluation Metrics:

- Classification: Accuracy, Error Rate, F-Score etc.
- Regression: Mean Absolute Error (MAE), Root Mean Squared Error (RMSE)

#### Splitting Training Data to Disjoint Training & Evaluation Sets:

- Random Subsampling: Randomly sample a fraction of the data for evaluation and the rest for training. Repeat this process for several iterations.
- Cross Validation: Divide the entire dataset into  $k$  subsets of the same size. Hold aside one subset for evaluation and use the others to build the model. Traverse through all the subsets, changing the training subset at each iteration. For example, a five-fold cross validation will have five iterations.
- Bootstrap: Similar to random subsampling, however, the training set is sampled with replacement (i.e. duplicate samples allowed).

# Chapter 3

## Bayesian Decision Theory

### 3.1 Bayesian Classifiers

In many applications, the relationship between the input features and output labels is non-deterministic (i.e. uncertain). Bayesian classifiers aim to model probabilistic relationships between the input features and output class.

### 3.2 Math Review - Probability Concepts

Let  $X$  and  $Y$  be random variables.

**Marginal probability** refers to the probability that  $Y$  will take on the value  $y$ .

$$P(Y = y)$$

$$\sum_{y_i} P(Y = y_i) = 1$$

**Joint probability** refers to the probability that variable  $X$  will take on the value  $x$  and variable  $Y$  will take on the value  $y$ .

$$P(X = x, Y = y)$$

**Conditional probability** refers to the probability that the variable  $Y$  will take on the value  $y$ , given that the variable  $X$  is observed to have the value  $x$ .

$$P(Y = y|X = x)$$

$$\sum_{y_i} P(Y = y_i|X = x) = 1$$

**Sum Rule** — The joint and marginal probabilities for  $X$  and  $Y$  are related.

$$P(X = x) = \sum_{y_i} P(X = x, Y = y_i)$$

$$P(X) = \sum_Y P(X, Y)$$

$$P(X = x) = \sum_{z_j} \sum_{y_i} P(X = x, Y = y_i, Z = z_j)$$

$$P(X) = \sum_Z \sum_Y P(X, Y, Z)$$

**Product Rule** — The joint and conditional probabilities for  $X$  and  $Y$  are related.

$$\begin{aligned} P(X = x, Y = y) &= P(Y = y|X = x) \times P(X = x) \\ &= P(X = x|Y = y) \times P(Y = y) \end{aligned}$$

$$P(X, Y) = P(Y|X) \times P(X) = P(X|Y) \times P(Y)$$

#### 3.2.1 Decision with Priors

A decision rule prescribes what action to take based on observed input.

Predict  $Y = y_i$  if:

$$P(Y = y_i) = \max_k P(Y = y_k)$$

### 3.3 Bayes Theorem

$$P(Y|X) = \frac{P(X|Y)P(Y)}{P(X)}$$

- $P(Y|X)$ : Posterior Probability
- $P(X|Y)$ : Likelihood
- $P(Y)$ : Prior Probability
- $P(X)$ : Evidence

### 3.3.1 General Bayesian Classifier

Predict  $Y = y_i$  if:

$$P(Y = y_i | \mathbf{X} = \mathbf{x}) = \max_k P(Y = y_k | \mathbf{X} = \mathbf{x})$$

Applying Bayes Theorem:

$$P(Y = y_k | \mathbf{X} = \mathbf{x}) = \frac{P(\mathbf{X} = \mathbf{x} | Y = y_k)P(Y = y_k)}{P(\mathbf{X} = \mathbf{x})}$$

Since  $P(\mathbf{X} = \mathbf{x})$  is constant w.r.t to different values of  $Y$ , predict  $Y = y_i$  if:

$$P(\mathbf{X} | Y = y_i)P(Y = y_i) = \max_k P(\mathbf{X} | Y = y_k)P(Y = y_k)$$

## 3.4 Losses & Risks

So far, the decisions are assumed to be equally good or costly. However, the cost of misclassification for different classes may be different. For example, misdiagnosing a healthy patient as ill versus misdiagnosing an ill patient as healthy.

### 3.4.1 Formulae

- Let  $a_i$  be the action of predicting  $Y = y_i$ .
- Let  $\lambda_{ik}$  be the loss of  $a_i$  when the class value is actually  $y_k$ .

**Expected Risk for Action  $a_i$ :**

$$R(a_i | X) = \sum_{k=1}^K \lambda_{ik} P(Y = y_k | X)$$

**Choose  $a_i$  if:**

$$R(a_i | X) = \min_k R(a_k | X)$$

### 3.4.2 0/1 Loss

$$\lambda_{ik} = \begin{cases} 0 & i = k \\ 1 & i \neq k \end{cases}$$

If all correct decisions have no loss and all errors are equally costly:

$$\begin{aligned} R(a_i | X) &= \sum_{k=1}^K \lambda_{ik} P(Y = y_k | X) \\ &= \sum_{k \neq i} P(Y = y_k | X) \\ &= 1 - P(Y = y_i | X) \end{aligned}$$

### 3.4.3 Reject or Doubt

If the automatic system has low certainty of its decision and making a wrong decision may have very high cost, a manual decision is required.

The solution is to define an additional action for *reject* or *doubt*,  $a_{K+1}$ , with  $a_i$ ,  $i = 1, \dots, K$ , being the usual actions of predicting on classes  $y_i$ .

The revised 0/1 loss function is now:

$$\lambda_{ik} = \begin{cases} 0 & i = k \\ \theta & i = K + 1 \\ 1 & i \neq k \end{cases}$$

where  $0 < \theta < 1$  is the loss incurred for choosing the reject action.

The expected risk of taking the reject action:

$$R(a_{K+1} | X) = \sum_{k=1}^K \theta P(Y = y_k | X) = \theta$$

### 3.4.4 Optimal Decision Rule

The following rules are meaningful if  $0 < \theta < 1$ . If  $\theta = 0$ , we will always reject and if  $\theta \geq 1$ , we will never reject.

#### 3.4.4.1 Decision Rule 1

Take action  $a_i$  if the following conditions are true:

$$\begin{aligned} R(a_i | X) &< R(a_k | X), \forall k | k \neq i \\ R(a_i | X) &< R(a_{K+1} | X) \end{aligned}$$

Choose reject if:

$$R(a_{K+1} | X) < R(a_i | X), i = 1, \dots, K$$

#### 3.4.4.2 Decision Rule 2

Predict  $Y = y_i$  if the following conditions are true:

$$\begin{aligned} P(Y = y_i|X) &> R(Y = y_k|X), \forall k|k \neq i \\ P(Y = y_i|X) &> 1 - \theta \end{aligned}$$

Choose reject otherwise.



# Chapter 4

## Naive Bayes Classifier

### 4.1 Bayesian Classifiers

Given a vector of features  $\mathbf{X}$  and a class label  $Y = y_k$ ,  $\mathbf{X}$  and  $Y$  are treated as random variables and their relationship is captured using  $P(Y|\mathbf{X})$  using training data.

A test record  $\mathbf{X} = \mathbf{x}^*$  or  $\mathbf{X}^*$  can be classified by finding the class  $y_i$  that maximizes the posterior  $P(Y|\mathbf{X}^*)$ . Predict  $Y = y_i$  if:

$$y_i = \underset{k}{\operatorname{argmax}} P(Y = y_k|\mathbf{X}^*)$$
$$y_i = \underset{k}{\operatorname{argmax}} P(\mathbf{X}^*|Y = y_k)P(Y = y_k)$$

### 4.2 Math Review - Independence & Conditional Independence

Let  $\mathbf{X}$  and  $\mathbf{Y}$  be two sets of random variables.

The variables in  $\mathbf{X}$  are said to be **independent** of  $\mathbf{Y}$  if the following condition holds:

$$P(\mathbf{X}, \mathbf{Y}) = P(\mathbf{X}|\mathbf{Y}) \times P(\mathbf{Y}) = P(\mathbf{X}) \times P(\mathbf{Y})$$

Let  $\mathbf{X}$ ,  $\mathbf{Y}$  and  $\mathbf{Z}$  be three sets of random variables.

The variables in  $\mathbf{X}$  are said to be **conditionally independent** of  $\mathbf{Y}$ , given  $\mathbf{Z}$ , if the following condition holds:

$$P(\mathbf{X}|\mathbf{Y}, \mathbf{Z}) = P(\mathbf{X}|\mathbf{Z})$$

The conditional independence between  $\mathbf{X}$  and  $\mathbf{Y}$  given  $\mathbf{Z}$  can be used to derive the following:

$$\begin{aligned} P(\mathbf{X}, \mathbf{Y}|\mathbf{Z}) &= \frac{P(\mathbf{X}, \mathbf{Y}, \mathbf{Z})}{P(\mathbf{Z})} \\ &= \frac{P(\mathbf{X}, \mathbf{Y}, \mathbf{Z})}{P(\mathbf{Y}, \mathbf{Z})} \times \frac{P(\mathbf{Y}, \mathbf{Z})}{P(\mathbf{Z})} \\ &= P(\mathbf{X}|\mathbf{Y}, \mathbf{Z}) \times P(\mathbf{Y}|\mathbf{Z}) \\ &= P(\mathbf{X}|\mathbf{Z}) \times P(\mathbf{Y}|\mathbf{Z}) \end{aligned}$$

### 4.3 Naive Bayes Classifier

Assuming that the features in  $\mathbf{X}$  are conditionally independent given the class label:

$$P(\mathbf{X}|Y = y_k) = \prod_{i=1}^d P(X_i|Y = y_k)$$

$$P(X_1, X_2, \dots, X_d|Y = y_k) = \prod_{i=1}^d P(X_i|Y = y_k)$$

To classify a test record  $\mathbf{X}^*$ , the posteriors for each class  $y_k$  need to be computed using:

$$P(Y = y_k|\mathbf{X}^*) = \frac{P(Y = y_k) \prod_{i=1}^d P(X_i^*|Y = y_k)}{P(\mathbf{X}^*)}$$

Since  $P(\mathbf{X}^*)$  is fixed for each  $y_k$ , it is sufficient to choose the class that maximizes the numerator:

$$\underset{k}{\operatorname{max}} P(Y = y_k) \prod_{i=1}^d P(\mathbf{X}_i^*|Y = y_k)$$

#### 4.3.1 Estimate Priors

$$P(Y = y_k) = \frac{|Y = y_k|}{N}$$

where  $N$  is the total number of training examples.

#### 4.3.2 Estimate Conditional Probabilities for Discrete Features

$$P(X_i = x_{ij}|Y = y_k) = \frac{|(X_i = x_{ij}) \cap (Y = y_k)|}{|Y = y_k|}$$

where  $x_{ij}$  is the  $j$ th distinct value of the feature  $X_i$ .

### 4.3.3 Estimate Conditional Probabilities for Continuous Features

- Assume the values of a feature given a specific  $y_k$  follows a Gaussian distribution i.e.  $P(X_i|Y = y_k)$  is a Gaussian distribution.
- Use the training data to estimate the mean ( $\mu$ ) and variance ( $\sigma^2$ ) of the distribution.

$$P(X_i|Y = y_j) = \frac{1}{\sqrt{2\pi\sigma_{ij}^2}} e^{-\frac{(X_i - \mu_{ij})^2}{2\sigma_{ij}^2}}$$

## 4.4 Laplace Estimate & M-Estimate

If one of the conditional probabilities is zero, then the entire expression becomes zero. Therefore, an alternative probability estimation is needed.

**Laplace Estimate:**

$$P(X_i = x_{ij}|Y = y_k) = \frac{|(X_i = x_{ij}) \cap (Y = y_k)| + 1}{|Y = y_k| + n_i}$$

where  $n_i$  is the number of possible values for  $X_i$ .

**M-Estimate:**

$$P(X_i = x_{ij}|Y = y_k) = \frac{|(X_i = x_{ij}) \cap (Y = y_k)| + m \times p}{|Y = y_k| + m}$$

where  $m$  and  $p$  are user-specified parameters based on prior information of  $P(X_i = x_{ij}|Y = y_k)$ .

## 4.5 Notes

- Naive Bayes Classifier is computationally efficient.
- The independence assumption may not hold for some features.
- Correlated features can degrade the performance of Naive Bayes Classifiers.

# Chapter 5

## Bayesian Belief Networks

When two features are correlated, the assumption for Naive Bayes Classifier does not hold true.

$$P(X_1, X_2|Y) \neq P(X_1|Y) \times P(X_2|Y)$$

BBNs provide a more flexible approach for modeling the likelihood probabilities  $P(\mathbf{X}|Y)$ . A Bayesian network provides a graphical representation of the probabilistic relationships among a set of random variables.

There are two key elements:

1. A directed acyclic graph (DAG) encoding the dependence relationships among a set of variables.
2. A probability table associating each node to its immediate parent nodes.

### 5.1 BBN Directed Acyclic Graph

Considering three random variables  $A$ ,  $B$  &  $C$ , where  $A$  and  $B$  are independent variables and each has a direct influence on  $C$ :

- There will be a directed arc from  $A$  to  $C$  and  $B$  to  $C$  in the DAG.
- $A$  &  $B$  are parents of  $C$ .
- $C$  is the child of  $A$  &  $B$ .

### 5.2 Conditional Independence

- A node in a Bayesian network is **conditionally independent** of its non-descendants, if its parents are known.
- A Naive Bayes Classifier can be represented using a Bayesian network where  $Y$  is the parent to all the features in  $\mathbf{X}$ .

### 5.3 BNN Representation

Besides the conditional independence conditions imposed by the network topology, each node is also associated with a

probability table.

- If a node  $X$  does not have any parents, then the table contains only the prior probability  $P(X)$ .
- If a node  $X$  has only one parent  $Z$ , then the table contains the conditional probability  $P(X|Z)$ .
- If a node  $X$  has multiple parents  $\{Z_1, Z_2, \dots, Z_k\}$ , then the table contains the conditional probability  $P(X|Z_1, Z_2, \dots, Z_k)$ .

### 5.4 BNN Model Building

- Step 1: Create the structure of the network.
  - Network topology can be obtained by encoding the subjective knowledge of domain experts;
  - Or can be learned from data (structure learning).
- Step 2: Estimate the probability values in the table associated with each node.

### 5.5 Inference from BNN

Given a BBN and an inference problem:

1. Translate the problem into a probabilistic expression.
2. Based on the property on conditional independence, find all dependent variables to the probabilities being estimated.
3. If the probabilities to be estimated cannot be obtained directly from the probability tables of the BBN, apply the Sum Rule, Product Rule and/or Bayes Theorem to decompose the probabilities until all decomposed probabilities can be obtained from the tables.

### 5.6 Notes

- BBN provides an approach for capturing the prior knowledge of a specific domain using a directed graphical model.
  - Other Examples of Directed Graphical Models: Hidden Markov Models, Dynamic Bayesian Networks etc.
  - Other Examples of Undirected Graphical Models: Markov Random Fields, Condition Random Fields etc.

- Network construction, however, is time consuming as it requires domain knowledge or a system that can learn structure from data.

# Chapter 6

## Decision Tree

A decision tree is used to query particular features at particular points in the tree to come to a decision about the classification of the current  $\mathbf{X}$ . The tree has to be *induced* from the training data and can then be used to *deduce* classifications.

### 6.1 Induction Algorithms

A tree can be learned by splitting training data into subsets based on outcomes of a feature test. This process is recursively applied on each derived subset until the subset at a node has the same value for the target variable or there is no improvement for prediction.

### 6.2 Hunt's Algorithm

Hunt's algorithm grows a decision tree in a recursive fashion by partitioning the training records into successively purer subsets. A pure subset is one where the label (i.e. value of  $Y$ ) is the same for all records.

Let  $D_t$  be the set of training records that reach a node  $t$ .

- If  $D_t$  contains records that belong to the same class  $y_t$ , then  $t$  is a leaf node labelled as  $y_t$ .
- Else, if  $D_t$  is an empty set, then  $t$  is a leaf node labelled by the default class  $y_d$ .
- Else, if  $D_t$  contains records that belong to more than one class of  $Y$ , then a feature is selected to conduct a conditional test to split the data into smaller subsets.
  - A child node is created for each outcome of the test condition and the records in  $D_t$  are distributed to the children based on the outcome.
  - Recursively apply this procedure to each subset.

### 6.3 Specifying the Split

- Splitting based on binary features will split the  $D_t$  into two subsets.
- Splitting based on categorical features can be done in one of two ways:

- A multi-way split uses as many partitions as distinct values.
- A binary split divides  $D_t$  into two subsets for two distinct sets of possible values of the feature.
- Splitting based on continuous features can be done in one of two ways:
  - A binary split divides  $D_t$  into two subsets for two distinct ranges of possible values of the feature. ( $X_j < v$ ) or ( $X_j \geq v$ ).
  - A multi-way split divides  $D_t$  into multiple distinct ranges of possible values of the feature. May be very computationally intensive.

### 6.4 Determining the Best Split

Intuitive Idea: Nodes with **homogeneous** class distribution is preferred.

A split criterion can be defined in terms of the difference in degrees of node impurity before and after splitting. This impurity can be measured using the entropy at a given node  $t$ :

$$\text{Entropy}(t) = - \sum_{y_k} P(Y = y_k | t) \log_2 P(Y = y_k | t)$$

- Maximum:  $\log_2 K$  where  $K$  is the total number of all possible values of  $Y$  i.e. records are equally distributed among all classes i.e. least information.
- Minimum: 0 i.e. when all records belong to one class i.e. most information.

#### 6.4.1 Best Split on Information Gain

Suppose a parent node  $t$  is split into  $P$  partitions, the information gain:

$$\delta_{\text{info}} = \text{Entropy}(t) - \sum_{j=1}^k \frac{n_j}{n} \text{Entropy}(i)$$

where  $n_j$  represents the number of examples at child  $j$  and  $n$  represents the number of examples at node  $t$ .

The feature test that maximizes the information gain must be chosen as it minimizes the weighted average impurity measures of the child nodes.

However, a disadvantage is that splitting based on entropy tends to prefer splits that result in a large number of partitions, each being small but pure.

### 6.4.2 Gain Ratio

$$\delta_{\text{InfoR}} = \frac{\delta_{\text{info}}}{\text{SplitINFO}}$$

$$\text{SplitINFO} = - \sum_{i=1}^k \frac{n_i}{n} \log_2 \frac{n_i}{n}$$

where  $n$  is total number of records in the parent node  $t$  and  $n_i$  is the number of records in partition  $i$ .

By using the gain ratio to adjust information gain, higher entropy partitioning (i.e. large number of small partitions) is penalized.

## 6.5 Stopping Criteria for Tree Induction

- Stop expanding a node when all the records belong to the same class.
- Stop expanding a node when all the records have similar attribute values.
- Use early termination.

## 6.6 Notes

- Decision tree classifiers are easy to interpret and efficient in both, training & testing.
- They are used as a base classifier in many ensemble learning approaches.

# Chapter 7

## Generalization

### 7.1 Underfitting & Overfitting

- Underfitting occurs when the model is too simple, and hence, both training and test error rates are large.
- Overfitting occurs when the test error rate begins to increase even though the training error rate continues to decrease.
  - Overfitting results in decision trees that are more complex than necessary.
  - The training error no longer provides a good estimate of how well the tree will perform on previously unseen records. Therefore, there is a need for new ways to estimate the error rate.

### 7.2 Generalization Error

Generalization error is a measure of how accurately an algorithm is able to predict outcome values of previously unseen data. A model with the ideal complexity is the one that produces the lowest generalization error. Since there is no knowledge of the test data and how well the model will perform, the generalization error of the induced model needs to be estimated.

#### 7.2.1 Occam's Razor

- Given two models of similar generalization errors, one should prefer the simpler model over the more complex model.
- For complex models, there is a greater chance that it was fitted accidentally by errors in data. Therefore, one should include model complexity when evaluating a model.

#### 7.2.2 Pessimistic Error Estimate

- Explicitly compute the generalization error as the sum of training error and a penalty term for model complexity.

$$e'(T) = e(T) + \Omega(T)$$

In a decision tree, we can define a penalty term of  $k > 0$  on each leaf node:

$$e'(T) = e(T) + N \times k$$

where  $N$  is the total number of leaf nodes.

#### 7.2.3 Using a Validation Set

- Divide the original training dataset into two subsets — one for training and the other (i.e. validation set) is for estimating the generalization error.
- The complexity of the best model can be estimated using the performance of the model on the validation set.

### 7.3 Addressing Overfitting

#### 7.3.1 Pre-Pruning (Early Stopping Rule)

- Stop the algorithm before it becomes a fully-grown tree.
- Typical stopping conditions for a node:
  - Stop if all instances belong to the same class.
  - Stop if all the feature values are the same.
- More restrictive stopping conditions for a node:
  - Stop if the number of instances is less than some user-specified threshold.
  - Stop if the class distribution of the instances is independent of the available features.
  - Stop if expanding the current node does not improve impurity measures.

#### 7.3.2 Post-Pruning

- Grow decision to its entirety and trim the nodes in a bottom-up fashion.
- If generalization error improves after trimming, replace sub-tree with a new leaf node. The class label of this leaf node is determined from the majority class of the instances in the sub-tree.

# Chapter 8

## Artificial Neural Networks

The human brain is a densely interconnected network of neurons, connected to others via axons. Axons are used to transmit nerve impulses from one neuron to another. The human brain learns by changing the strength of the synaptic connection between neurons. An Artificial Neural Network is composed of an interconnected assembly of nodes and directed links.

### 8.1 Perceptron

- Every input node is connected via a weighted link to the output node. Weights can be positive, negative or zero (no connection).
- The model is an assembly of interconnected nodes and weighted links.
- Output node first sums up each of its input value according to the weights of its links. The weighted sum is compared against some threshold  $\theta$  and an output is produced based on the sign of the result.

$$\text{sign}(z) \begin{cases} 1 & \text{if } z \geq 0 \\ -1 & \text{otherwise} \end{cases}$$

$$y = \text{sign}\left(\sum_{i=1}^d w_i X_i - \theta\right)$$

Alternatively,

$$y = \text{sign}(\mathbf{w} \cdot \mathbf{x})$$

where:

$$\mathbf{w} = (w_0, w_1, \dots, w_d)$$

$$\mathbf{x} = (X_0, X_1, \dots, X_d)$$

$$w_0 = -\theta \text{ and } X_0 = 1$$

#### 8.1.1 Learning (Gradient Descent Approach)

During training, the weight parameters  $\mathbf{w}$  are adjusted until the outputs of the perceptron become consistent with the true outputs of the training data. The weight parameters  $\mathbf{w}$  are updated iteratively with every training example using the gradient descent approach.

$$\mathbf{w}^{(t+1)} = \mathbf{w}^t - \lambda \frac{\partial E(\mathbf{w})}{\partial \mathbf{w}}$$

where  $\lambda \in (0, 1]$  is the learning rate and  $E(\mathbf{w})$  is the error function to be minimized.

$$\mathbf{w}^* = \min_{\mathbf{w}} E(\mathbf{w})$$

Let  $h_i$  be the predicted output for  $\mathbf{x}_i$  and consider the loss function for each training example as:

$$\begin{aligned} E &= \frac{1}{2} (y_i - h_i)^2 \\ &= \frac{1}{2} (y_i - \text{sign}(\mathbf{w}^{(t)} \cdot \mathbf{x}_i))^2 \end{aligned}$$

The weight update rule can be rewritten as:

$$\begin{aligned} \mathbf{w}^{t+1} &= \mathbf{w}^t - \lambda \frac{\partial E(\mathbf{w})}{\partial \mathbf{w}} \\ &= \mathbf{w}^t - \lambda \frac{\partial E(h)}{\partial h} \frac{\partial h(z)}{\partial z} \frac{\partial z(\mathbf{w})}{\partial \mathbf{w}} \end{aligned}$$

where:

$$h = \text{sign}(z)$$

$$z = \mathbf{w} \cdot \mathbf{x}$$

Thus, after solving:

$$\mathbf{w}^{(t+1)} = \mathbf{w}^{(t)} + \lambda (y_i - h_i) \mathbf{x}_i$$



### 8.1.1.1 Learning Model

- If the prediction is correct,  $(y - h) = 0$ , then the weight remains unchanged i.e.  $\mathbf{w}^{(k+1)} = \mathbf{w}^{(k)}$ .
- If  $y = +1$  and  $h = -1$ , then  $(y - h) = 2$ .
  - The weights of all links with positive inputs need to be updated by increasing their values.
  - The weights of all links with negative inputs need to be updated by decreasing their weights.
- If  $y = -1$  and  $h = +1$ , then  $(y - h) = -2$ .
  - The weights of all links with positive inputs need to be updated by decreasing their values.
  - The weights of all links with negative inputs need to be updated by increasing their weights.

### 8.1.2 Notes

- The decision boundary of a perceptron is a linear hyper-plane.
- The perceptron learning algorithm is guaranteed to converge to an optimal solution for linear classification problems.
- However, if the problem is not linearly separable, the algorithm fails to converge.

## 8.2 Multilayer Neural Networks

- Multilayer ANNs are feed-forward neural networks i.e. the nodes in one layer are only connected to the nodes in the next layer.
- Each node takes in the inputs and passes it through an **integration function** followed by an **activation function**.

### 8.2.1 Integration Functions

Weighted Sum:

$$\sum_{i=1}^d w_i X_i - \theta$$

Quadratic Function:

$$\sum_{i=1}^d w_i X_i^2 - \theta$$

Spherical Function:

$$\sum_{i=1}^d (X_i - w_i)^2 - \theta$$

### 8.2.2 Activation Functions

Sign Function (Threshold Function):

$$a(z) = \text{sign}(z) = \begin{cases} 1 & \text{if } z \geq 0 \\ -1 & \text{if } z < 0 \end{cases}$$

Unipolar Sigmoid Function:

$$a(z) = \frac{1}{1 + e^{-\lambda z}}$$

When  $\lambda = 1$ , it is called the sigmoid function.

### 8.2.3 General Algorithm for Learning

Just like the perceptron, the learning algorithm for multilayer ANNs starts with initializing the weights of the links and adjusting these weights such that the output of the ANN is consistent with the class labels of the training examples.

Loss function for each training instance:

$$E = \frac{1}{2}(y_i - h_i)^2$$

The objective is to find the weights  $w'_i$  that minimize the error function:

$$\mathbf{w}^{(t+1)} = \mathbf{w}^t - \lambda \frac{\partial E(\mathbf{w})}{\partial \mathbf{w}}$$

However, in multilayer ANNs, the errors cannot be computed directly for the hidden nodes.

### 8.2.4 Backpropagation Algorithm

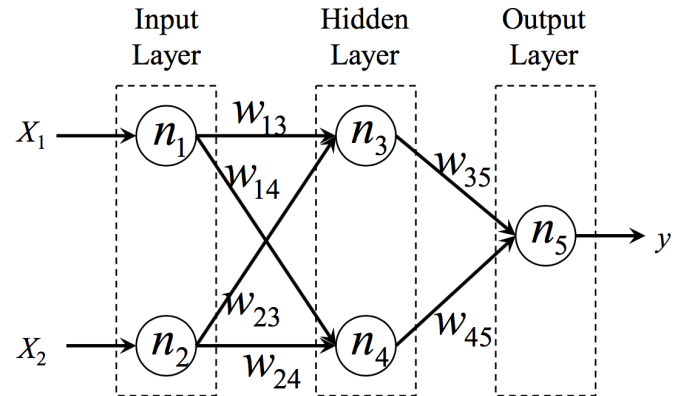


Figure 8.1: Multilayer ANN

After initializing the weights, the training examples are *forward passed* through the neural network to generate outputs.

*Backpropagation* involves starting with the output layer to propagate error back to the previous layer in order to update the weights between the two layers. Since it is not possible to calculate the error in the hidden layers, the error is decomposed using the weights and propagated back when a hidden layer is reached.

For example, in Figure 8.1,  $E_5$  or the error in  $n_5$  will be decomposed using the weights and propagated back. Thus, the value of  $E_3$  will be  $E_5 \times w_{35}$  and the value of  $E_4$  will be  $E_5 \times w_{45}$ .

Revisiting the gradient descent rule:

$$\begin{aligned}\mathbf{w}^{(t+1)} &= \mathbf{w}^{(t)} + \lambda(y_i - h_i) \frac{\partial a(z)}{\partial z} \mathbf{x}_i \\ &= \mathbf{w}^{(t)} + \lambda E_i \frac{\partial a(z)}{\partial z} \mathbf{x}_i \\ &= \mathbf{w}^{(t)} + \lambda \Delta_i \mathbf{x}_i\end{aligned}$$

where  $a(z)$  is the activation function,  $\Delta_i$  is the modified error and  $\mathbf{x}_i$  is the output of the previous layer.

For example, in Figure 8.1, assuming the activation function is the sign function:

$$\frac{\partial a(z)}{\partial z} = 1$$

Then:

$$\mathbf{w}_{35}^{(t+1)} = \mathbf{w}_{35}^{(t)} + \lambda E_5 o_3$$

$$\mathbf{w}_{45}^{(t+1)} = \mathbf{w}_{45}^{(t)} + \lambda E_5 o_4$$

$$\begin{aligned}\mathbf{w}_{13}^{(t+1)} &= \mathbf{w}_{13}^{(t)} + \lambda E_3 o_1 \\ &= \mathbf{w}_{13}^{(t)} + \lambda(\mathbf{w}_{35}^{(t+1)} \times E_5) o_1\end{aligned}$$

$$\begin{aligned}\mathbf{w}_{14}^{(t+1)} &= \mathbf{w}_{14}^{(t)} + \lambda E_4 o_1 \\ &= \mathbf{w}_{14}^{(t)} + \lambda(\mathbf{w}_{45}^{(t+1)} \times E_5) o_1\end{aligned}$$

$$\begin{aligned}\mathbf{w}_{24}^{(t+1)} &= \mathbf{w}_{24}^{(t)} + \lambda E_4 o_2 \\ &= \mathbf{w}_{24}^{(t)} + \lambda(\mathbf{w}_{45}^{(t+1)} \times E_5) o_2\end{aligned}$$

where  $o_i$  is the output of  $n_i$ .

## 8.3 Design Issues for ANN

- The number of nodes in the input layer.
  - Assign an input node to each numerical or binary input variable.
- The number of nodes in the output layer.
  - Binary class problem  $\rightarrow$  single node
  - $k$ -class problem  $\rightarrow k$  output nodes
- Training examples with missing values should be removed or replaced with most likely values.

# Chapter 9

## Support Vector Machines

SVMs are based on statistical learning theory that has shown promising empirical results in many practical applications, such as computer vision, sensor networks and text mining. The basic idea behind SVMs is the **maximum margin hyperplane**.

### 9.1 Maximum Margin

The goal is to learn a binary classifier by finding a hyperplane (decision boundary) such that all the squares reside on one side of the hyperplane and all the circles reside on the other side. Although there are many hyperplanes that can separate the training examples perfectly, their generalization errors may be different.

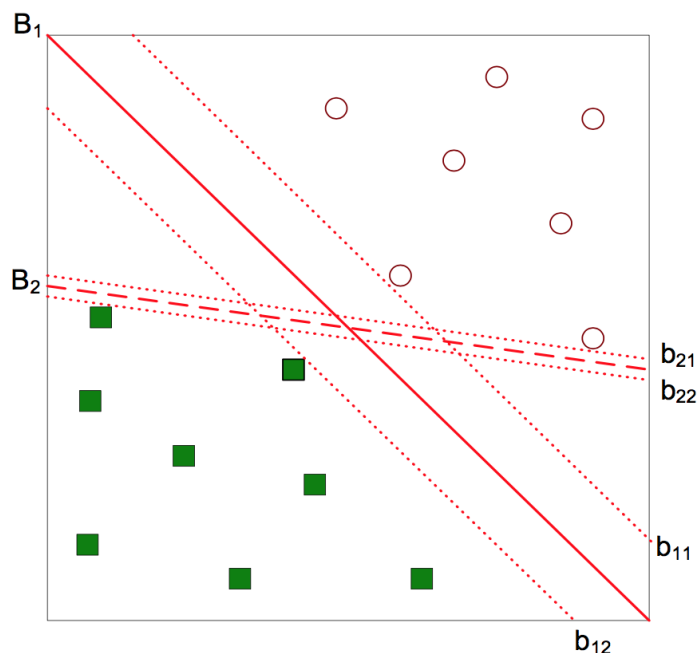


Figure 9.1: Maximum Margin Example (2)

Each decision boundary  $B_1$  is associated with a pair of parallel hyperplanes —  $b_{i1}$  and  $b_{i2}$ .  $b_{i1}$  is obtained by moving the hyperplane until it touches the closest circle(s) and  $b_{i2}$  is

obtained by moving a hyperplane away from the decision boundary until it touches the closest square(s).

The assumption is that larger margins imply better generalization errors. Since the margin of  $B_1$  is larger than the margin of  $B_2$ ,  $B_1$  is better than  $B_2$ .

### 9.2 Math Review

#### 9.2.1 Direction of Vectors

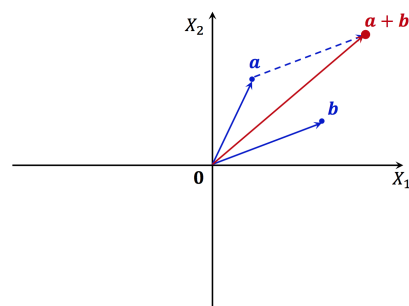


Figure 9.2:  $a + b$

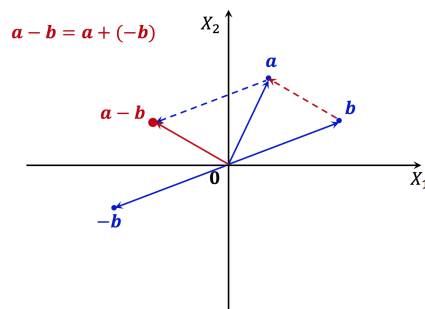


Figure 9.3:  $a - b$

### 9.2.2 Inner Product

The inner product of two vectors,  $\mathbf{u}$  and  $\mathbf{v}$  of  $d$  dimensions is defined as:

$$\mathbf{u} \cdot \mathbf{v} = \sum_{i=1}^d (u_i \times v_i)$$

From a geometry viewpoint:

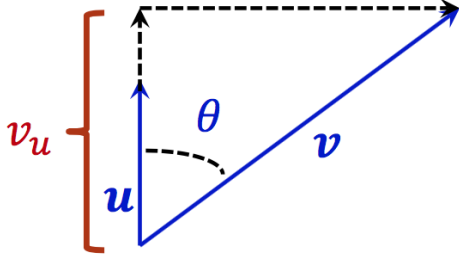


Figure 9.4:  $\mathbf{u}$  &  $\mathbf{v}$

$$\mathbf{u} \cdot \mathbf{v} = \|\mathbf{u}\|_2 \times \|\mathbf{v}\|_2 \times \cos(\theta)$$

$$\|\mathbf{u}\|_2^2 = \mathbf{u} \cdot \mathbf{u} = \sum_{i=1}^d (u_i \times u_i)$$

Thus, when  $\mathbf{u} \cdot \mathbf{v} = 0$ ,  $\theta = 90^\circ$  i.e.  $\mathbf{u}$  and  $\mathbf{v}$  are orthogonal.

The value of  $v_u$  can be derived using the inner product:

$$\begin{aligned} \mathbf{u} \cdot \mathbf{v} &= \|\mathbf{u}\|_2 \times \|\mathbf{v}\|_2 \times \cos(\theta) \\ &= \|\mathbf{u}\|_2 \times (\|\mathbf{v}\|_2 \times \cos(\theta)) \\ &= \|\mathbf{u}\|_2 \times v_u \end{aligned}$$

where  $v_u$  represents the length of  $\mathbf{v}$  in the direction of  $\mathbf{u}$ .

## 9.3 Linear SVM: Separable Case

Given a binary classification task, denoting  $y_i = +1$  as the circle class and  $y_i = -1$  as the square class, the decision boundary is defined as:

$$\mathbf{w} \cdot \mathbf{x} + b = 0$$

For any test example  $\mathbf{x}^*$ :

$$\mathbf{x}^* : \begin{cases} f(\mathbf{x}^*) = +1 & \text{if } \mathbf{w} \cdot \mathbf{x} + b \geq 0 \\ f(\mathbf{x}^*) = -1 & \text{if } \mathbf{w} \cdot \mathbf{x} + b < 0 \end{cases}$$

### 9.3.1 Direction of $\mathbf{w}$

Suppose  $\mathbf{x}_a$  and  $\mathbf{x}_b$  are two points on the decision boundary:

$$\mathbf{w} \cdot \mathbf{x}_a + b = 0$$

$$\mathbf{w} \cdot \mathbf{x}_b + b = 0$$

$$\mathbf{w} \cdot (\mathbf{x}_a - \mathbf{x}_b) = 0$$

Since  $(\mathbf{x}_a - \mathbf{x}_b)$  is a vector on the decision boundary, based on the definition of inner product, the direction of  $\mathbf{w}$  is orthogonal to the decision boundary.

### 9.3.2 Equations for Parallel Hyperplanes (i.e. Support Vectors)

For any circle  $\mathbf{x}_c$  located above the decision boundary:

$$\mathbf{w} \cdot \mathbf{x}_c + b = k, \text{ where } k > 0$$

For any square  $\mathbf{x}_s$  located below the decision boundary:

$$\mathbf{w} \cdot \mathbf{x}_s + b = k', \text{ where } k' < 0$$

The two parallel hyperplanes passing the closest circle(s) and square(s) can be written as:

$$\mathbf{w} \cdot \mathbf{x}_c + b = \bar{k}$$

$$\mathbf{w} \cdot \mathbf{x}_s + b = -\bar{k}$$

$$\text{where } \bar{k} > 0$$

After rescaling  $\mathbf{w}$  and  $b$ , the two parallel hyperplanes can be further rewritten as:

$$\mathbf{w} \cdot \mathbf{x}_c + b = +1$$

$$\mathbf{w} \cdot \mathbf{x}_s + b = -1$$

### 9.3.3 Deriving the Margin

Let  $\mathbf{x}_1$  be a point on  $b_{11}$  and  $\mathbf{x}_2$  be a point on  $b_{12}$ :

$$b_{11} : \mathbf{w} \cdot \mathbf{x}_1 + b = +1$$

$$b_{12} : \mathbf{w} \cdot \mathbf{x}_2 + b = -1$$

$$\mathbf{w} \cdot (\mathbf{x}_1 - \mathbf{x}_2) = 2$$

Based on the definition of inner product:

$$\|w\|_2 \times d = 2$$

$$d = \frac{2}{\|w\|_2}$$

where  $d$  represents the margin i.e.  $(\|x_1 - x_2\|_2 \times \cos(\theta))$ .

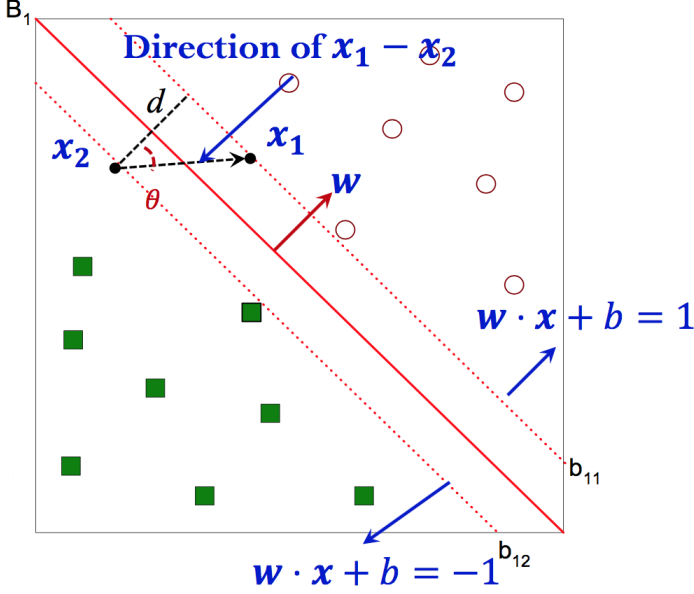


Figure 9.5: Deriving the Margin

### 9.3.4 Learning in Linear SVM

The goal is to maximize the margin  $d$ , or in other words, minimize the following:

$$\min_w \frac{\|w\|_2^2}{2}$$

Given the following constraints:

$$\begin{aligned} w \cdot x_i + b &\geq +1, \text{ if } y_i = 1 \\ w \cdot x_i + b &\leq -1, \text{ if } y_i = -1 \end{aligned}$$

Or, in other words:

$$y_i \times (w \cdot x_i + b) \geq 1$$

Thus, the optimization problem of linear SVM is:

$$\begin{aligned} \min_w \frac{\|w\|_2^2}{2} \\ \text{s.t. } y_i \times (w \cdot x_i + b) \geq 1, i = 1, \dots, N \end{aligned}$$

The optimization is convex and there are many numerical approaches that can be applied to solve it.

## 9.4 Linear SVM: Nonseparable Case

In separable cases, there is no training data within the margin. However, in nonseparable cases, there may be training data that lies within the margin. Thus, slack variables  $\xi \geq 0$  need to be introduced to absorb errors.

$$\begin{aligned} w \cdot x_i + b &\geq +1 - \xi_i, \text{ if } y_i = 1 \\ w \cdot x_i + b &\leq -1 + \xi_i, \text{ if } y_i = -1 \end{aligned}$$

Or, in other words:

$$y_i \times (w \cdot x_i + b) \geq 1 - \xi_i$$

- If  $\xi_i = 0$ , there is no problem with  $x_i$ .
- If  $0 < \xi_i < 1$ ,  $x_i$  is correctly classified but in the margin.
- If  $\xi_i = 1$ ,  $x_i$  is on the decision boundary (random guess).
- If  $\xi_i > 1$ ,  $x_i$  is misclassified.

### 9.4.1 Soft Error

- Number of Misclassifications =  $\#\{\xi_i > 1\}$
- Number of Nonseparable Points =  $\#\{\xi_i > 0\}$

Soft Errors:

$$\sum_{i=1}^N \xi_i$$

Learning with soft errors:

$$\min_w \frac{\|w\|_2^2}{2} + C \left( \sum_{i=1}^N \xi_i \right)$$

$$\text{s.t. } y_i(w \cdot x_i + b) \geq 1 - \xi_i, i = 1, \dots, N, \xi_i \geq 0$$

where  $C \geq 0$  is a parameter to tradeoff the impact of margin maximization and tolerable errors and a nonnegative  $\xi_i$  provides an estimate of the error of the decision boundary on the training example  $x_i$ .

## 9.5 Non-linear SVM

To generalize linear decision boundary to become non-linear,  $x_i$  has to be transformed to a higher dimensional space using a function  $\varphi(x_i)$ . The original input space is mapped to a higher dimensional feature space where the training set is separable.

$$\varphi : \mathbf{x} \rightarrow \varphi(\mathbf{x})$$

The assumption is that in a higher dimensional space, it is easier to find a linear hyperplane to classify the data.

Optimization problem of non-linear SVM:

$$\begin{aligned} \min_w \quad & \frac{\|\mathbf{w}\|_2^2}{2} \\ \text{s.t. } & y_i \times (\mathbf{w} \cdot \varphi(\mathbf{x}_i) + b) \geq 1, i = 1, \dots, N \end{aligned}$$

where  $\mathbf{w} \cdot \varphi(\mathbf{x}_i) + b = 0$  is the hyperplane in feature space.

However, computation in the feature space can be costly because it is high dimensional since the feature space is typically very high dimensional.

### 9.5.1 Kernel Trick

Suppose  $\varphi(\cdot)$  is given as follows, mapping an instance from two-dimensional space to six-dimensional space:

$$\varphi([X_1, X_2]) = [1, \sqrt{2}X_1, \sqrt{2}X_2, X_1^2, X_2^2, \sqrt{2}X_1X_2]$$

Given two data instances,  $\mathbf{a}$  and  $\mathbf{b}$ :

$$\begin{aligned} \varphi(\mathbf{a}) \cdot \varphi(\mathbf{b}) &= 1 + 2A_1B_1 + 2A_2B_2 + A_1^2B_1^2 + A_2^2B_2^2 + 2A_1A_2B_1B_2 \\ &= (1 + A_1B_1 + A_2B_2)^2 \end{aligned}$$

So, if we define the kernel function as follows, there is no need to carry out  $\varphi(\cdot)$  explicitly:

$$\begin{aligned} k(\mathbf{a}, \mathbf{b}) &= (1 + A_1B_1 + A_2B_2)^2 \\ &= (1 + \mathbf{a} \cdot \mathbf{b})^2 \end{aligned}$$

The use of the kernel function to avoid carrying out  $\varphi(\cdot)$  explicitly is known as the **kernel trick**.

Thus, if  $\varphi(\cdot)$  satisfies some conditions, then we can find a function  $k(\cdot, \cdot)$  such that:

$$k(\mathbf{x}_i, \mathbf{x}_j) = \varphi(\mathbf{x}_i) \cdot \varphi(\mathbf{x}_j)$$

#### 9.5.1.1 Well-Known Kernel Functions

Linear Kernel:

$$k(\mathbf{x}_i, \mathbf{x}_j) = \mathbf{x}_i \cdot \mathbf{x}_j$$

Radial Basis Kernel Function w/ Width  $\sigma$ :

$$k(\mathbf{x}_i, \mathbf{x}_j) = \exp\left(-\frac{\|\mathbf{x}_i - \mathbf{x}_j\|_2^2}{2\sigma^2}\right)$$

Polynomial Kernel Function w/ Degree  $d$ :

$$k(\mathbf{x}_i, \mathbf{x}_j) = (\mathbf{x}_i \cdot \mathbf{x}_j + 1)^d$$

### 9.5.2 Lagrange Multiplier Method

In its primal form, it is not possible to solve the optimization problem for non-linear SVMs using the kernel trick as instances in the feature space do not appear in the form of inner products. Thus, its dual form needs to be used.

Using the Lagrange Multiplier Method, we can transform the original optimization problem to its dual form where the mapping function  $\varphi(\mathbf{x})$  does not need to be computed explicitly.

Given an objective function to be minimized, with a set of inequality constraints:

$$\begin{aligned} \min_w \quad & f(\mathbf{w}) \\ \text{s.t. } & h_i(\mathbf{w}) \leq 0, i = 1, 2, \dots, q \end{aligned}$$

The Lagrangian for the optimization problem will be:

$$L(\mathbf{w}, \boldsymbol{\lambda}) = f(\mathbf{w}) + \sum_{i=1}^q \lambda_i h_i(\mathbf{w})$$

where  $\boldsymbol{\lambda} = \{\lambda_1, \dots, \lambda_q\}$  are the Lagrange multipliers.

Thus,

$$\begin{aligned} \min_w \quad & \frac{\|\mathbf{w}\|_2^2}{2} \\ \text{s.t. } & y_i \times (\mathbf{w} \cdot \varphi(\mathbf{x}_i) + b) \geq 1, i = 1, \dots, N \end{aligned}$$

can be transformed to its dual form:

$$\max_{\boldsymbol{\lambda}} L_D(\boldsymbol{\lambda}) = -\left(\frac{1}{2} \sum_{i,j} \lambda_i \lambda_j y_i y_j (\varphi(\mathbf{x}_i) \cdot \varphi(\mathbf{x}_j)) - \sum_{i=1}^N \lambda_i\right)$$

#### 9.5.2.1 Dual Optimization Problem

- The dual Lagrangian involves only the Lagrange multipliers and the training data.
- The negative sign in the dual Lagrangian transforms a minimization problem of the primal form to a maximization problem of the dual form.
- The objective is to maximize  $L_D(\boldsymbol{\lambda})$ .
  - Can be solved using numerical techniques such as quadratic programming.

Once the  $\lambda_i$  values are found, we can obtain the feasible solutions for  $\mathbf{w}$  and  $b$  from the two equations below:

$$\mathbf{w} = \sum_{i=1}^N \lambda_i y_i \varphi(\mathbf{x}_i)$$

$$\lambda_i (y_i (\mathbf{w} \cdot \varphi(\mathbf{x}_i) + b) - 1) = 0$$

The decision boundary can be expressed as:

$$\begin{aligned} \mathbf{w} \cdot \varphi(\mathbf{x}_i) + b &= 0 \\ \left( \sum_{i=1}^N \lambda_i y_i \varphi(\mathbf{x}_i) \cdot \varphi(\mathbf{x}) \right) + b &= 0 \end{aligned}$$

A test instance  $\mathbf{x}^*$  can be classified using:

$$f(\mathbf{x}^*) = \text{sign} \left( \sum_{i=1}^N \lambda_i y_i \varphi(\mathbf{x}_i) \cdot \varphi(\mathbf{x}^*) + b \right)$$

If  $\mathbf{x}_i$  is a support vector, then the corresponding  $\lambda_i > 0$ , otherwise,  $\lambda_i = 0$ .

### 9.5.3 Non-linear SVM via Kernel Trick

Training:

$$\max_{\lambda} \left( \sum_{i=1}^N \lambda_i - \frac{1}{2} \sum_{i,j} \lambda_i \lambda_j y_i y_j (\varphi(\mathbf{x}_i) \cdot \varphi(\mathbf{x}_j)) \right)$$

Decision Boundary:

$$\sum_{i=1}^N \lambda_i y_i (\varphi(\mathbf{x}_i) \cdot \varphi(\mathbf{x}^*)) + b$$

Thus, the data points only appear as inner product in feature space and can be replaced by the kernel function.

Training:

$$\max_{\lambda} \left( \sum_{i=1}^N \lambda_i - \frac{1}{2} \sum_{i,j} \lambda_i \lambda_j y_i y_j k(\mathbf{x}_i, \mathbf{x}_j) \right)$$

Decision Boundary:

$$\sum_{i=1}^N \lambda_i y_i k(\mathbf{x}_i, \mathbf{x}^*) + b$$

If  $\mathbf{x}_i$  is a support vector, then the corresponding  $\lambda_i > 0$ , otherwise,  $\lambda_i = 0$ .

## Chapter 10

# Additional Notes for Multi-Class Classification Problems

Given a 3-class classification problem —  $C_1$ ,  $C_2$  &  $C_3$ , the following binary classifications are performed:

1. Binary Classification 1: Positive ( $C_1$ ) v.s. Negative ( $C_2$  &  $C_3$ )
2. Binary Classification 2: Positive ( $C_2$ ) v.s. Negative ( $C_1$  &  $C_3$ )
3. Binary Classification 3: Positive ( $C_3$ ) v.s. Negative ( $C_1$  &  $C_2$ )

The results of each binary classification need to be combined for a test instance  $\mathbf{x}^*$  to make a final prediction.

- Case 1: Suppose  $f_i$  is a probabilistic model that outputs how likely an instance belongs to class  $C_i$ .
  - The final prediction will be based on which  $f_i$  had the highest probability output.
- Case 2: Suppose  $f_i$  only generates 0/1.
  - A voting table needs to be generated from the results of each  $f_i$  and the class with the most votes is chosen.
  - Each  $f_i$  will either generate a vote for  $C_i$  or a vote for every class except  $C_i$ .



# Chapter 11

## $k$ -Nearest Neighbor Classifier

NN Classifier uses  $k$  *closest* points (nearest neighbors) for performing classification. It requires three things:

1. Set of stored labeled instances.
2. Distance metric to compute the distance between instances.
3. The value of  $k$ , the number of nearest neighbors to retrieve.

To classify an unknown instance:

1. Compute the distance to other training instances.
2. Identify the  $k$  nearest neighbors.
3. Use class labels of nearest neighbors to determine the class label of unknown instances.

### 11.1 Distance Metric

One way to compute the distance between two points is by computing the Euclidean distance:

$$d(\mathbf{x}_i, \mathbf{x}_j) = \sqrt{\sum_{k=1}^d (x_{ik} - x_{jk})^2}$$

To determine the class from the nearest neighbor list, a majority vote of class labels among the  $k$ -nearest neighbors can be used.

$$y^* = \operatorname{argmax}_k \sum_{(\mathbf{x}_i, y_i) \in N_{\mathbf{x}^*}} I(y = y_i)$$

### 11.2 Value of $k$

- If  $k$  is too small, sensitive to noise points.
- If  $k$  is too large, neighborhood may include points from other classes.

### 11.3 Voting Approaches

In simple majority voting, every neighbor has the same impact on the classification which makes the algorithm sensitive to the

choice of  $k$ . The solution is to implement **distance-weight voting**:

$$y^* = \operatorname{argmax}_k \sum_{(\mathbf{x}_i, y_i) \in N_{\mathbf{x}^*}} w_i \times I(y = y_i)$$

where  $w_i$  is the weight according to the distance of the nearest neighbor:

$$w_i = \frac{1}{d(\mathbf{x}^*, \mathbf{x}_i)^2}$$

### 11.4 Normalization

Another issue is that the feature may need to be scaled to prevent the distance from being dominated by some features. For example, the height of a person may vary from 1.5m to 1.8m but the income may vary from \$10K to \$10M. The solution is to normalize the features on different scales.

#### 11.4.1 Min-Max Normalization

$$v_{new} = \frac{v_{old} - \min_{old}}{\max_{old} - \min_{old}} (\max_{new} - \min_{new}) + \min_{new}$$

where *old* represents the original scale and *new* represents the new scale to be normalized to.

#### 11.4.2 Standardization

$$v_{new} = \frac{v_{old} - \mu_{old}}{\sigma_{old}}$$

The above results in  $\mu_{new} = 0$  and  $\sigma_{new} = 1$ .

### 11.5 Notes

- $k$ -NN Classifiers are lazy learners as it does not build models explicitly.

- *Training* is very efficient but classifying unknown test instances is relatively expensive.

# Chapter 12

## Model Evaluation

The most widely-used evaluation metric is:

$$\begin{aligned}\text{Accuracy} &= \frac{\text{Number of Correct Predictions}}{\text{Total Number of Predictions}} \\ &= \frac{\text{TP} + \text{TN}}{\text{TP} + \text{FN} + \text{FP} + \text{TN}}\end{aligned}$$

$$\begin{aligned}\text{Error Rate} &= \frac{\text{Number of Wrong Predictions}}{\text{Total Number of Predictions}} \\ &= \frac{\text{FP} + \text{FN}}{\text{TP} + \text{FN} + \text{FP} + \text{TN}}\end{aligned}$$

However, this metric has limitations. For example, if the training set has 99 examples of Class 0 and 1 example of Class 1, even if the model predicts everything as Class 0, it has an accuracy of 99%, which is misleading.

In many real-world applications, datasets may have imbalanced class distributions. Since accuracy measure treats every class as equally important, it may not be suitable for analyzing imbalanced datasets, where the rare class is considered more interesting than the majority class.

For binary classification, the rare class is often denoted as the positive class while the majority class is denoted as the negative class.

**Precision** and **Recall** are two widely used metrics in applications where successful detection of one of the classes is considered more significant than detection of the other classes. A good model should have both high precision and recall.

### 12.1 Precision

Precision is to measure among all the predicted positive instances, how many are true positive.

$$p = \frac{\text{TP}}{\text{TP} + \text{FP}}$$

### 12.2 Recall

Recall is to measure among all the true positive instances, how many are predicted correctly by the classifier.

$$r = \frac{\text{TP}}{\text{TP} + \text{FN}}$$

### 12.3 $F_1$ -measure

Recall and Precision can be summarized into another metric known as  $F_1$ -measure:

$$F_1 = \frac{2rp}{r + p}$$

A high value of  $F_1$ -measure ensures that precision and recall are reasonably high.

### 12.4 Receiver Operating Characteristic (ROC)

ROC is a graphical approach to displaying the tradeoff between true positive rate and false positive rate of a classifier.

$$\text{True Positive Rate: TPR} = \frac{\text{TP}}{\text{TP} + \text{FN}}$$

$$\text{False Positive Rate: FPR} = \frac{\text{FP}}{\text{TN} + \text{FP}}$$

The ROC curve graphs (FPR, TPR).

- A value of (0, 0) declares everything to be negative class.
- A value of (1, 1) declares everything to be positive class.
- A value of (0, 1) is ideal.
- A value on the diagonal is random guessing.
- A value below the diagonal means the prediction is the opposite of the true class.

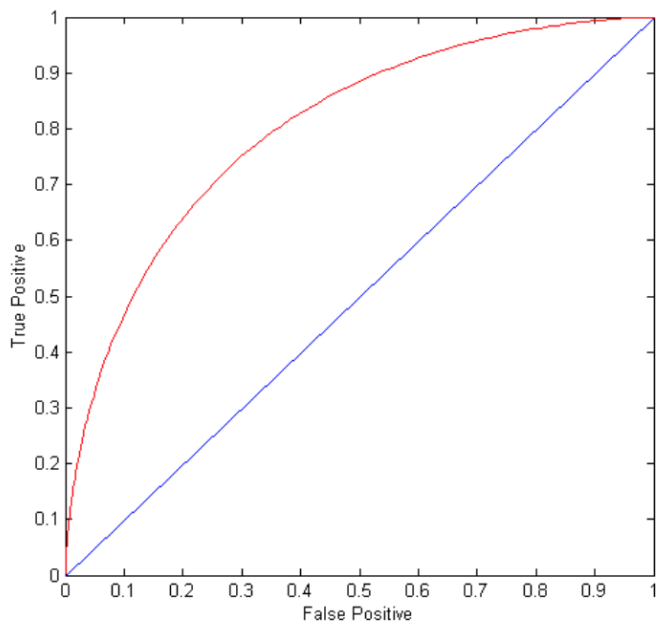


Figure 12.1: ROC Curve

To compare two classifiers using the ROC, the area under the ROC curve must be compared as no classifier consistently outperforms the other. The area under the ROC curve should ideally be 1.

# Chapter 13

## Regression

### 13.1 Linear Regression Model

In a special case where an instance is represented by one input feature, the goal is to learn a linear function  $f(x)$  (where  $x$  is a scalar) in terms of  $w$  (also a scalar) from  $\{x_i, y_i\}$  such that the difference (i.e. error) between the predicted values of  $f(x_i)$ 's and the ground truth values  $y_i$ 's is as small as possible.

$$f(x) = x \times w$$

Suppose sum-of-squares error is used:

$$\begin{aligned} E(w) &= \frac{1}{2} \sum_{i=1}^N (f(x_i) - y_i)^2 \\ &= \frac{1}{2} \sum_{i=1}^N (w \times x_i - y_i)^2 \end{aligned}$$

The linear model is learnt in terms of  $w$  by minimizing the error:

$$w^* = \underset{w}{\operatorname{argmin}} E(w)$$

#### 13.1.1 Linear Regression Model Learning

To solve the unconstrained minimization problem, the derivative of  $E(w)$  with respect to  $w$  can be set to zero.

$$\begin{aligned} \frac{\partial E(w)}{\partial w} &= 0 \\ \frac{\partial \left( \frac{1}{2} \sum_{i=1}^N (w \times x_i - y_i)^2 \right)}{\partial w} &= 0 \\ \sum_{i=1}^N (w \times x_i - y_i) \times x_i &= 0 \\ w \sum_{i=1}^N x_i^2 - \sum_{i=1}^N y_i \times x_i &= 0 \end{aligned}$$

$$w = \frac{\sum_{i=1}^N y_i \times x_i}{\sum_{i=1}^N x_i^2}$$

### 13.2 Math Review

#### 13.2.1 Matrix Transformation

If a matrix/vector  $\mathbf{X}$  is transposed to  $\mathbf{X}^T$ , each element in the matrix/vector switches from  $e_{ij}$  to  $e_{ji}$  where  $i$  is the original row number and  $j$  is the original column number of the element.

- If  $\mathbf{X}$  is a **square matrix**, its number of rows and columns is the same.
- If  $\mathbf{X}$  is a **symmetric matrix**, it is a square matrix and  $\mathbf{X}^T = \mathbf{X}$ .

$$\begin{aligned} \mathbf{X}\mathbf{Y}^T &= \mathbf{Y}^T\mathbf{X}^T \\ \mathbf{X}\mathbf{w}^T &= \mathbf{w}^T\mathbf{X}^T \\ \mathbf{x}^T\mathbf{w}^T &= \mathbf{w}^T\mathbf{x} \\ \mathbf{x}^T &= x \end{aligned}$$

#### 13.2.2 Identity Matrix

- For a square matrix, if  $\mathbf{X}$  is invertible, then  $\mathbf{X}\mathbf{X}^{-1} = \mathbf{I}$ , where  $\mathbf{I}$  is the identity matrix.
- Any vector (or matrix)  $\mathbf{x}$  (or  $\mathbf{X}$ ) times the identity matrix  $\mathbf{I}$  equals to the vector (or matrix) itself.

### 13.3 Linear Regression Model (cont.)

To learn a linear function  $f(\mathbf{x})$  in a more general form in terms of  $\mathbf{w}$  and  $b$ :

$$f(\mathbf{x}) = \mathbf{w} \cdot \mathbf{x} + b$$

By defining  $w_0 = b$  and  $X_0 = 1$ ,  $\mathbf{w}$  and  $\mathbf{x}$  are of  $d + 1$  dimensions:

$$f(\mathbf{x}) = \mathbf{w} \cdot \mathbf{x}$$

Suppose sum-of-squares error is used:

$$\begin{aligned} E(\mathbf{w}) &= \frac{1}{2} \sum_{i=1}^N (f(\mathbf{x}_i) - y_i)^2 \\ &= \frac{1}{2} \sum_{i=1}^N (\mathbf{w} \cdot \mathbf{x}_i - y_i)^2 \end{aligned}$$

Learn the linear model using in terms of  $\mathbf{w}$  by minimizing the error:

$$\mathbf{w}^* = \underset{\mathbf{w}}{\operatorname{argmin}} E(\mathbf{w}) + \frac{\lambda}{2} \|\mathbf{w}\|_2^2$$

where  $\lambda$  is a positive tradeoff parameter and  $\|\mathbf{w}\|_2^2$  is a regularization term to control the complexity of the model.

### 13.3.1 Linear Regression Model Learning

To solve the unconstrained minimization problem, the derivate of  $E(\mathbf{w})$  with respect to  $\mathbf{w}$  can be set to zero.

$$\begin{aligned} \frac{\partial \left( E(\mathbf{w}) + \frac{\lambda}{2} \|\mathbf{w}\|_2^2 \right)}{\partial \mathbf{w}} &= 0 \\ \frac{\partial \left( \frac{1}{2} \sum_{i=1}^N (\mathbf{w} \cdot \mathbf{x}_i - y_i)^2 + \frac{\lambda}{2} \mathbf{w} \cdot \mathbf{w} \right)}{\partial \mathbf{w}} &= 0 \end{aligned}$$

A closed-form solution for the above can be obtained.

Denoting  $\mathbf{X} = (\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_N)^T$ :

$$\mathbf{X} = \begin{pmatrix} x_{10} & \dots & x_{N0} \\ \vdots & \ddots & \vdots \\ x_{1d} & \dots & x_{Nd} \end{pmatrix}^T = \begin{pmatrix} x_{10} & \dots & x_{1d} \\ \vdots & \ddots & \vdots \\ x_{N0} & \dots & x_{Nd} \end{pmatrix}$$

And  $\mathbf{y}$ :

$$\mathbf{y} = \begin{pmatrix} y_1 \\ \vdots \\ y_N \end{pmatrix}$$

Solving the minimization problem:

$$\begin{aligned} \frac{\partial \left( \frac{1}{2} \sum_{i=1}^N (\mathbf{w} \cdot \mathbf{x}_i - y_i)^2 + \frac{\lambda}{2} \mathbf{w} \cdot \mathbf{w} \right)}{\partial \mathbf{w}} &= 0 \\ \sum_{i=1}^N (\mathbf{w} \cdot \mathbf{x}_i - y_i) \mathbf{x}_i + \lambda \mathbf{w} &= 0 \\ \sum_{i=1}^N (\mathbf{w} \cdot \mathbf{x}_i) \mathbf{x}_i - \sum_{i=1}^N y_i \mathbf{x}_i + \lambda \mathbf{w} &= 0 \\ (\mathbf{X}^T \mathbf{X}) \mathbf{w} - \mathbf{X}^T \mathbf{y} + \lambda \mathbf{I} \mathbf{w} &= 0 \\ (\mathbf{X}^T \mathbf{X} + \lambda \mathbf{I}) \mathbf{w} &= \mathbf{X}^T \mathbf{y} \end{aligned}$$

$$\mathbf{w} = (\mathbf{X}^T \mathbf{X} + \lambda \mathbf{I})^{-1} \mathbf{X}^T \mathbf{y}$$

As long as  $\lambda$  is positive,  $(\mathbf{X}^T \mathbf{X} + \lambda \mathbf{I})$  is always invertible.

## 13.4 Linear Basis Function Models

Linear basis function models can be used for non-linear fitting. The linear function  $f(\mathbf{x})$  in terms of a set of basis functions is written as:

$$f(\mathbf{x}) = \mathbf{w} \cdot \boldsymbol{\phi}(\mathbf{x})$$

where  $\boldsymbol{\phi}(\mathbf{x}) = (\phi_0(\mathbf{x}), \phi_1(\mathbf{x}), \dots, \phi_m(\mathbf{x}))$  and each  $\phi_i(\mathbf{x})$  maps the instance  $\mathbf{x}$  to a scalar.

Typically,  $\phi_0(\mathbf{x}) = 1$ , then  $w_0$  acts as a bias.

In the simplest case (if  $d = m$  and  $\phi_i(\mathbf{x}) = x_i$ ), it is reduced to a standard linear model.

### 13.4.1 Examples of Basis Functions

**Polynomial Basis Functions:**

$$\phi_j(\mathbf{x}) = \|\mathbf{x}\|_2^j$$

In a special case  $x$  with one input feature:

$$\phi_j(x) = x^j$$

Then, the polynomial curve will be:

$$f(\mathbf{x}) = \sum_{i=0}^m w_i x^i$$

**Gaussian Basis Functions:**

$$\phi_j(\mathbf{x}) = \exp\left(-\frac{\|\mathbf{x} - \mathbf{u}_j\|_2^2}{2\sigma^2}\right)$$

**Sigmoid Basis Functions:**

$$\phi_j(\mathbf{x}) = g\left(\frac{\|\mathbf{x} - \mathbf{u}_j\|^2}{\sigma}\right)$$

where:

$$g(a) = \frac{1}{1 - \exp(-a)}$$

where  $\mathbf{u}_j$  and  $\sigma$  control location and scale (width).

### 13.4.2 Linear Basis Function Learning

The goal is to minimize the following error:

$$E(\mathbf{w}) = \frac{1}{2} \sum_{i=1}^N (\mathbf{w} \cdot \phi(\mathbf{x}_i) - y_i)^2 + \frac{\lambda}{2} \|\mathbf{w}\|_2^2$$

A closed-form solution for  $\mathbf{w}$  can be obtained:

Denoting  $\Phi = (\phi(\mathbf{x}_1), \phi(\mathbf{x}_1), \dots, \phi(\mathbf{x}_N))^T$ :

$$\begin{aligned} \Phi &= \begin{pmatrix} \phi_0(\mathbf{x}_1) & \dots & \phi_m(\mathbf{x}_1) \\ \vdots & \ddots & \vdots \\ \phi_0(\mathbf{x}_N) & \dots & \phi_m(\mathbf{x}_N) \end{pmatrix}^T \\ &= \begin{pmatrix} \phi_0(\mathbf{x}_1) & \dots & \phi_m(\mathbf{x}_1) \\ \vdots & \ddots & \vdots \\ \phi_0(\mathbf{x}_N) & \dots & \phi_m(\mathbf{x}_N) \end{pmatrix} \end{aligned}$$

The closed-form solution for  $\mathbf{w}$  can be written as:

$$\mathbf{w} = (\Phi^T \Phi + \lambda \mathbf{I})^{-1} \Phi^T \mathbf{y}$$

where:

$$\mathbf{y} = \begin{pmatrix} y_1 \\ \vdots \\ y_N \end{pmatrix}$$

### 13.4.3 Limitations

- In practice, to learn a precise regression model, we may need a large number of basis functions.
- The computational cost is expensive.

### 13.4.4 Kernelized Regression

By using the kernel trick  $k(x_i, x_j) = \phi(x_i) \cdot \phi(x_j)$ :

$$\begin{aligned} f(\mathbf{x}) &= \mathbf{w} \cdot \phi(\mathbf{x}) \\ f(\mathbf{x}) &= \mathbf{k}(\mathbf{x})(\mathbf{K} + \lambda \mathbf{I})^{-1} \mathbf{y} \end{aligned}$$

where:

$$\begin{aligned} \mathbf{K} &= \begin{pmatrix} k(\mathbf{x}_1, \mathbf{x}_1) & \dots & k(\mathbf{x}_1, \mathbf{x}_N) \\ \vdots & \ddots & \vdots \\ k(\mathbf{x}_N, \mathbf{x}_1) & \dots & k(\mathbf{x}_N, \mathbf{x}_N) \end{pmatrix} \\ &= \begin{pmatrix} \phi(\mathbf{x}_1) \cdot \phi(\mathbf{x}_1) & \dots & \phi(\mathbf{x}_1) \cdot \phi(\mathbf{x}_N) \\ \vdots & \ddots & \vdots \\ \phi(\mathbf{x}_N) \cdot \phi(\mathbf{x}_1) & \dots & \phi(\mathbf{x}_N) \cdot \phi(\mathbf{x}_N) \end{pmatrix} \end{aligned}$$

## 13.5 Evaluation

**Root Mean Square Error (RMSE):**

$$\sqrt{\frac{1}{N} \sum_{i=1}^N (f(\mathbf{x}_i) - y_i)^2}$$

**Mean Absolute Error (MAE):**

$$\frac{1}{N} \sum_{i=1}^N |f(\mathbf{x}_i) - y_i|$$

# Chapter 14

## Ensemble Learning

### What?

Ensemble learning refers to a collection of methods that learn a target function by training a number of individual learners and combining their predictions.

### Why?

- Accuracy: a more reliable mapping can be obtained by combining the output of multiple *experts* (i.e. classifiers).
- Efficiency: a complex problem can be decomposed into multiple sub-problems that are easier to understand and solve (divide-and-conquer approach).
- Multiple representations and multiple models: There is not a single model that works for all pattern recognition problems!

### When?

When you can build component classifiers that are more accurate than chance and, more importantly, that are independent from each other.

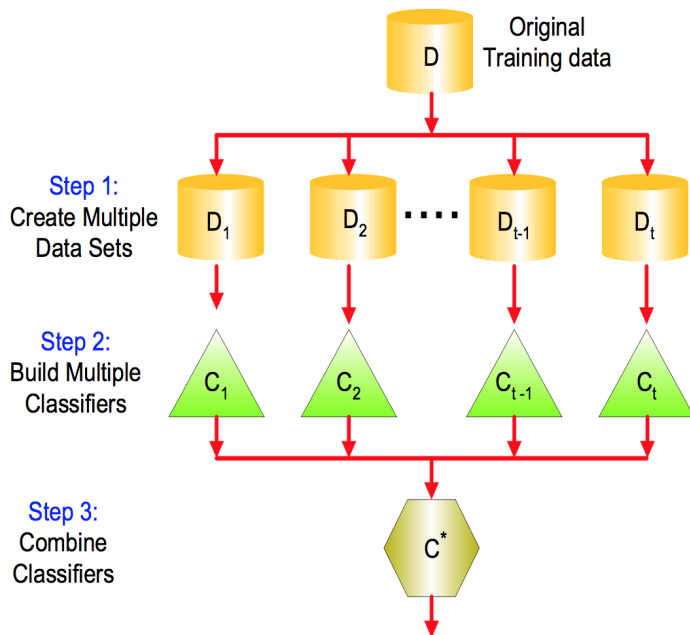


Figure 14.1: Ensemble Learning

## 14.1 Why do ensembles work?

### 14.1.1 Increased Accuracy

If the error of individual classifiers is independent, we can eliminate the error of combined classifier. For example:

Assume a binary classification problem for which you can train individual classifiers with an error rate of 0.3. Assume that you build an ensemble by combining the prediction of 21 such classifiers with a majority vote. If there are  $i$  classifiers out of 21 making an error in prediction, the error of classification is:

$$\binom{21}{i} 0.3^i (1 - 0.3)^{21-i}$$

where:

$$\binom{21}{i} = \frac{21!}{(21-i)!i!}$$

In order for the ensemble to misclassify an example, 11 or more classifiers have to be in error, or a probability of 0.026.

$$\sum_{i=11}^{21} \binom{21}{i} 0.3^i (1 - 0.3)^{21-i} = 0.026$$

### 14.1.2 Approximation by Ensemble Averaging

The desired target function may not be implementable with individual classifiers, but may be approximated by ensemble averaging. For example:

Assume that we use a single discriminant function as the classifier, whose decision boundary is linear. However, the true boundary may be non-linear. If we combine many linear discrimination functions, we can approximate the non-linear boundary.



## 14.2 Methods for Constructing Ensembles

1. Subsampling the training examples
  - a. Multiple hypotheses are generated by training individual classifiers on different datasets obtained by resampling a common training set (Bagging, Boosting).
2. Manipulating the input features
  - a. Multiple hypotheses are generated by training individual classifiers on different representations or different subsets of a common feature vector.
3. Manipulating the output targets
  - a. The output targets for  $C$  classes are encoded with an  $L$ -bit codeword and an individual classifier is built to predict each one of the bits in the codeword.
  - b. Additional “auxiliary” targets may be used to differentiate classifiers.
4. Modifying the learning parameters of the classifier
  - a. A number of classifiers are built with different learning parameters, such as number of neighbors in a  $k$ -Nearest Neighbor rule, initial weights in a Multi-Layer Perceptron etc.

## 14.3 Structure of Ensemble Classifiers

### 14.3.1 Parallel

All the individual classifiers are invoked independently and their results are fused with a combination rule (e.g. average, weighted voting) or a meta-classifier (e.g. stacked generalization).

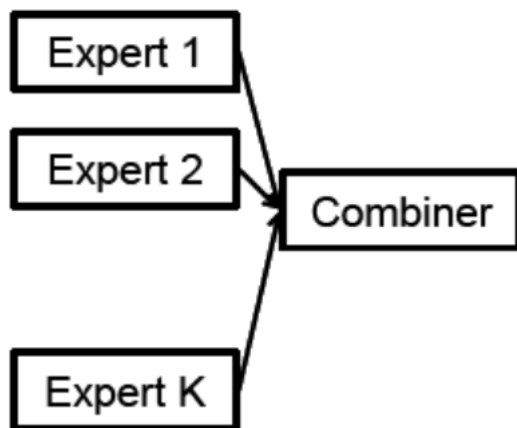


Figure 14.2: Parallel Ensemble Learning

### 14.3.2 Cascading or Hierarchical

Classifiers are invoked in a sequential or tree-structured fashion. For the purpose of efficiency, inaccurate but fast methods are invoked first (maybe using a small subset of the features) and computationally more intensive but accurate methods are left for the latter stages.



Figure 14.3: Cascading Ensemble Learning

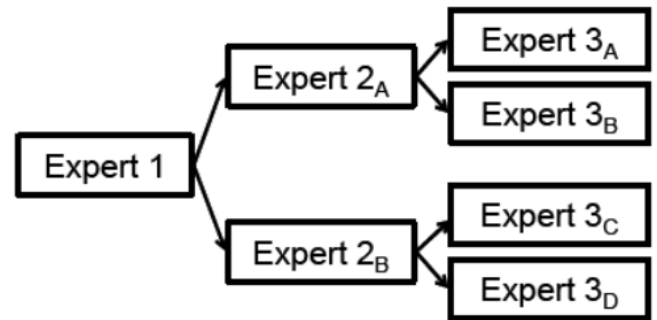


Figure 14.4: Hierarchical Ensemble Learning

## 14.4 Combination Strategies

### 14.4.1 Static Combiners

The combiner decision rule is independent of the feature vector. Static approaches can be broadly divided into non-trainable and trainable.

- Non-trainable: The voting is performed independent of the performance of each individual classifier.
  - Voting: used when each classifier produces a single class label. In this case, each classifier “votes” for a particular class and the class with the majority vote on the ensemble wins.
  - Averaging: used when each classifier produces a confidence estimate (e.g. a posterior). In this case, the winner is the class with the highest average posterior across the ensemble.
- Trainable: The combiner undergoes a separate training phase to improve the performance of the ensemble machine. Two noteworthy approaches are:
  - Weighted Averaging: the output of each classifier is weighted by a measure of its own performance, for example, prediction accuracy on a separate validation set.
  - Stacked Generalization: the output of the ensemble serves as a feature vector to a meta-classifier.

#### 14.4.1.1 Stacked Generalization

In stacked generalization, the output pattern of an ensemble of trained experts serves as an input to a second-level expert.

Training of this modular ensemble can be performed as follows:

1. From a dataset  $X$  with  $N$  examples, leave out one test example and train each of the Level-0 experts on the remaining  $(N - 1)$  examples.
2. Generate a prediction for the test example. The output pattern  $y = [y_1, y_2, \dots, y_K]$  across the Level-0 experts, along with the target  $t$  for the test example, becomes a training example for the Level-1 expert.
3. Repeat this process in a leave-one-out fashion. This yields a training set  $Y$  with  $N$  examples, which is used to train the Level-1 expert separately.
4. To make full use of the training data, re-train all the Level-0 experts one more time using all  $N$  examples in  $X$ .

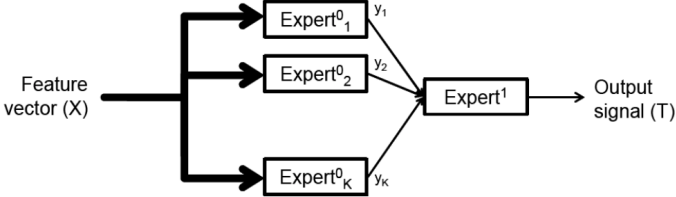


Figure 14.5: Stacked Generalization

#### 14.4.2 Adaptive Combiners

The combiner is a function that depends on the input feature vector. Thus, the ensemble implements a function that is local to each region in the feature space.

This divide-and-conquer approach leads to modular ensembles where relatively simple classifiers specialize in different parts of the input-output space. In contrast with static-combiner ensembles, the individual experts here do not need to perform well for all inputs, only in their region of expertise.

Representative examples of this approach are Mixture of Experts (ME) and Hierarchical ME.

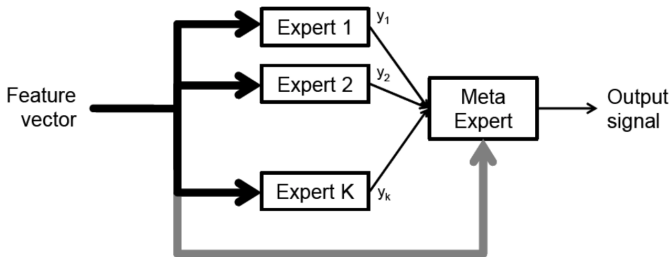


Figure 14.6: Adaptive Learning

#### 14.4.2.1 Mixture of Experts

- A gating network is used to partition the feature space into different regions, with one expert in the ensemble being responsible for generating the correct output within one region.
- The experts in the ensemble and the gating network are trained simultaneously, which can be efficiently performed with the expectation-maximization algorithm.
- ME can be extended to a multi-level hierarchical structure, where each component is itself a ME. In this case, a linear network can be used for the terminal classifiers without compromising the modeling capabilities of the machine.

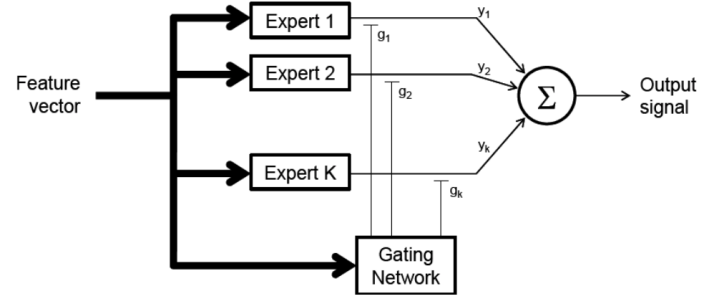


Figure 14.7: Mixture of Experts

### 14.5 Bagging

This method is also known as “arc-ing” (adaptive re-weighting and combining). One of the simplest is “bagging” (bootstrap aggregation).

#### Algorithm:

1. Given  $N$  labeled data points  $\{x_i\}$ , use random sampling (with replacement) to create  $K$  new data sets, each with  $N' < N$ .
  2. Learn a classifier (of whatever type) based on each of the  $K$  data sets giving  $K$  classifiers.
  3. Classify a new point  $x$  based on a majority vote among the  $K$  classifiers.
- The perturbation in the training set due to the bootstrap resampling causes different hypotheses to be built, particularly if the classifier is unstable.
    - A classifier is said to be unstable if a small change in the training data (e.g. order of presentation of examples) can lead to a radically different hypothesis. This is the case of decision trees and, arguably, neural networks.
  - Bagging can be expected to improve accuracy if the induced classifiers are uncorrelated.
    - In some cases, such as  $k$ -Nearest Neighbors, bagging has been shown to degrade performance as compared to individual classifiers as a result of an effectively smaller training set.

## 14.6 Boosting

Boosting takes a different resampling approach than bagging, which maintains a constant probability of  $1/N$  for selecting each individual example. In boosting, this probability is adapted over time based on performance. The component classifiers are built sequentially and examples that are mislabeled by previous components are chosen more often than those that are correctly classified.

Boosting is based on the concept of a “weak learner” (i.e. an algorithm that performs slightly better than chance). An example is a binary classifier with a 50% classification rate. A weak learner can be converted into a strong learner by changing the distribution of the training examples. Boosting can also be used with classifiers that are highly accurate but the benefits in this case will be very small.

A popular variant of boosting is AdaBoost (Adaptive Boosting), which allows the designer to continue adding components until an arbitrarily small error rate is obtained on the training set.

### 14.6.1 AdaBoost

Training:

- For all  $\{x^t, r^t\}_{t=1}^N \in \mathcal{X}$ , initialize  $p_1^t = 1/N$
- For all base-learners  $j = 1, \dots, L$ 
  - Randomly draw  $\mathcal{X}_j$  from  $\mathcal{X}$  with probabilities  $p_j^t$
  - Train  $d_j$  using  $\mathcal{X}_j$
  - For each  $(x^t, r^t)$ , calculate  $y_j^t \leftarrow d_j(x^t)$
  - Calculate error rate:  $\epsilon_j \leftarrow \sum_t p_j^t \cdot 1(y_j^t \neq r^t)$
  - If  $\epsilon_j > 1/2$ , then  $L \leftarrow j - 1$ ; stop
  - $\beta_j \leftarrow \epsilon_j / (1 - \epsilon_j)$
  - For each  $(x^t, r^t)$ , decrease probabilities if correct:
    - If  $y_j^t = r^t$ ,  $p_{j+1}^t \leftarrow \beta_j p_j^t$  Else  $p_{j+1}^t \leftarrow p_j^t$
  - Normalize probabilities:
    - $Z_j \leftarrow \sum_t p_{j+1}^t$ ;  $p_{j+1}^t \leftarrow p_{j+1}^t / Z_j$

Testing:

- Given  $x$ , calculate  $d_j(x), j = 1, \dots, L$
- Calculate class outputs,  $i = 1, \dots, K$ :
$$y_i = \sum_{j=1}^L \left( \log \frac{1}{\beta_j} \right) d_{ji}(x)$$

Figure 14.8: AdaBoost Algorithm

# Chapter 15

## Cluster Analysis

### 15.1 What is Cluster Analysis?

Finding groups of objects such that the objects in a group will be similar (or related) to one another and different from (or unrelated to) the objects in the other groups. Intra-cluster distances are minimized whereas inter-cluster distances are maximized.

#### 15.1.1 Measurements & Distances

Any technique for clustering needs a measure of distance between items. The distance  $d_i$  between two objects  $x_i$  and  $x_j$  can be calculated using the Euclidean distance between two points of  $p$  dimensions.

$$d_{ij} = \sqrt{(x_{i1} - x_{j1})^2 + (x_{i2} - x_{j2})^2 + \dots + (x_{ip} - x_{jp})^2}$$

The Euclidean distance is an example of a dissimilarity measure. However, it may be more convenient to work with a similarity measure, such as the square of the correlation coefficient:

$$r_{ij}^2 = \frac{\sum_{m=1}^p (x_{im} - \bar{x}_m)(x_{jm} - \bar{x}_m)}{\sqrt{\sum_{m=1}^p (x_{im} - \bar{x}_m)^2 \sum_{m=1}^p (x_{jm} - \bar{x}_m)^2}}$$

where  $\bar{x}_m$  is the mean of the  $m$  dimension between the two points.

#### 15.1.2 Application of Clustering in Image Segmentation

- There is a need for data reduction i.e. obtaining a compact representation for *interesting* image data in terms of a set of components.
- The aim is to find components that belong together (form clusters).

#### 15.1.3 Types of Clusterings

A **clustering** is a set of clusters. There are two main types of clusterings:

1. Partitional Clustering - non-overlapping subsets, each data object is in exactly one subset.
2. Hierarchical Clustering - nested clusters, organized as a hierarchical tree.

### 15.2 K-means Clustering

K-means clustering is a partitional clustering approach, where each cluster is associated with a centroid (i.e. center point). Each point is assigned to the cluster with the closest centroid and the number of clusters (i.e. K) must be specified.

Algorithm:

1. Select K points as the initial centroids.
2. Repeat:
  - a. Form K clusters by assigning all points to the closest centroid.
  - b. Recompute the centroid of each cluster.
3. Stop when the centroids no longer change.

#### 15.2.1 Further Details

- The initial centroids are often chosen randomly and the clusters produced vary from one run to another.
- The centroid is (usually) the mean of the points in the cluster.
- ‘Closeness’ is measured by Euclidean distance, cosine similarity, correlation etc.
- K-means will converge for common similarity measures and most of the convergence happens in the first few iterations. Often, the stopping condition may be changed to “Stop when relatively few points change clusters”.
- Complexity is  $O(n \times K \times I \times d)$ , where  $n$  = number of points,  $K$  = number of clusters,  $I$  = number of iterations,  $d$  = number of attributes.

## 15.2.2 Problems with Selecting Initial Points

If there are  $K$  ‘real’ clusters, then the chance of selecting one centroid from each cluster is small. The chance is relatively small when  $K$  is large and it is not guaranteed that the centroids will readjust themselves in the right way. To solve this, the following methods can be used:

1. Multiple runs.
2. Use hierarchical clustering to determine initial centroids.
3. Select more than  $K$  initial centroids (that are widely separated) and then select among these initial centroids.

## 15.2.3 Evaluating K-means Clusters

The most common measure is Sum of Squared Error (SSE), where the error for each point is the distance to the nearest cluster.

$$SSE = \sum_{i=1}^K \sum_{x \in C_i} d(m_i, x)^2$$

where  $x$  is a data point in cluster  $C_i$  and  $m_i$  is the representative point for cluster  $C_i$ .

## 15.2.4 Limitations of K-means

K-means has problems when:

1. Clusters are of different sizes or densities.
2. Data contains outliers.
3. The number of clusters (i.e.  $K$ ) is difficult to determine.

## 15.3 Hierarchical Clustering

Hierarchical clustering produces a set of nested clusters organized as a hierarchical tree and can be visualized as a dendrogram.

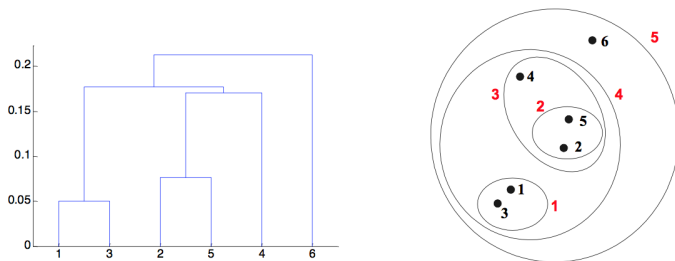


Figure 15.1: Hierarchical Clustering

## 15.3.1 Strengths of Hierarchical Clustering

- There is no assumption about the number of clusters and any desired number of clusters can be obtained by *cutting* the dendrogram at the appropriate level.
- The clustering may correspond to meaningful taxonomies (e.g. animal kingdom).

## 15.3.2 Types of Hierarchical Clustering

1. Agglomerative: Start with the points as individual clusters and at each step, the closest pair of clusters is merged until one cluster (or  $k$  clusters) are left.
2. Divisive: Start with one, all-inclusive cluster and at each step, split a cluster until each cluster contains one point (or there are  $k$  clusters).

## 15.3.3 Agglomerative Hierarchical Clustering

Algorithm:

1. Compute the proximity matrix.
2. Let each data point be a cluster.
3. Repeat:
  - a. Merge the two closest clusters.
  - b. Update the proximity matrix.
4. Stop when only one cluster remains.

The key operation is the computation of proximity between two clusters. The different approaches to calculate this proximity distinguish the different algorithms.

### 15.3.3.1 Inter-Cluster Similarity

1. MIN (Single Linkage): Similarity of two clusters is based on the two most similar (i.e. closest) points in the different clusters. Hence, this inter-cluster similarity is only determined by one pair of points i.e. by one link in the proximity graph.

$$d(A, B) = \min_{a \in A, b \in B} d(a, b)$$

2. MAX (Complete Linkage): Similarity of two clusters is based on the two least similar (i.e. farthest) points in the different clusters. Hence, this inter-cluster similarity is determined by all pairs of points in the two clusters and a cluster is only formed when all points in the cluster are completely linked.

$$d(A, B) = \max_{a \in A, b \in B} d(a, b)$$

3. Group Average: Proximity of two clusters is the average of pairwise proximity between points in the two clusters. There is a need to use average connectivity for scalability since total proximity favors large clusters. This method is less susceptible to noise and outliers but is biased towards globular clusters.

$$d(A, B) = \frac{1}{|A||B|} \sum_{a \in A, b \in B} d(a, b)$$

4. Distance Between Centroids:

$$d(A, B) = d(c_A, c_B)$$

#### 15.3.3.2 Time & Space Complexity

- $O(N^2)$  space complexity due to the proximity matrix (where  $N$  is the number of points).
- $O(N^3)$  time complexity as there are  $N$  steps and at each step,  $N^2$  time is needed to search and update the proximity matrix. In some approaches, the time complexity can be reduced to  $O(N^2 \log(N))$ .