

Lecture 10:

Regression models

Outline

- Motivation
- Correlated vs. uncorrelated variables
- Correlation coefficient
- Linear regression
- Nonlinear models (regression trees, RBF networks)

Motivation

Problem: Let us suppose that we know some information about a car (e.g. cylinders, horsepower, weight, acceleration, model etc) and we would like to estimate the fuel consumption (e.g. expressed as miles per gallon)

Example [autoMpg.arff from <http://archive.ics.uci.edu/ml/datasets.html>]

@relation autoMpg

@attribute cylinders { 8, 4, 6, 3, 5} @attribute displacement real

@attribute horsepower real @attribute weight real @attribute acceleration real

@attribute model { 70, 71, 72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82}

@attribute origin { 1, 3, 2}

@attribute class real

@data

8,307,130,3504,12,70,1,18

8,350,165,3693,11.5,70,1,15

4,113,95,2372,15,70,3,24

6,198,95,2833,15.5,70,1,22

6,199,97,2774,15.5,70,1,18

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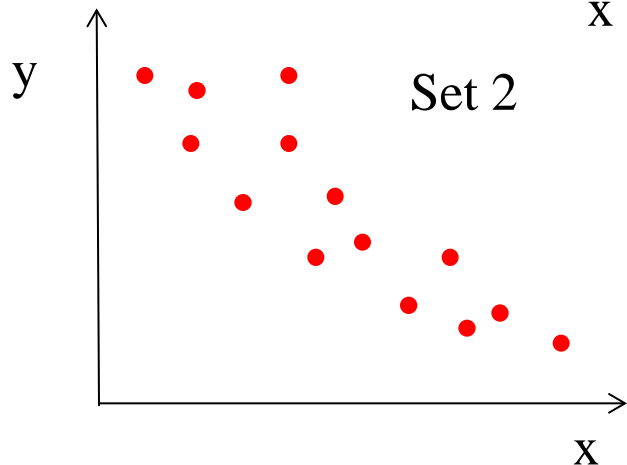
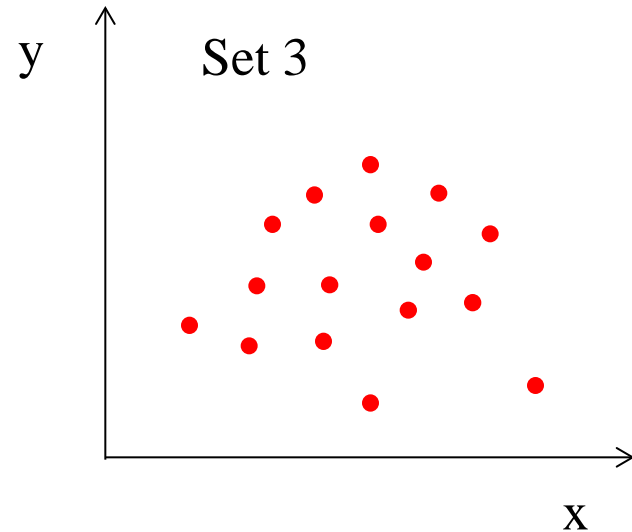
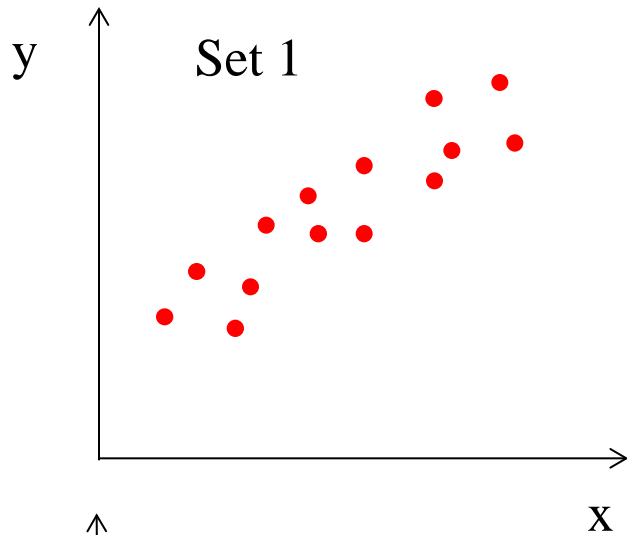
6,198,95,2833,15.5,70,1,22

6,199,97,2774,15.5,70,1,18

We are looking for a dependence between the fuel consumption (class attribute in the dataset) the car characteristics (first 7 attributes in the dataset)

A simpler example

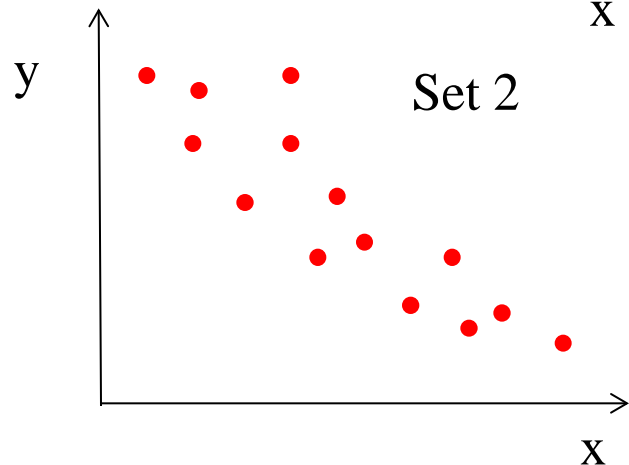
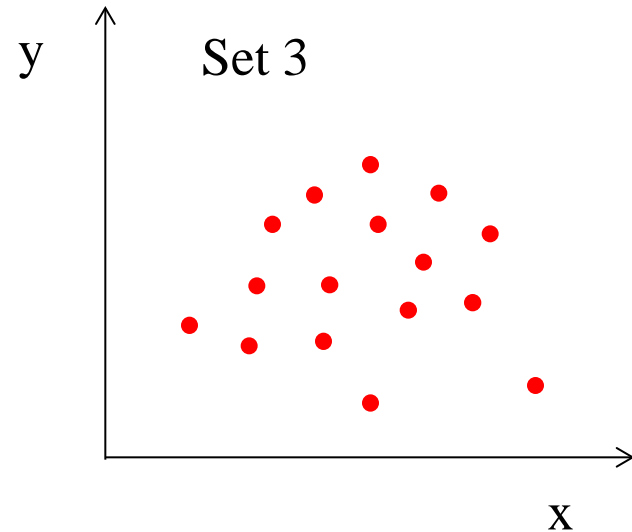
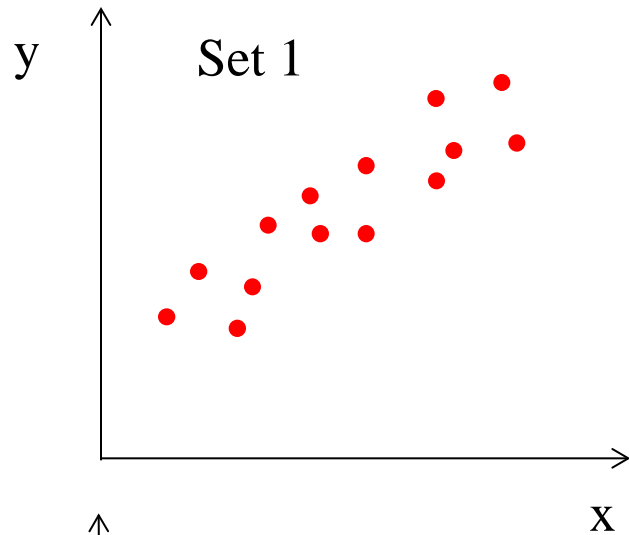
Some synthetic 2D data



What can we say about the data in each set?

A simpler example

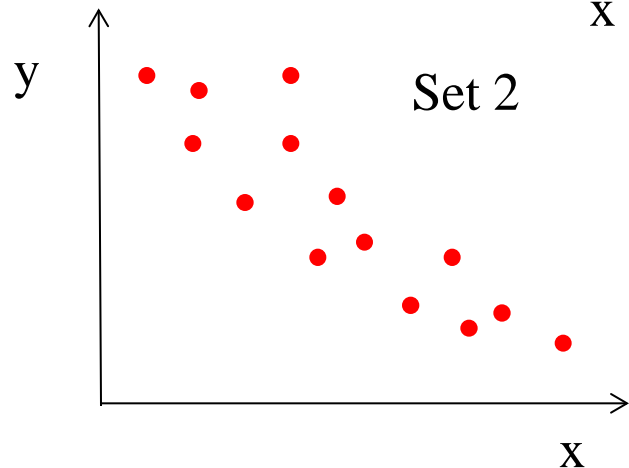
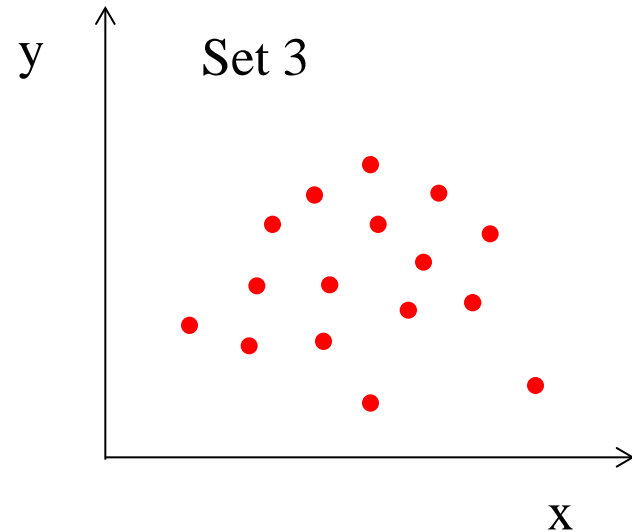
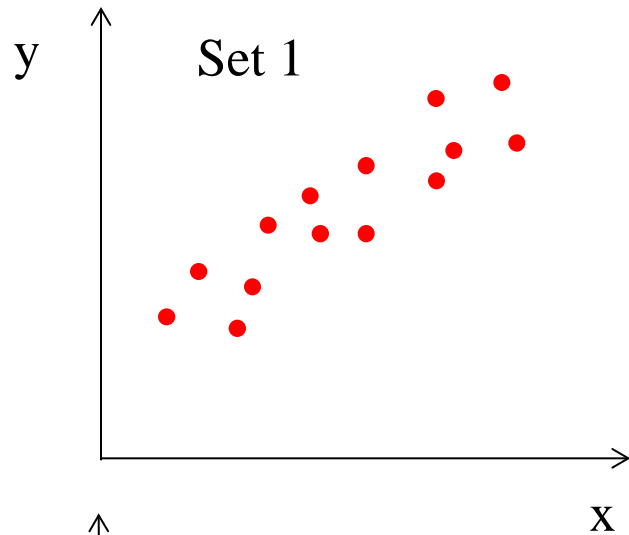
Some synthetic 2D data



Set 1: the data seem to be “positively correlated” = when x increases y also increases

A simpler example

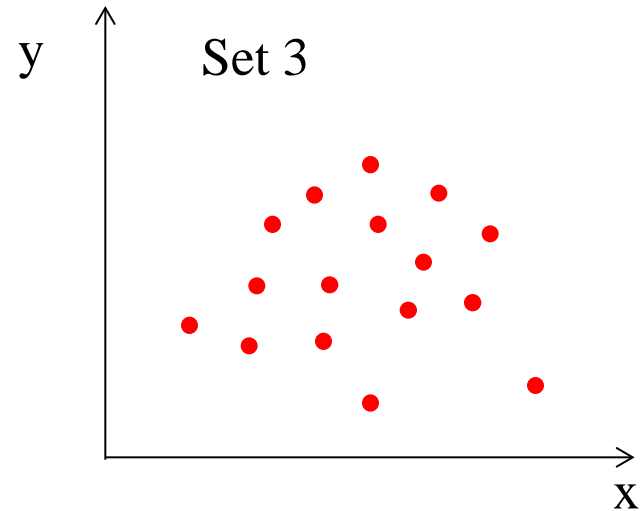
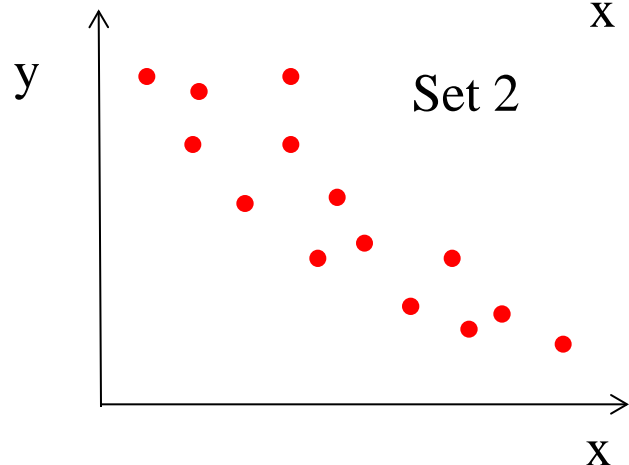
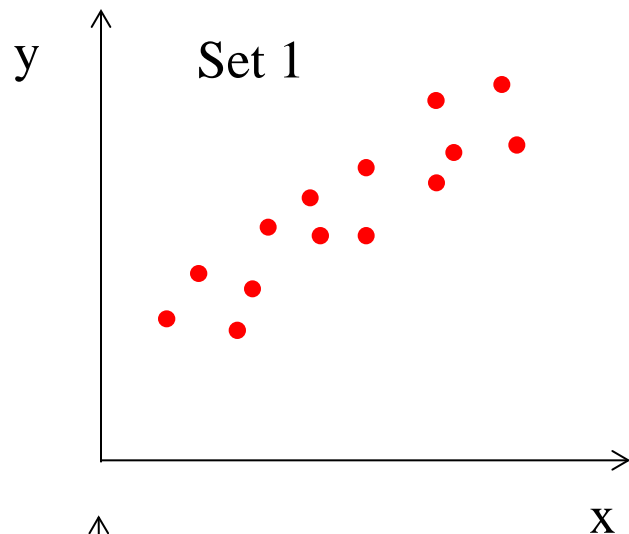
Some synthetic 2D data



Set 2: the data seem to be “negatively correlated” = when x increases y decreases

A simpler example

Some synthetic 2D data



Set 3: the data does not seem to be correlated (it seems to be just a cloud of points)

Questions:

- How can be measured the degree of correlation?
- What kind of correlation?

Correlation coefficient

How can be measured the degree of correlation?

[reminder – Probability and Statistics]

- For instance, by using the **Pearson correlation coefficient** – it expresses the **degree of linear correlation** between two variables

$$R(X, Y) = \frac{\frac{1}{n} \sum_{i=1}^n (x_i - \text{avg}(X))(y_i - \text{avg}(Y))}{\text{stdev}(X)\text{stdev}(Y)}$$

$$\text{stdev}(X) = \sqrt{\frac{1}{n} \sum_{i=1}^n (x_i - \text{avg}(X))^2}$$

$$\text{stdev}(Y) = \sqrt{\frac{1}{n} \sum_{i=1}^n (y_i - \text{avg}(Y))^2}$$

$$\text{avg}(X) = \frac{1}{n} \sum_{i=1}^n x_i, \quad \text{avg}(Y) = \frac{1}{n} \sum_{i=1}^n y_i$$

Remark: $-1 \leq R(X, Y) \leq 1$

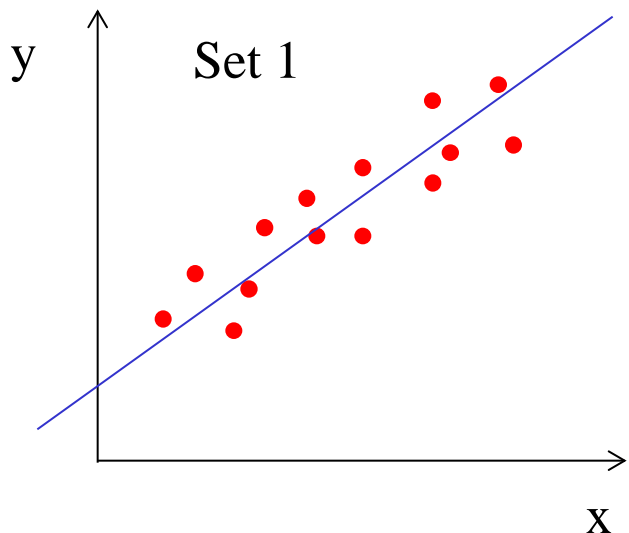
- $R(X, Y)$ close to 1: positive linear correlation
- $R(X, Y)$ close to -1: negative linear correlation
- $R(X, Y)$ close to 0: no linear correlation (however, X and Y could be nonlinearly correlated)

Linear regression

What kind of correlation? [reminder – Probability and Statistics]

Simplest case: **Linear dependence** between two variables: $Y = w_1 X + w_0$

- X = predictor (independent, input, explanatory) variable
- Y = predicted (dependent, response, explained) variable
- Aim of **linear regression**: estimate the parameters w_1 and w_0 such that the available data for the variables X (i.e. x_1, x_2, \dots, x_n) and Y (i.e. y_1, y_2, \dots, y_n) are well explained by the linear function, i.e. the sum of squared errors is minimized



$$SSE(w_1, w_0) = \sum_{i=1}^n (y_i - (w_1 x_i + w_0))^2$$

$$= \sum_{i=1}^n (y_i - \overline{w} x_i)^2$$

$$(\overline{w} = (w_1, w_0), \overline{x}_i = (x_i, 1)^T)$$

row vector

column vector

Simple linear regression

Reminder: some linear algebra

$$w = (w_1, w_0), D = \begin{pmatrix} x_1 & x_2 & \dots & x_n \\ 1 & 1 & \dots & 1 \end{pmatrix}^T, y = (y_1, y_2, \dots, y_n)^T$$

$$\begin{aligned} SSE(w) &= \|y - Dw^T\|^2 = (y - Dw^T)^T (y - Dw^T) \\ &= y^T y - 2wD^T y + wD^T Dw^T \end{aligned}$$

Finding the vector w which minimizes $SSE(w)$ is equivalent with finding the critical point of SSE , i.e. solving the following equation with respect to w :

$$D^T Dw^T = D^T y \Rightarrow w^T = (D^T D)^{-1} D^T y = D^+ y$$

$D^+ = (D^T D)^{-1} D^T$ is called the pseudoinverse of D

Multiple linear regression

Remark: the same approach can be extended in the case when there are d predicting variables (e.g. as in the autoMPG dataset)

$$w = (w_1, w_2, \dots, w_d, w_0), D = \begin{pmatrix} x_{11} & x_{12} & \dots & x_{1n} \\ x_{21} & x_{22} & \dots & x_{2n} \\ \dots & \dots & \dots & \dots \\ x_{d1} & x_{d2} & \dots & x_{dn} \\ 1 & 1 & \dots & 1 \end{pmatrix}^T, y = (y_1, y_2, \dots, y_n)^T$$

$$\begin{aligned} SSE(w) &= \|y - Dw^T\|^2 = (y - Dw^T)^T (y - Dw^T) \\ &= y^T y - 2wD^T y + wD^T Dw^T \end{aligned}$$

$$D^T Dw^T = D^T y \Rightarrow w^T = (D^T D)^{-1} D^T y$$

Linear regression - regularization

Remark: if the matrix $D^T D$ is singular (the inverse cannot be computed) then the objective function (SSE) is modified by adding a so-called **regularization** term which will modify the matrix of the linear system in such a way that it becomes invertible).

Examples:

- Tikhonov regularization (ridge regression)

$$SSE'(w) = SSE(w) + \lambda \|w\|^2$$

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

$I = (d + 1) \times (d + 1)$ identity matrix

Remarks:

- the parameter of the regularization term (**lambda**) is usually chosen adaptively based on cross-validation
- the penalty term “discourages” the large values of the weights

Linear regression - regularization

Remark: if the matrix $D^T D$ is singular (the inverse cannot be computed) then the objective function (SSE) is modified by adding a so-called regularization term which will modify the matrix of the linear system in such a way that it becomes invertible).

Examples:

- Lasso regularization

$$SSE'(w) = SSE(w) + \lambda \sum_{i=1}^d |w_i|$$

(no closed form solution for w)

Remarks:

- In this case the optimization problem is solved by using numerical methods
- Is useful for high dimensional data with many irrelevant features (leading to sparse models)

Generalized linear models

Main idea: instead of $y_i = w_1 x_i + w_0$ the output (y_i) is modelled through a random variable with a distribution having a mean $f(w_1 x_i + w_0)$

Main elements of a GLM (generalized linear model):

- Mean function: f
- Link function: f^{-1}
- Probability distribution

Mean function	Link function	Distribution
$f(u) = u$	identity	normal
$f(u) = -1/u$	inverse	exponential, gamma
$f(u) = \exp(u)$	Log	Poisson
$f(u) = 1/(1 + \exp(-u))$	Logit	Bernoulli

Generalized linear models

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least
squares
regression

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Logistic regression

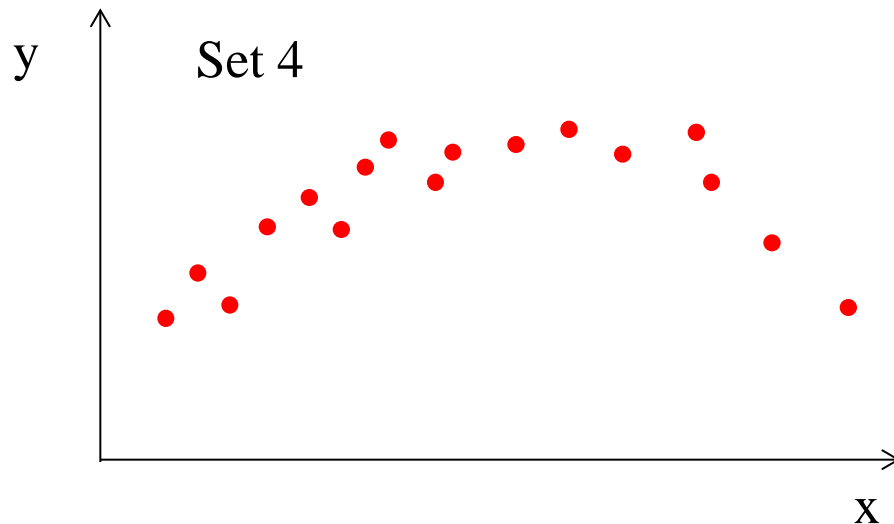
Nonlinear regression

What about the cases when the dependence between the predicted variable and the predictor(s) is not linear?

Other models are needed

Examples:

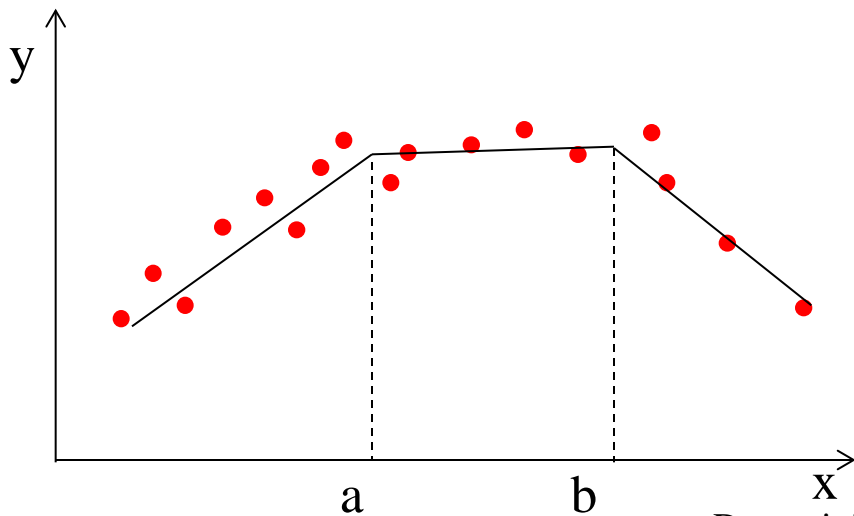
- Regression trees
- Nonlinear neural networks



Nonlinear regression

Main idea:

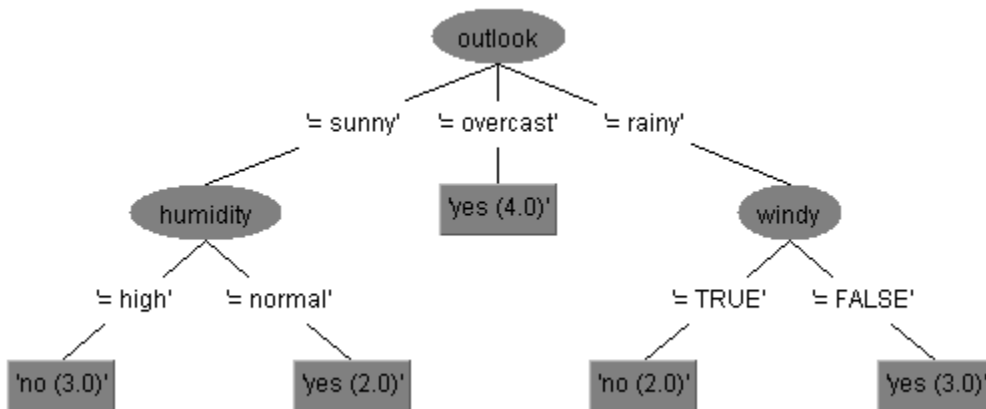
- A nonlinear relationship can be modelled through local linear functions (one linear function per region)
- The regression process would then consist of two steps:
 - Identify the regions by splitting the space of the decision variables
 - Identify a regression model (e.g. a linear one) for each of the identified regions



Regression trees

Reminder:

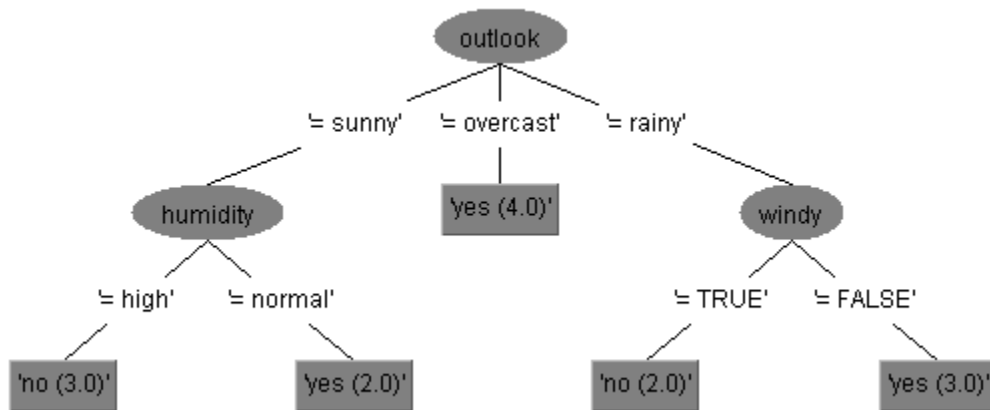
Decision tree = hierarchical structure containing in the internal nodes conditions on the predictor variables and on the leaf nodes information on the predicted variables (e.g. class); if the predicted variable is discrete (categorical/ nominal) then the decision tree is in fact a **classification tree**



Regression trees

Reminder:

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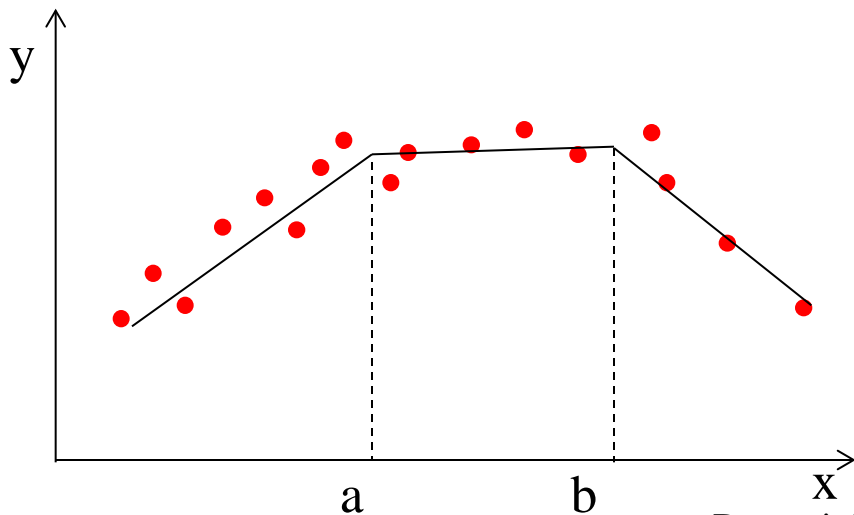
Question:

- What about the case when the predicting variable is continuous? (e.g. we would like to obtain not only a yes/no answer to the “weather-play” problem but a value in $[0,1]$ expressing the degree of decision between 0 (no) and 1 (yes))

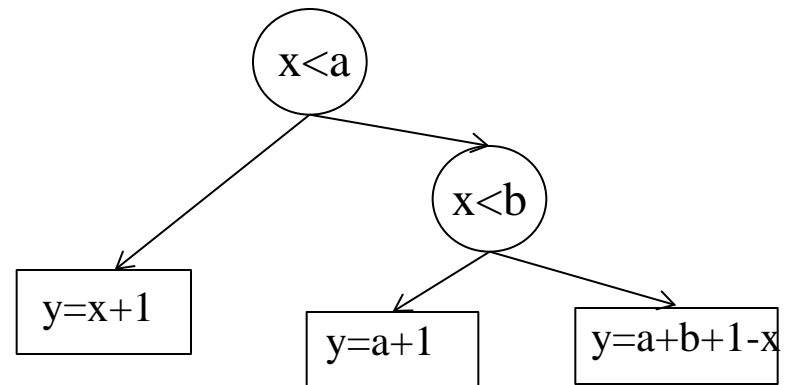
Regression trees

Main idea:

- Use a similar process of splitting the space of the decision (predictor) variables as in the case of trees used for classification
 - In the case of continuous predictor variables the splitting condition is of the one of the types: $\text{variable} < \text{value}$ or $\text{variable} > \text{value}$ or $\text{variable in } [\text{min}, \text{max}]$
- Infer a regression model (e.g. a linear model) for each region identified by the splitting procedure



(Very) simple example -> piecewise linear model:



Nonlinear regression

Beyond piecewise linear models:

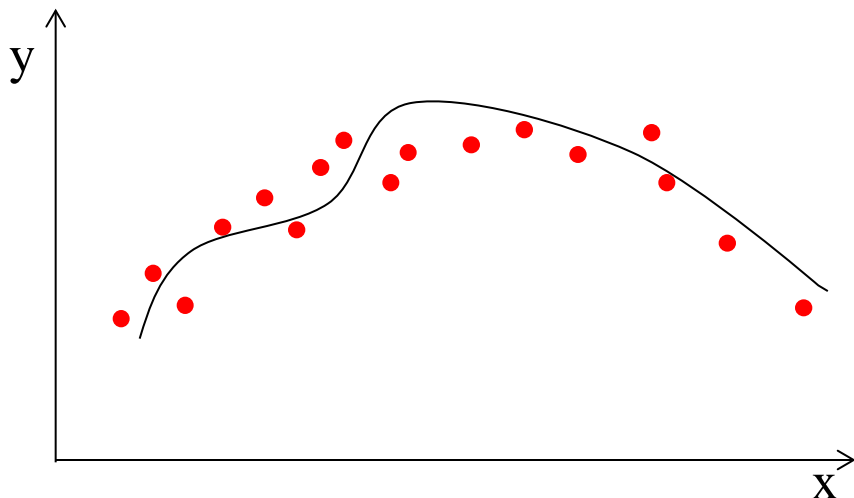
- Extending basic linear regression by using derived input features:

$$y = w_0 + w_1 h_1(x) + w_2 h_2(x) + \dots + w_m h_m(x)$$

(x can be a vector and h_i a function associating a scalar/vector to another vector)

Particular case 1. Polynomial models: $y = w_0 + w_1 x + w_2 x^2 + \dots + w_m x^m$

(x is a scalar)



Particular case 2.

Kernel-based models: h_i are functions which can take significant values only for a limited region of the input space.

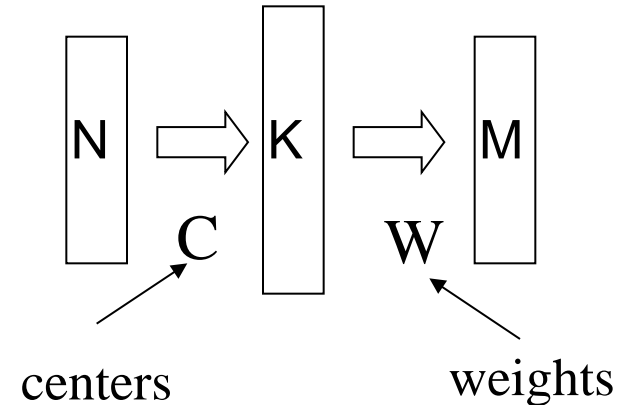
- when these functions are with radial symmetry (e.g. gaussian functions) then we obtain the so-called RBF networks (a particular case of neural networks)

RBF networks

RBF - “Radial Basis Function”:

Architecture:

- Two levels of functional units
- **Aggregation functions:**
 - Hidden units: distance between the input vector and the corresponding center vector
 - Output units: weighted sum



$$G(X, C^k) = \|X - C^k\| = \sum_{i=1}^N (x_i - c_i^k)^2$$

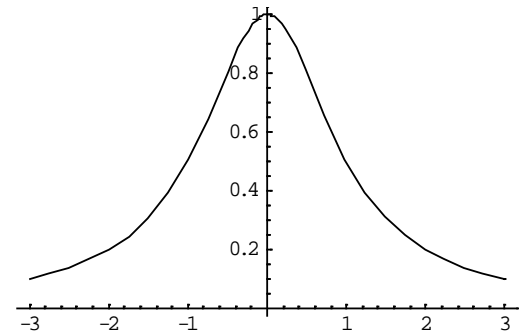
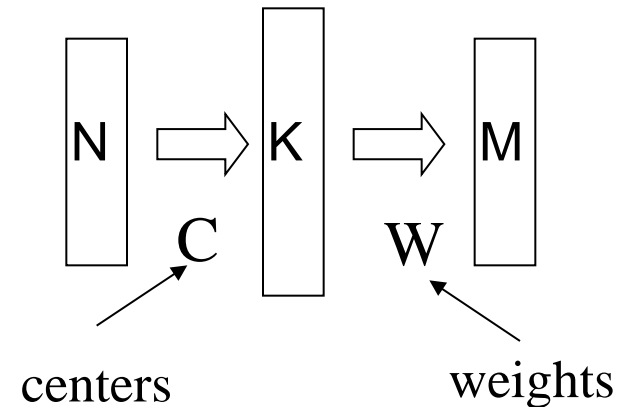
Rmk: hidden units do not have bias values (activation thresholds)

RBF networks

The **activation functions** for the hidden neurons are functions with radial symmetry

- Hidden units generates a **significant output** signal only for input vectors which are **close** enough to the corresponding **center vector**

The activation functions for the output units are usually linear functions



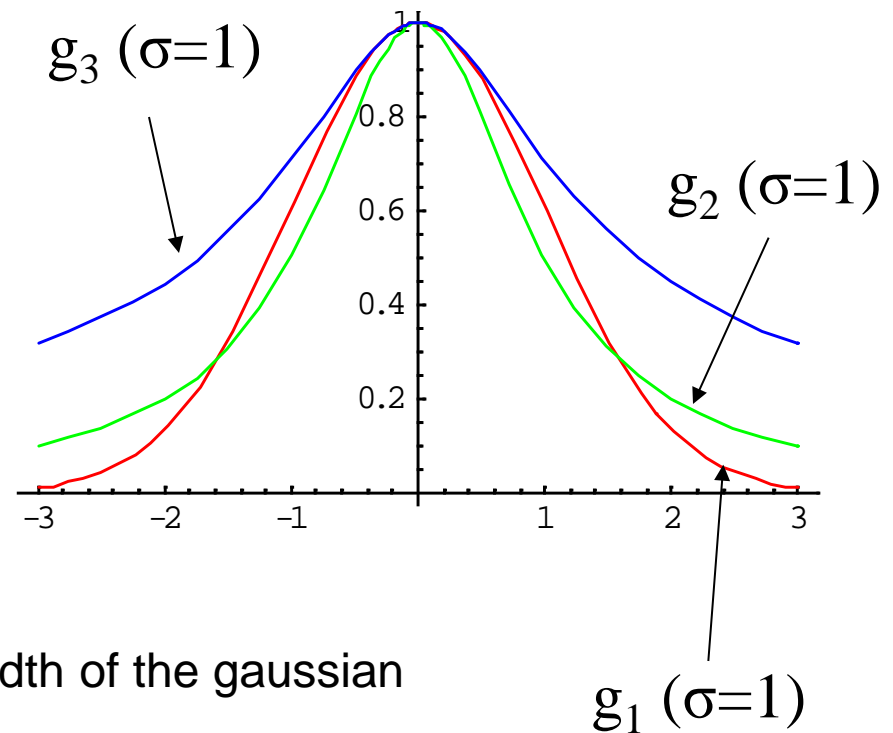
RBF networks

Examples of functions with radial symmetry:

$$g_1(u) = \exp\left(-\frac{u^2}{2\sigma^2}\right)$$

$$g_2(u) = 1/(u^2 + \sigma^2)$$

$$g_3(u) = 1/\sqrt{u^2 + \sigma^2}$$



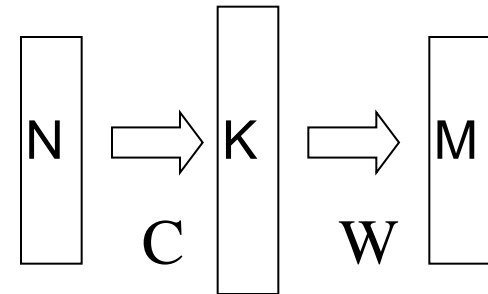
Rmk: the parameter σ controls the width of the gaussian

RBF networks

Computation of the output signal:

$$y_i = \sum_{k=1}^K w_{ik} g(\|X - C^k\|) - w_{i0}, \quad i = \overline{1, M}$$

$$y_i = \sum_{k=1}^K w_{ik} z_k - w_{i0}, \quad z_k = g(\|X - C^k\|)$$



Center matrix

Weight matrix

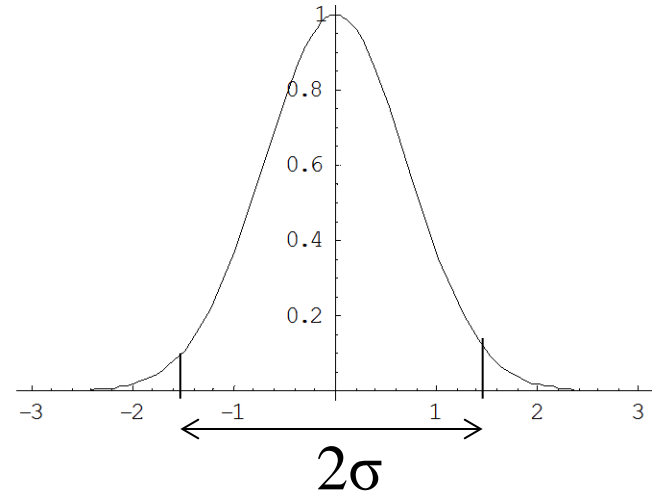
The vectors C^k can be interpreted as prototypes;

- only input vectors similar to the prototype of the hidden unit “activate” that unit
- the output of the network for a given input vector will be influenced only by the output of the hidden units having centers close enough to the input vector

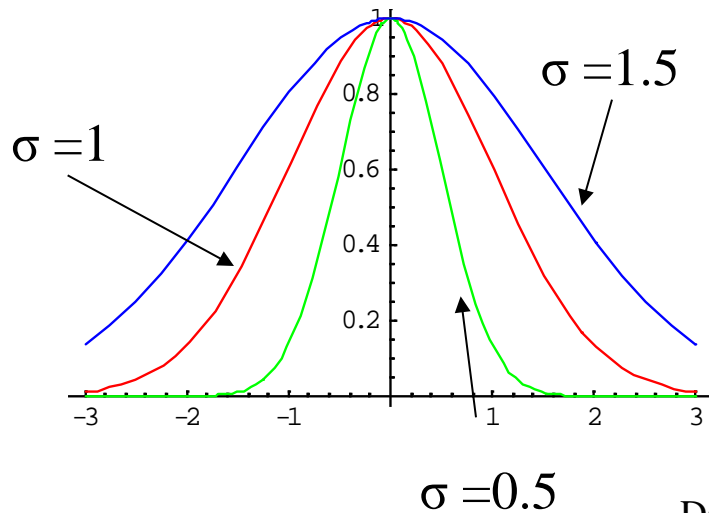
RBF networks

Each hidden unit is “sensitive” to a region in the input space corresponding to a neighborhood of its center. This region is called **receptive field**

The size of the receptive field depends on the parameter σ



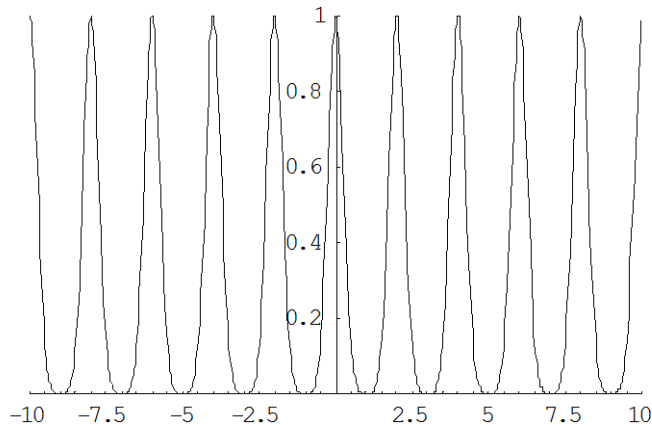
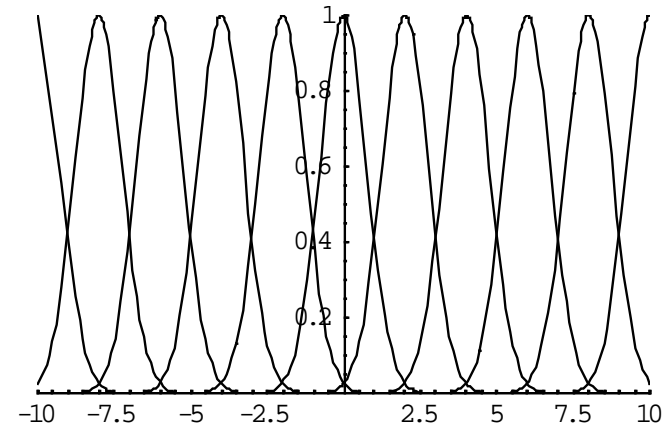
$$g(u) = \exp\left(-\frac{u^2}{2\sigma^2}\right)$$



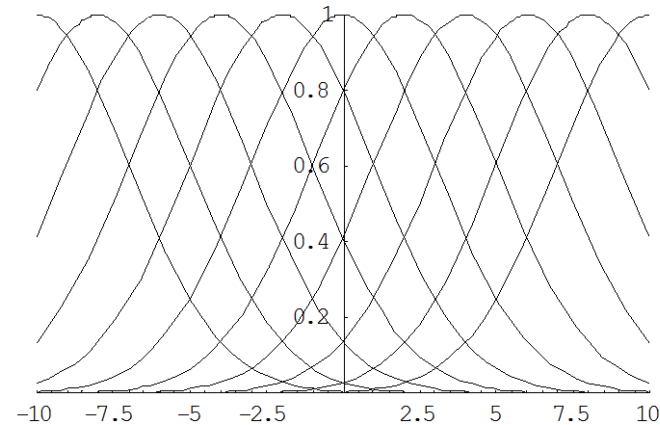
RBF networks

- The receptive fields of all hidden units covers the input space
- A good covering of the input space is essential for the approximation power of the network
- Too small or too large values of the width of the radial basis function lead to inappropriate covering of the input space

appropriate covering



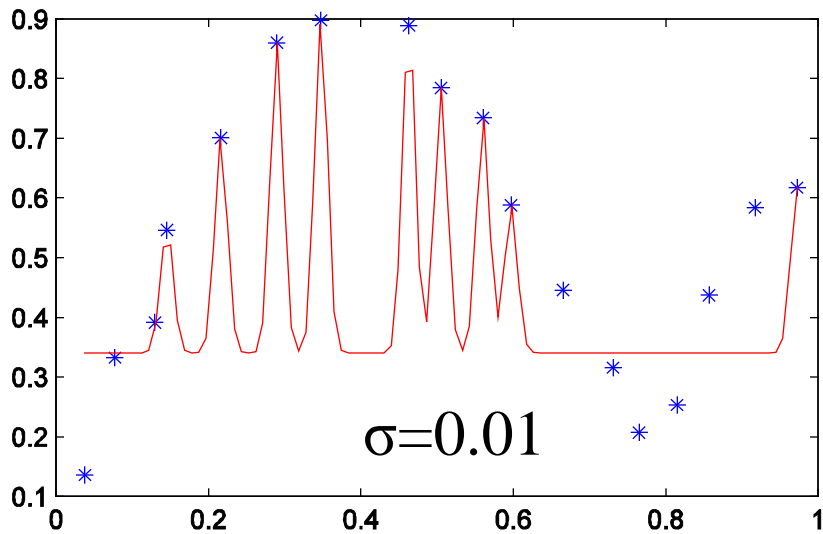
undercovering



overcovering

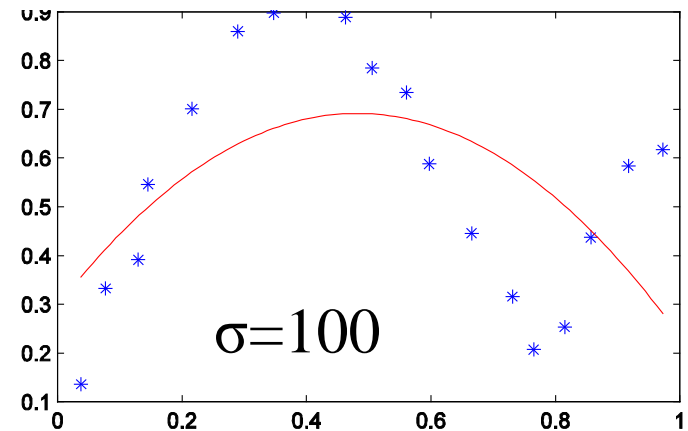
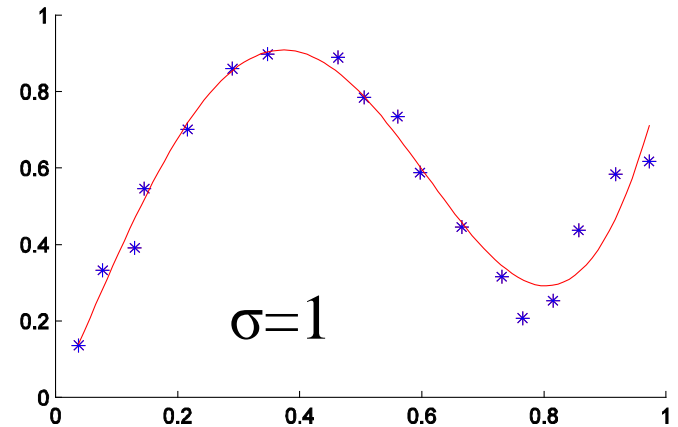
RBF networks

- The receptive fields of all hidden units covers the input space
- A good covering of the input space is essential for the approximation power of the network
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undercovering

appropriate covering



overcovering

RBF networks

RBF networks are universal approximators:

a network with N inputs and M outputs can approximate any function defined on \mathbb{R}^N , taking values in \mathbb{R}^M , as long as there are enough hidden units

The theoretical foundations of RBF networks are:

- Theory of approximation
- Theory of regularization

RBF networks

Adaptive parameters:

- Centers (prototypes) corresponding to hidden units
- Receptive field widths (parameters of the radial symmetry activation functions)
- Weights associated to connections between the hidden and output layers

Learning variants:

- **Simultaneous learning of all parameters** (similar to BackPropagation)
 - **Rmk:** same drawbacks as multilayer perceptron's BackPropagation
- **Separate learning of parameters:** centers, widths, weights

RBF networks

Separate learning :

Training set: $\{(x^1, d^1), \dots, (x^L, d^L)\}$

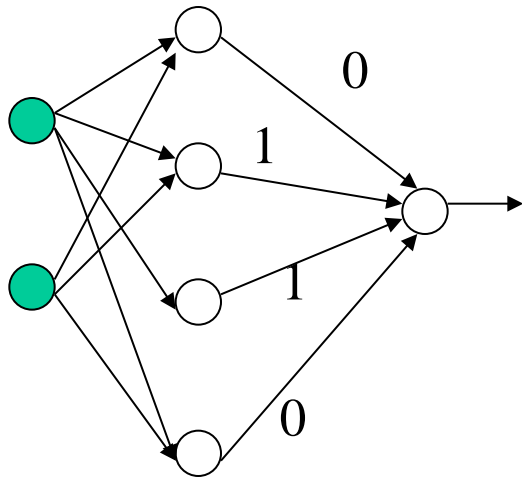
1. Estimating of the centers: simplest variant

- $K=L$ (nr of centers = nr of examples),
 - $C^k=x^k$ (this corresponds to the case of exact interpolation: see the example for XOR)

RBF networks

Example (particular case) : RBF network to represent XOR

- 2 input units
- 4 hidden units
- 1 output unit



Centers:

Hidden unit 1: (0,0)

Hidden unit 2: (1,0)

Hidden unit 3: (0,1)

Hidden unit 4: (1,1)

Weights:

w1: 0

w2: 1

w3: 1

w4: 0

Activation function:

$g(u)=1$ if $u=0$

$g(u)=0$ if $u \neq 0$

This approach cannot be applied for general approximation problems

RBF networks

Separate learning :

Training set: $\{(x^1, d^1), \dots, (x^L, d^L)\}$

1. Estimating of the centers

- $K < L$: the centers are established
 - by random selection from the training set
 - simple but not very effective
 - by systematic selection from the training set (Orthogonal Least Squares)
 - by using a clustering method

RBF networks

Orthogonal Least Squares:

- Incremental selection of centers such that the error on the training set is minimized
- The new center is chosen such that it is orthogonal on the space generated by the previously chosen centers (this process is based on the Gram-Schmidt orthogonalization method)
- This approach is related with regularization theory and ridge regression

RBF networks

Clustering:

- Identify K groups in the input data $\{X^1, \dots, X^L\}$ such that data in a group are sufficiently similar and data in different groups are sufficiently dissimilar
- Each group has a representative (e.g. the mean of data in the group) which can be considered the center
- The algorithms for estimating the representatives of data belong to the class of partitional clustering methods
- Classical algorithm: K-means

RBF networks

Incremental variant:

- Start with a small number of centers, randomly initialized
- Scan the set of input data:
 - If there is a center close enough to the data then this center is slightly adjusted in order to become even closer to the data
 - if the data is dissimilar enough with respect to all centers then a new center is added (the new center will be initialized with the data vector)

RBF networks

Incremental variant:

$$K := K_0$$

$$C_i^k := \text{rand}(\text{min}, \text{max}), i = 1..N; k = 1..K$$

$$t := 0$$

REPEAT

FOR $l := 1, L$ DO

find $k^* \in \{1, \dots, K\}$ such that $d(X^l, C^{k^*}) \leq d(X^l, C^k)$

IF $d(X^l, C^{k^*}) < \delta$ THEN $C^{k^*} := C^{k^*} + \eta \cdot (X^l - C^{k^*})$

ELSE $K := K + 1; C^K := X^l$

$$t := t + 1$$

$$\eta := \eta_0 t^{-\alpha}$$

UNTIL $t > t_{\max}$ OR $\eta < \varepsilon$

δ is a dissimilarity threshold

α controls the decrease of the learning rates

RBF networks

2. Estimating the receptive fields widths.

Heuristic rules:

$$\sigma = \frac{d_{\max}}{\sqrt{2K}}, \quad d_{\max} = \text{maximal distance between centers}$$

$$\sigma_k = \gamma d(C^k, C^j), \quad C^j = \text{the closest center to } C^k, \gamma \in [0.5, 1]$$

$$\sigma_k = \frac{1}{m} \sum_{j=1}^m d(C^k, C^j), \quad C^1, \dots, C^m : \text{the closest } m \text{ centers to } C^k$$

$$\sigma_k = \frac{1}{q_k} \sum_{j=1}^{q_k} d(C^k, X^j), \quad X^1, \dots, X^{q_k} : \text{input vectors represented by unit } k$$

RBF networks

3. Estimating the weights of connections between hidden and output layers:

- This is equivalent with the problem of training one layer linear network
- Variants:
 - Apply linear algebra tools (pseudo-inverse computation)
 - Apply Widrow-Hoff learning (training based on the gradient method applied to one layer neural networks)

Initialization:

$w_{ij}(0) := \text{rand}(-1, 1)$ (the weights are randomly initialized in $[-1, 1]$),
 $k := 0$ (iteration counter)

Iterative process

REPEAT

FOR $l := 1, L$ DO

 Compute $y_i(l)$ and $\text{delta}_i(l) = d_i(l) - y_i(l)$, $i = 1, M$

 Adjust the weights: $w_{ij} := w_{ij} + \text{eta} * \text{delta}_i(l) * x_j(l)$

 Compute the $\text{SSE}(W)$ for the new values of the weights

$k := k + 1$

UNTIL $\text{SSE}(W) < E^*$ OR $k > k_{\text{max}}$

(E^* = approximation error, k_{max} = maximal number of iterations)

RBF vs. BP networks

RBF networks:

- 1 hidden layer
- Distance based aggregation function for the hidden units
- Activation functions with radial symmetry for hidden units
- Linear output units
- Separate training of adaptive parameters
- Similar with local approximation approaches

BP networks:

- many hidden layers
- Weighted sum as aggregation function for the hidden units
- Sigmoidal activation functions for hidden neurons
- Linear/nonlinear output units
- Simultaneous training of adaptive parameters
- Similar with global approximation approaches