# Machine Learning

Revision



### Definition of Machine Learning

"A computer program is said to **learn** from experience E with respect to some class of tasks T and performance measure P if its performance at tasks in T, as measured by P, improves with experience E."

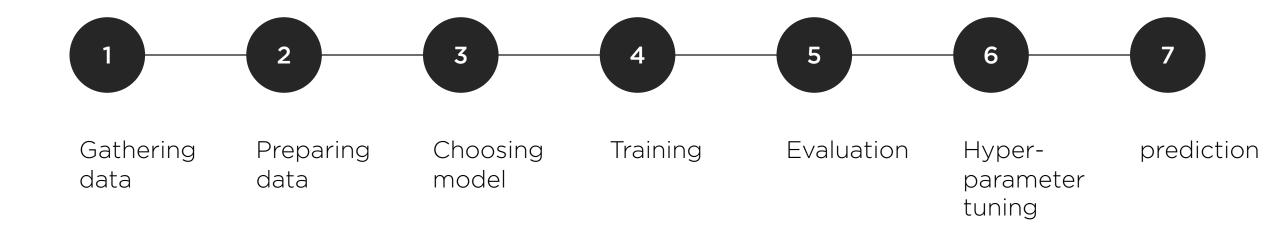
-- Tom M. Mitchell, 1997

Machine learning is the study of algorithms that

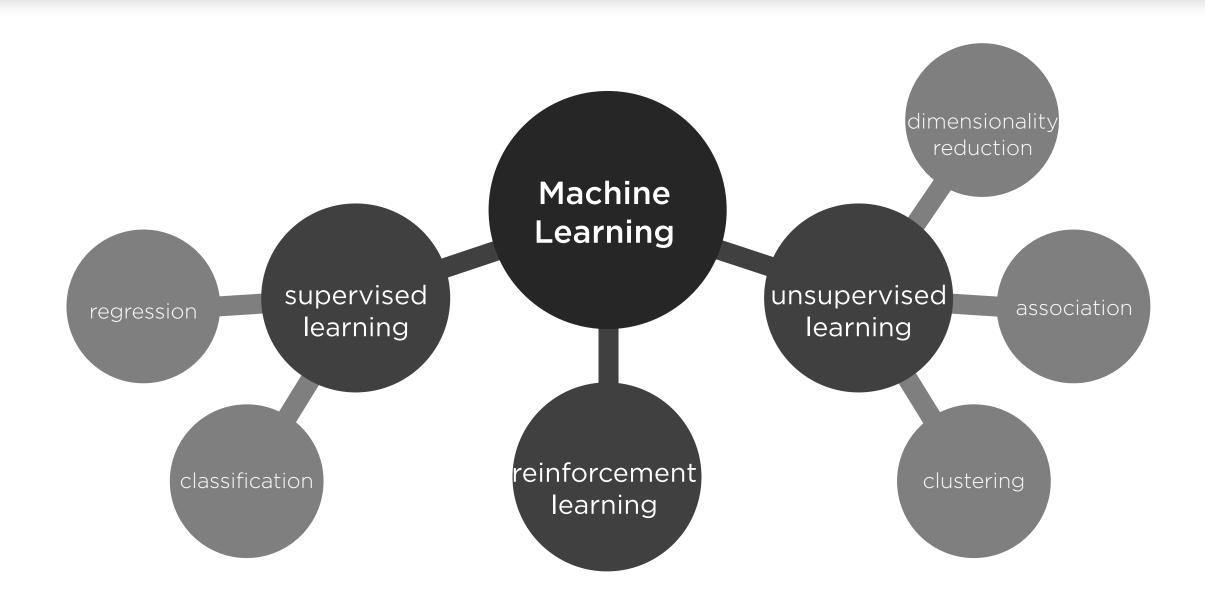
- improve their performance P
- at some task T
- with experience *E*

A well-defined learning task is given by <P, T, E>.

# Machine Learning Lifecycle



## Categories of Machine Learning



# Key Terminologies

- Label is the variable that we are predicting typically represented by the variable y
- Features are input variables that describe our data typically represented by the variables  $\{x_1, x_2, x_3, ..., x_n\}$
- Example is a particular instance of data, x
  - Labelled example has {features, label}: (x, y) used to train the model
  - Unlabelled example has {feature, ?}: (x,?) used for making prediction on new data
- Model maps examples to predict labels:  $\hat{y}$  defined by internal parameters, which are learned
  - Training creating or learning the model.
  - Inference applying the trained model to unlabelled examples.

A method to find the straight line or hyperplane that best fits a set of points.

$$\hat{y} = b + w_1 x_1$$

- $\hat{y}$  the predicted label (a desired output).
- b the bias (the y-intercept), sometimes referred to as  $w_0$ .
- $w_1$  the weight of feature 1 (slope).
- $x_1$  a feature (a known input).

#### Training & Loss

**Training a model**: learning (determining) good values for all weights and the bias from labelled examples.

**Goal of training**: to find a set of weights and biases that have <u>low loss</u>, on average, across all examples.

Loss: the penalty for a bad prediction.

**Empirical Risk Minimisation**: the process of examining many examples and attempting to find a model that minimise loss.

#### Training & Loss

Mean Square Error (MSE) = 
$$\frac{1}{N} \sum_{(x,y) \in D} (y - prediction(x))^2$$

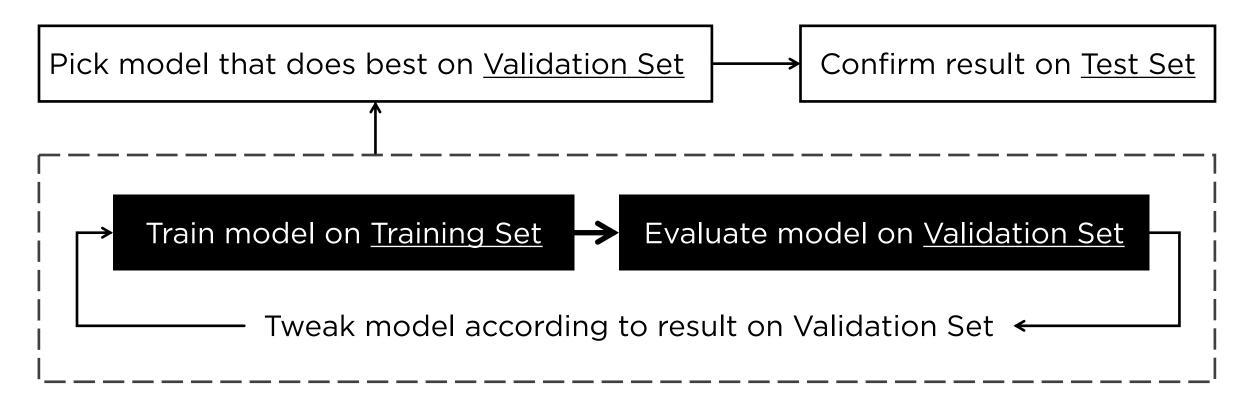
- (x, y) is an example where
  - x is the set of features used by the model to make predictions.
  - y is the example's label
- prediction(x) is a function of the weights & bias in combination with the set of features x.
- D is a dataset containing many labelled examples (x, y) pairs.
- N is the number of examples in D.

#### Reducing Loss

- Hyperparameters are the configuration settings used to tune how the model is trained.
- Derivative of  $(y \hat{y})^2$  with respect to the weights and biases tells us how loss changes for a given example
  - Simple to compute and convex
- So we repeatedly take small steps in the direction that minimises loss
  - We call these Gradient Steps
  - This strategy is called Gradient Descent
- Learning rate  $\frac{1}{f(x)''}$  is one of the hyperparameters of the training process.

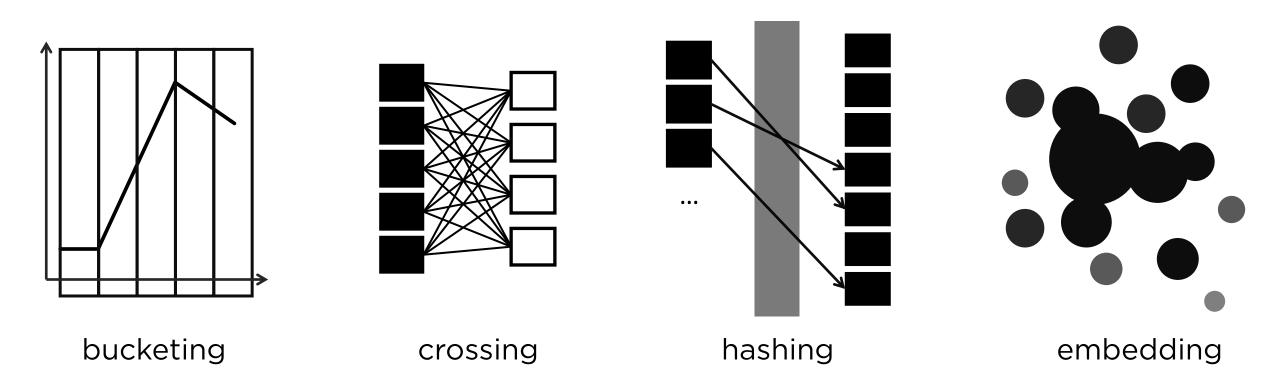
# Training & Test Set

Better Workflow: Use a Validation Set



- Keeping the test data way off to the side (completely unused).
- Pick the model that does best on the validation set.
- Double-check that model against the test set.

## Representation

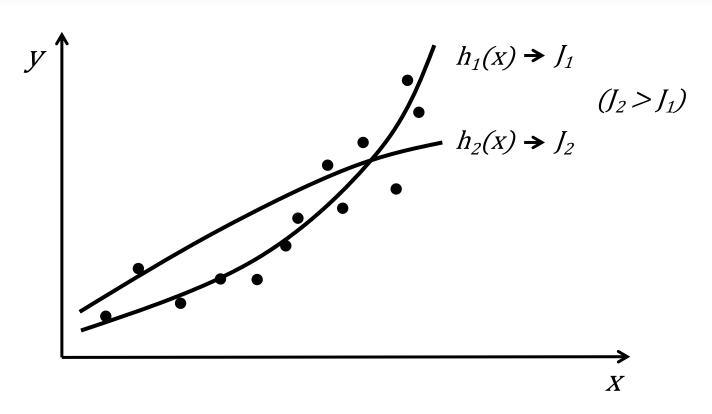


Numeric (Bucketing) / Categorical (Feature Crossing, Hashing, Embedding)

### Cost Functions

 Describes how well the current response surface h(x) fits the available data (on a given data set):

$$J(y_i, h(x_i))$$
observed predicted



- Smaller values of the cost function correspond to a better fit.
- Machine learning goal: construct h(x) such that J is minimised.
- In regression, h(x) is usually directly interpretable as predicted response.

### Cost Functions

• Least Squares Deviation Cost  $J(y_i, h(x_i)) = \frac{1}{n} \sum_{i=1}^{n} (y_i - h(x_i))^2$ 

• Least Absolute Deviation Cost  $J(y_i, h(x_i)) = \frac{1}{n} \sum_{i=1}^{n} |y_i - h(x_i)|$ 

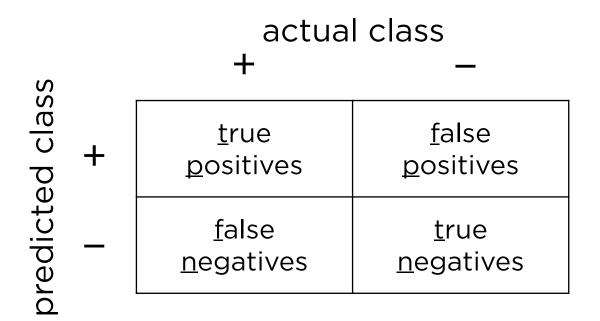
• Huber-M Cost  $J(y_i, h(x_i)) = \frac{1}{n} \sum_{i=1}^{n} \begin{cases} 0.5(y_i - h(x_i))^2, if |y_i - h(x_i)| < \delta \\ \delta(|y_i - h(x_i)| - 0.5\delta), otherwise \end{cases}$ 

# Binary Classifier

- Observed response y takes only two possible values + and -
- Define relationship between h(x) and y
- Use the decision rule:  $\hat{y} = \begin{cases} +, & h(x) \ge t \\ -, & otherwise \end{cases}$

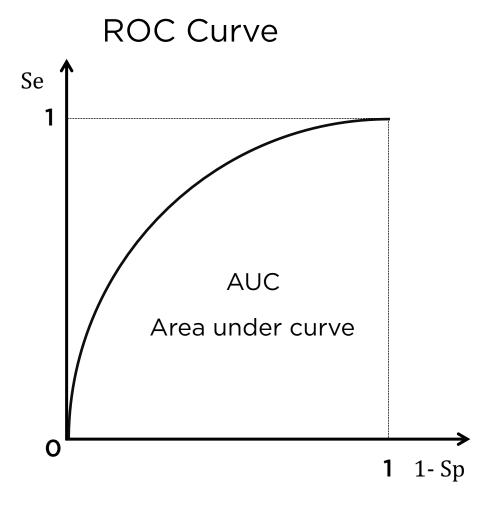
### Performance Measures

#### Prediction Success (Confusion Matrix)



• Precision, Sensitivity (Recall), Specificity

$$SPr = \frac{tp}{tp + fp}$$
  $Se = \frac{tp}{tp + fn}$   $Sp = \frac{tn}{tn + fp}$ 



### Logistic Regression

$$\log(\frac{p}{1-p}) = \log(odds)$$

Input: probability

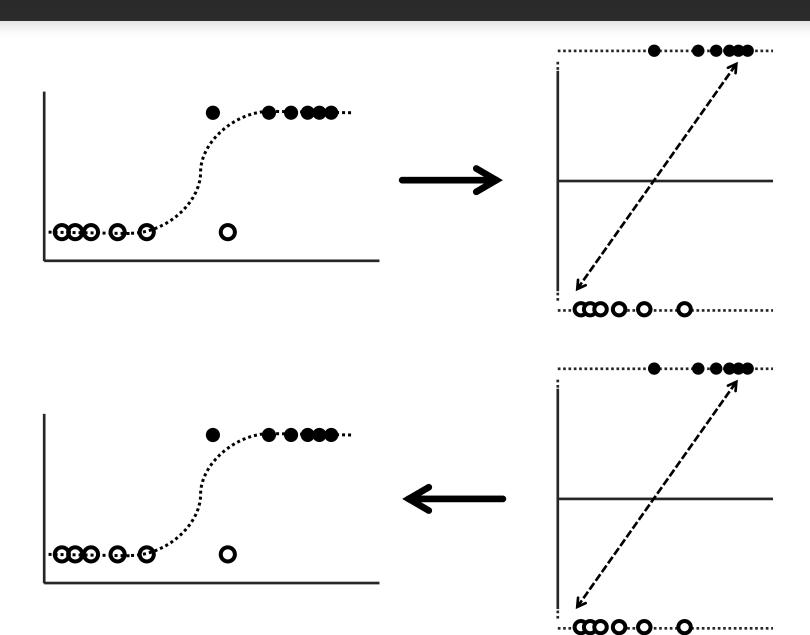
Output: log(odds)

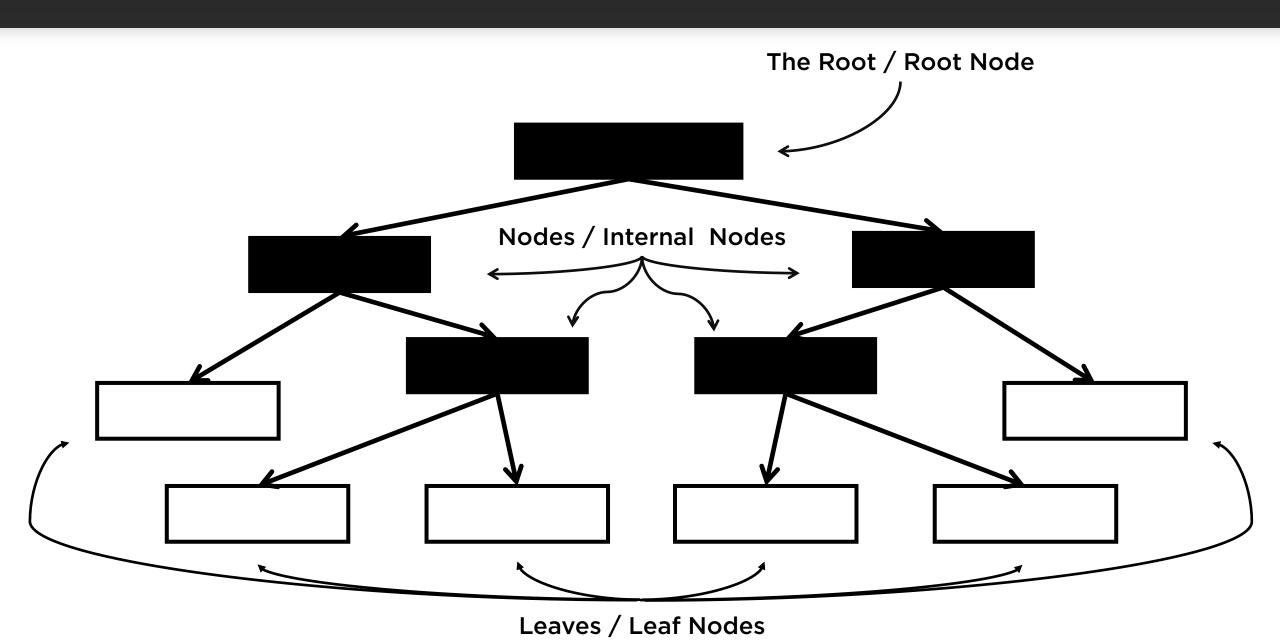
to

$$p = \frac{e^{\log(odds)}}{1 + e^{\log(odds)}}$$

Input: log(odds)

Output: probability

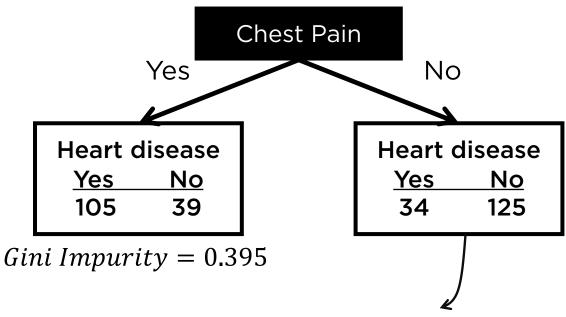




#### Building a Decision Tree

- 1. Calculate all of the Gini impurity values.
- 2. If the node itself has the lowest value, leave it as a Leaf node.
- 3. If separating the data results in an improvement, then pick the separation with the lowest Gini impurity value.

#### Calculating Gini Impurity



Gini Impurity =  $1 - (the probability of Yes)^2 - (the probability of No)^2$ 

$$= 1 - \left(\frac{34}{34 + 125}\right)^2 - \left(\frac{125}{34 + 125}\right)^2$$
$$= 0.336$$

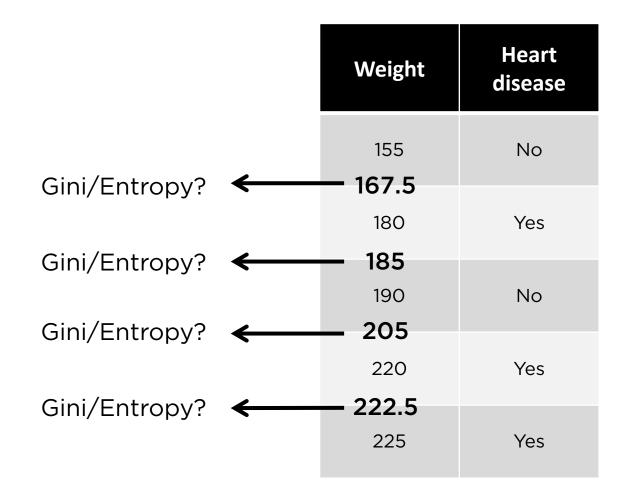
#### Gini Impurity vs Entropy

Gini impurity is a measure of how often a randomly chosen element from the set would be incorrectly labelled if it was randomly labelled according to the distribution of labels in the subset. It measures how heterogeneous some value is over a set. In **decision trees**, we find criteria that make a set into more homogeneous subsets. Information gain is often used to describe this difference that's being maximised though that term goes with **entropy** - entropy is the alternative to Gini impurity.

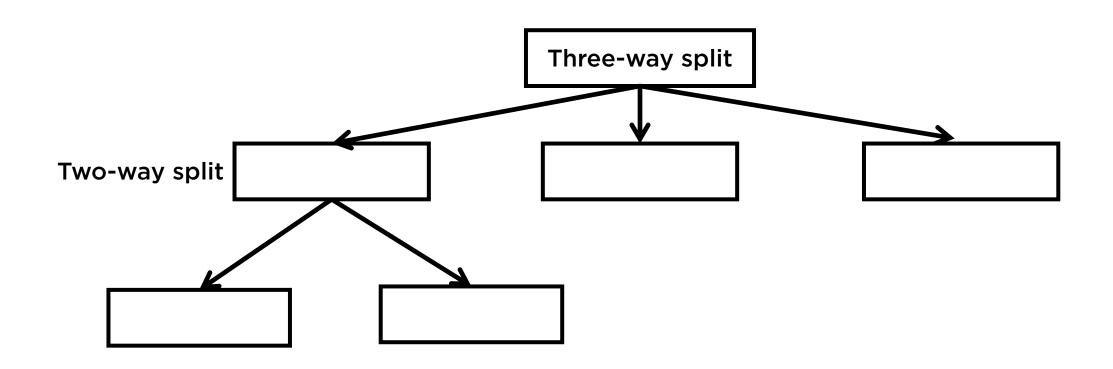
Gini: 
$$Gini(E) = 1 - \sum_{j=1}^{c} p_j^2$$
 Entropy:  $H(E) = -\sum_{j=1}^{c} p_j \log p_j$ 

#### Building a Decision Tree from Numeric Data

- 1. Sort data
- 2. Calculate the average value for all adjacent data
- 3. Calculate the Gini/Entropy values for each average value.
- 4. Find the cutoff to split data



A node can be a two-way (binary) split, a three-way split, and more.



#### Advantages

- Easy to understand
- Easy to generate rules
- Little effort for data preparation
- Less data cleaning required
- Data type is not constraint

#### Disadvantages

- May suffer from <u>overfitting</u>
- Does not easily handle nonnumeric data
- Can be quite large pruning is necessary

### Random Forest

#### Building a Random Forest

- **Step 1**: create a "bootstrapped" dataset
- Step 2: build a Decision Tree using the "bootstrapped" dataset, but only use a random subset of variables
- Step 3: go back to Step 1 and repeat: make a new "bootstrapped" dataset and build a tree considering a subset of variables at each step (ideally 100's of times)

### Random Forest

### Measuring accuracy of a Random Forest

- Measure accuracy of Random Forest is by the proportion of Out-Of-Bag samples that were correctly classified by the Random Forest.
- The proportion of Out-Of-Bag samples that were incorrectly classified is

the "Out-Of-Bag Error"

#### Definition

- More formally, a Support Vector Machine constructs a hyperplane or set of hyperplanes in a high- or infinite-dimensional space, that can be used for classification, regression, or other tasks like outliers detection.
- Intuitively, a good separation is achieved by the hyperplane that has
  the largest distance to the nearest training-data point of any class (socalled functional margin), since in general the larger the margin, the
  lower the generalization error of the classifier.

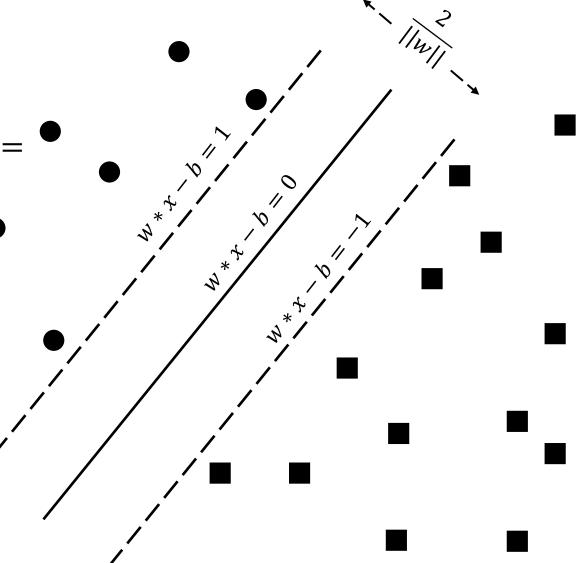
### Support Vector Machine

Linear Separable SVM

• Hyperplane: a set of Hilbert space  $\mathcal{H}_{w,b} = \{x \in \mathbb{R}^d : w^Tx + b = 0\}$  parameterised by  $w \in \mathbb{R}^d$  and  $b \in \mathbb{R}$ .

• Data are **linearly separable**, if there exist  $w \in \mathbb{R}^d$  and  $b \in \mathbb{R}$  such that  $y_i(w^Tx_i+b)>0, \ i=1,...,m$ 

• Then,  $\mathcal{H}_{w,b}$  is a separating hyperplane.



#### Linear Separable SVM

The **distance**  $\rho_x(w,b)$  of a point x from a hyperplane  $\mathcal{H}_{w,b}$  is:

$$\rho_{\mathcal{X}}(w,b) = \frac{|w^T x + b|}{\|w\|}$$

If  $\mathcal{H}_{w,b}$  separates the training set S we define its **margin** as:

$$\rho_{\chi}(w,b) = \min_{i=1:m} \rho_{\chi_i}(w,b)$$

If  $\mathcal{H}_{w,b}$  is a hyperplane (separating or not) we also define the margin of a point x as  $w^Tx + b$  (note that this can be positive )

#### Linear Separable SVM - Hard Margin SVM

- If the training data is linearly separable, we can select two parallel
  hyperplanes that separate the two classes of data, so that the distance
  between them is as large as possible.
- The region bounded by these two hyperplanes is called the "margin", and the maximum-margin hyperplane is the hyperplane that lies halfway between them.

#### Linear Separable SVM - Soft Margin SVM

If the data is not linearly separable:

Minimise 
$$\frac{1}{2}w^Tw + C\sum_{i=1}^m \xi_i$$

Subject to 
$$y_i(w^Tx_i + b) \ge 1 - \xi_i, \ \xi_i \ge 0, \ i = 1, ..., m$$

The idea is to introduce the **slack variable**  $\xi_i$  to relax the separation constraints ( $\xi_i > 0$ )

Linear Separable SVM - Soft Margin SVM

The role of the parameter C (slack penalty, a regularisation parameter)

- Small C allows constraints to be easily ignored, hence results in large margin.
- Large C makes constraints hard to ignore, hence results in narrow margin.
- When  $C = \infty$ , it enforces all constraints to become hard margin problem.

#### Motivation

- Explores the unknown natures of the data that are integrated with little or no prior information
- Saves effort of data labelling, which can be extremely expensive and time consuming
- Provides a compressed representation of the data and is useful in largescale data analysis

#### K-means

#### Optimisation algorithm:

- 1. Initialise  $\{m{\mu}_k\}$ , k $\epsilon$ 1, ..., K, then keep  $m{\mu}_k$  fixed and minimise E with respect to  $r_{nj}$
- 2. Assign  $r_{nj}$  to each data point  $x_n$  based on equation (2), then keep  $r_{nj}$  fixed and minimise J with respect to  $\mu_k$
- 3. Repeat step 1 and 2 until convergence

#### K-means

Estimating the number of clusters - Silhouette analysis

- Study the separation distance between the resulting clusters
- The silhouette plot displays a measure of how close each point in one cluster is to points in the neighbouring clusters
- This measure has a range of [-1, 1].

#### K-means

- Initialise cluster means and assign data points to clusters
- Iteratively repeat the two phases computation:
  - Re-assigning data points to clusters
  - Re-computing the cluster means
- Until there is no further change in the assignments (or until some maximum number of iterations is exceeded)

K-means

#### Issues

- Convergence
- The number of K
- Robustness
- Extension of the Definition of Means

# Good Luck!