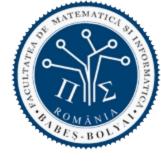


### BABEŞ-BOLYAI UNIVERSITY Faculty of Computer Science and Mathematics



# ARTIFICIAL INTELLIGENCE

#### **Intelligent systems**

Machine learning
Support Vector Machines
K-means

### **Topics**

#### A. Short introduction in Artificial Intelligence (AI)

#### A. Solving search problems

- A. Definition of search problems
- B. Search strategies
  - A. Uninformed search strategies
  - B. Informed search strategies
  - c. Local search strategies (Hill Climbing, Simulated Annealing, Tabu Search, Evolutionary algorithms, PSO, ACO)
  - D. Adversarial search strategies

#### c. Intelligent systems

- A. Rule-based systems in certain environments
- B. Rule-based systems in uncertain environments (Bayes, Fuzzy)

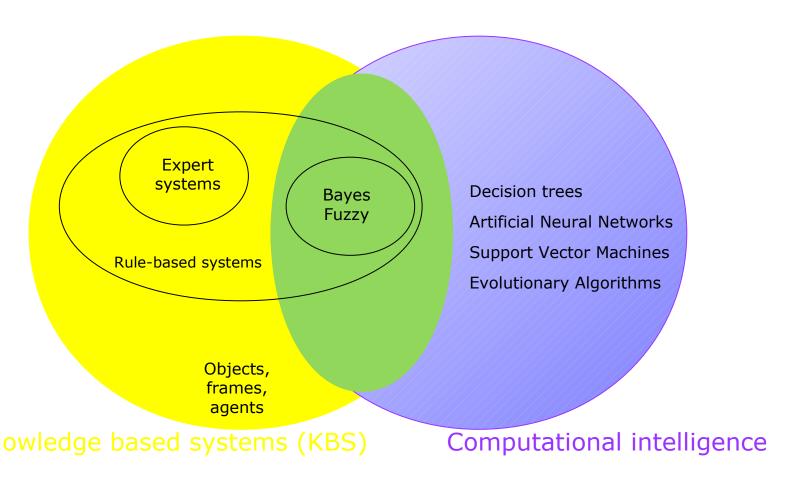
#### c. Learning systems

- A. Decision Trees
- **B.** Artificial Neural Networks
- c. Evolutionary algorithms
- D. Support Vector Machines
- **E.** K-means
- D. Hybrid systems

### Useful information

- Chapter 15 of C. Groşan, A. Abraham, Intelligent Systems: A Modern Approach, Springer, 2011
- Chapter 9 of T. M. Mitchell, Machine Learning, McGraw-Hill Science, 1997
- Documents from svm folder

### Intelligent systems



### Intelligent systems – Machine Learning

#### Typology

#### Experience criteria:

- Supervised learning
- Unsupervised learning
- Active learning
- Reinforcement learning

#### Algorithm criteria

- Decision trees
- Artificial Neural Networks
- Evolutionary Algorithms
- Support Vector Machines
- Hidden Markov Models
- K-means

- Support Vector Machines (SVMs)
  - Definition
  - Solved problems
  - Advantages
  - Difficulties
  - Tools

#### Definition

- Developed by Vapnik in 1970
- Popularised after 1992
- Linear classifiers that identify the hyper-plane that separates the positive and negative classes
- Have a theoretical foundation
- Work very well for large data (text mining, image analysis)

#### Remember

- Supervised learning problem a data set:
  - (x<sup>d</sup>, t<sup>d</sup>), with:
  - $X^d \in \mathbb{R}^m \rightarrow X^d = (X^d_1, X^d_2, \dots, X^d_m)$
  - $t^d \in \mathbb{R}$  →  $t^d \in \{1, -1\}, 1$  → positive class, -1 → negative class
  - where d = 1,2,...,n,n+1,n+2,...,N
- First n data (x<sup>d</sup> and t<sup>d</sup> are known) are used as training data
- Last N-n data (x<sup>d</sup> is known, t<sup>d</sup> is unknown) are used as testing data

Inteligentă artificială - sisteme inteligente (SVM, k-means)

#### Definition

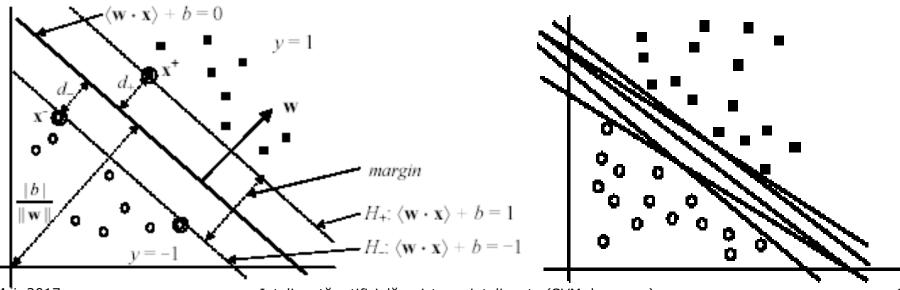
■ SVM finds a linear function  $f(\mathbf{x}) = \langle \mathbf{w} \cdot \mathbf{x} \rangle + b$ ,  $(\mathbf{w} - \text{weight vector})$  such as

$$y_i = \begin{cases} 1 & if \langle \mathbf{w} \cdot \mathbf{x}_i \rangle + b \ge 0 \\ -1 & if \langle \mathbf{w} \cdot \mathbf{x}_i \rangle + b < 0 \end{cases}$$

■  $\langle \mathbf{w} \cdot \mathbf{x} \rangle + b = 0$  → decision hyper-plane that separates the two classes

#### Definition

- There are more hyper-planes
  - Which is the best hyper-plane?
- SVM searches the hyper-plane with the largest margin (that minimises the generalisation error)
  - □ SMO (Sequential minimal optimization) algorithm



#### Solved problems

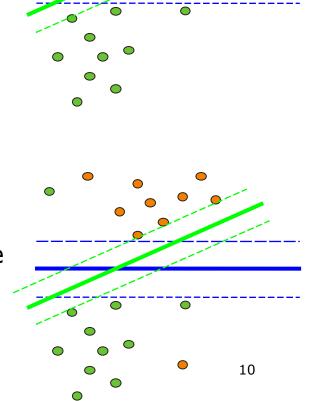
- Classification problems → more cases (based on the data type):
  - Linear separable



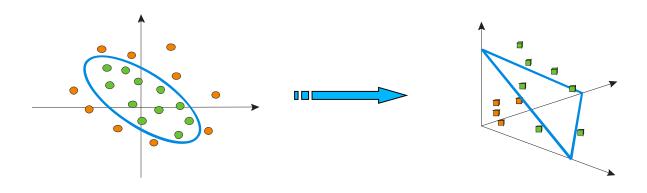
Error = 0



- Constrains are relaxed → some error are allowed
- C penalisation coefficient



- □ Solved problems → classification problems → data cases:
  - Non-linear separable
    - Input space is transformed (mapped) into a space of more dimensions (feature space) by using kernel function – in this new space the data becomes linear separable
    - In SVMs the kernel function computes the distance among 2 points
      - → kernel ~ similarity function

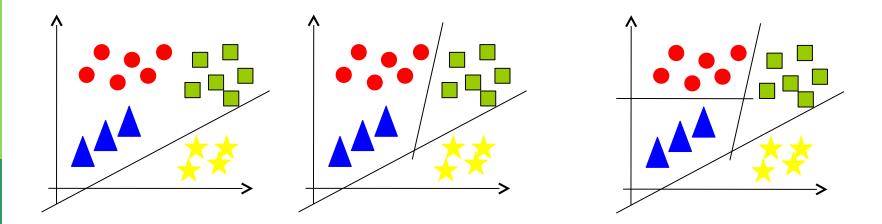


- □ Solved problems → classification problems → data cases:
  - Non-linear separable → possible kernels
    - Classic kernels
      - Polynomial kernel:  $K(\mathbf{x}^{d1}, \mathbf{x}^{d2}) = (\mathbf{x}^{d1}, \mathbf{x}^{d2} + 1)^p$
      - RBF kernel:  $K(\mathbf{x}^{d1}, \mathbf{x}^{d2}) = exp(-||\mathbf{x}^{d1} \mathbf{x}^{d2}||^2/2\sigma^2)$
    - Multiple Kernels
      - Linear :  $K(\mathbf{x}^{d1}, \mathbf{x}^{d2}) = \sum w_i K_i (\mathbf{x}^{d1}, \mathbf{x}^{d2})$
      - Non-linear
        - Without coefficients:  $K(\mathbf{x}^{d1}, \mathbf{x}^{d2}) = K_1(\mathbf{x}^{d1}, \mathbf{x}^{d2}) + K_2(\mathbf{x}^{d1}, \mathbf{x}^{d2}) * \exp(K_3(\mathbf{x}^{d1}, \mathbf{x}^{d2}))$
        - With coefficients:  $K(\mathbf{x}^{d1}, \mathbf{x}^{d2}) = K_1(\mathbf{x}^{d1}, \mathbf{x}^{d2}) + c_1 * K_2 * (\mathbf{x}^{d1}, \mathbf{x}^{d2}) exp(c_2 + K_3(\mathbf{x}^{d1}, \mathbf{x}^{d2}))$
    - Kernels for strings
    - Kernels for images
    - Kernels for graphs

### SVM setting

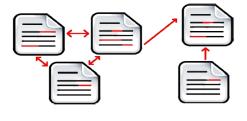
- SVM's parameters
  - Penalisation coefficient C
    - C- small → slowly convergence
    - C large → fast convergence
  - Kernel parameters (what kernel and what parameters)
    - If m (# of attributes) is larger than n (# of data)
      - SVM by a linear kernel (SVM without kernel)  $\rightarrow$   $K(\mathbf{x}^{d1}, \mathbf{x}^{d2}) = \mathbf{x}^{d1} \cdot \mathbf{x}^{d2}$
    - If m (# of attributes) is large and n (# of data) is medium
      - SVM with Gaussian kernel  $K(\mathbf{x}^{d1}, \mathbf{x}^{d2}) = \exp(-||\mathbf{x}^{d1} \mathbf{x}^{d2}||^2/2\sigma^2)$
      - σ dispersion of training data
      - Attributes must be normalised (scalled to (0,1))
    - If m (# of attributes) is small and n (# of data) is large
      - Ad new attributes and than SVM with linear kernel

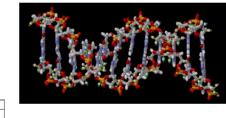
- SVM for multi-class classification problems (more than 2 classes)
  - one *vs.* all



- Structured SVMs
  - Machine Learning
    - □ Simple SVM  $f: X \rightarrow R$ 
      - Any type of inputs
      - Numerical outputs (natural numbers, integers, real numbers)
    - □ Structured SVM: X → Y
      - Any type of inputs
      - Any type of outputs (numerical or structured outputs)

- Structured information
  - Texts and hyper-texts
  - Molecules and molecular structures
  - Images





- Structured SVMs
  - Applications
    - Natural Language Processing
      - Automatic translation (outputs → sentences)
      - Syntactic and/or morphologic analysis of sentences (outputs
         → syntactic and/or morphologic tree)
    - Bioinformatic
      - Prediction of secondary structures (outputs → bi=partite graphs)
      - Prediction of enzyme function (outputs → paths in trees)
    - Speech processing
      - Automatic transcriptions (outputs → sentences)
      - Transformation of texts in voice (outputs → audio signal)
    - Robotics
      - Planning (outputs → sequences of actions)

#### Advantages

- Can work with any type of data (linear or non-linear separable, uniform distributed or not, with known or unknown distribution)
  - □ Kernel function that creates new attributes (features) → hidden layers of an ANN
- If the problem is convex SVM finds a unique solution → global optima
  - ANNs can associates more solutions → local optima
- Automatic selection of the learnt model (by support vectors)
  - In ANNs hidden layers have to be configured a priori
- Avoid over fitting
  - ANNs have over fitting problems even the cross-validation is involved

#### Difficulties

- Real attributes only
- Binary classification problems only
- Difficult mathematical background

#### Tools

- LibSVM → http://www.csie.ntu.edu.tw/~cjlin/libsvm/
- Weka → SMO
- SVMLight → http://svmlight.joachims.org/
- SVMTorch → http://www.torch.ch/
- http://www.support-vector-machines.org/

### Intelligent systems – Machine Learning

#### Typology

#### Experience criteria:

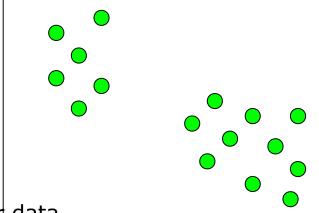
- Supervised learning
- Unsupervised learning
- Active learning
- Reinforcement learning

#### Algorithm criteria

- Decision trees
- Artificial Neural Networks
- Evolutionary Algorithms
- Support Vector Machines
- Hidden Markov Models
- K-means

### Unsupervised learning

- Aim
  - Finds a model or a structure of data
- Solved problems
  - Identification of groups (clusters)
    - Analysis of genes
    - Image processing
    - Analysis of social networks
    - Market segmentation
    - Analysis of astronomic data
    - Clusters of computers
  - Dimension reduction
  - Identification of causes (explanations) for data
  - Modelling the data densities
- Specific
  - Data are not annotated (labelled)



Separates the un-labeled examples in disjoint sub-sets (clusters) such as:

- Examples of the same cluster are similar
- Examples of different clusters are different

#### Definition

- Given
  - A set of data (examples, instances, cases)
    - Training data
      - As atribute\_data<sub>i</sub>, where
        - i = 1, N (N = # of training data)
        - atribute\_data<sub>i</sub>= (atr<sub>i1</sub>, atr<sub>i2</sub>, ..., atr<sub>im</sub>), m # of attributes (characteristics, properties) of data
    - Testing data
      - As (atribute\_data<sub>i</sub>), i =1,n (n = # of testing data)

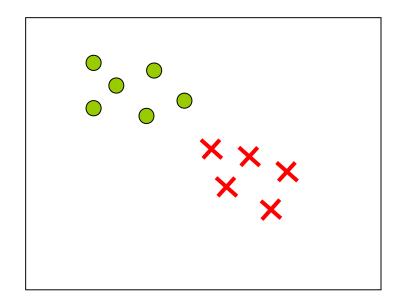
#### Determine

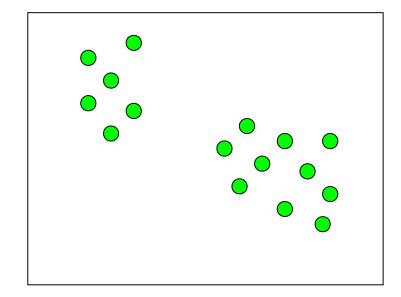
- An unknown function that groups the training data in more classes
  - # of classes can be pre-defined (k) or un-known
  - Data of the same class are similar
- The class of a new testing data by using the learnt grouping (on training data)

#### Other names

Clustering

### Supervised vs. un-supervised





- Distance between 2 elements  $\boldsymbol{p}$  and  $\boldsymbol{q} \in R^m$ 
  - Euclid distance

Manhattan distance

$$d(p,q) = \sum_{j=1,2,...,m} |p_j - q_j|$$

- Mahalanobis distance
  - $d(\mathbf{p},\mathbf{q}) = sqrt(\mathbf{p}-\mathbf{q})S^{-1}(\mathbf{p}-\mathbf{q}),$   $\text{Where S is the covariance matrix } (S = E[(\mathbf{p}-E[\mathbf{p}])(\mathbf{q}-E[\mathbf{q}])])$
- Internal product

Cosine distance

$$d(\mathbf{p},\mathbf{q}) = \sum_{j=1,2,...,m} p_j q_j / (sqrt(\sum_{j=1,2,...,m} p_j^2) * sqrt(\sum_{j=1,2,...,m} q_j^2))$$

- Hamming distance
  - # of differences between p and q
- Levenshtein distance
  - Minimal # of operations required for transforming p in q
- Distance vs. similarity
  - Distance → minimisation
  - Similarity → maximisation

**Application** 

Gene clustering

Market segmentation (for client clustering)

news.google.com

### Unsupervised learning – process

#### **Process**

- 2 steps:
  - Learning
    - Determine (learn), by using an algorithm, the existing clusters
  - Testing
    - Include a new data in one of the identified (during training) clusters

#### Learning quality (clustering validation)

- Internal criteria
  - Large similarity inside the cluster and reduce similarity between clusters
- External criteria
  - Using benchmarks composed of apriori grouped data

#### Performance measures

- Internal criteria
  - Distance inside the cluster
  - Distance between clusters
  - Davies-Bouldin index
  - Dunn index
- External criteria
  - Comparison with known data impossible in real-world applications
  - Precision
  - Recall
  - F-measure

#### Performance measures

- Internal criteria
  - Distance inside cluster  $c_j$  that contains  $n_j$  instances
    - Average distance (among instances)

$$D_a(c_j) = \sum_{x_{i1}, x_{i2} \in c_j} ||x_{i1} - x_{i2}|| / (n_j(n_j-1))$$

Nearest neighbour distance

$$D_{nn}(c_j) = \sum_{xilecj} \min_{xi2ecj} ||x_{il} - x_{i2}|| / n_j$$

Distance between centroids

• 
$$D_c(c_j) = \sum_{x_i, ec_j} ||x_i - \mu_j|| / n_j$$
, unde  $\mu_j = 1 / n_j \sum_{x_i ec_j} x_i$ 

#### Performance measure

- Internal criteria
  - Distance between 2 clusters  $c_{j1}$  and  $c_{j2}$ 
    - Simple link

• 
$$d_s(c_{j1}, c_{j2}) = min_{xilecjl, xi2ecj2} \{||x_{il} - x_{i2}||\}$$

Complete link

• 
$$d_{co}(c_{i1}, c_{i2}) = \max_{xilecil, xi2eci2} \{||x_{i1} - x_{i2}||\}$$

Average link

• 
$$d_a(c_{j1}, c_{j2}) = \sum_{xilecjl, xi2ecj2} \{||x_{il} - x_{i2}||\} / (n_{jl} * n_{j2})$$

Link between centroids

• 
$$d_{ce}(c_{j1}, c_{j2}) = ||\mu_{j1} - \mu_{j2}||$$

#### Performance measures

- Internal criteria
  - Davies-Bouldin index → min → compact clusters

$$DB = 1/nc*\sum_{i=1,2,...,nc} max_{j=1,2,...,nc,j\neq i} ((\sigma_i + \sigma_j)/d(\mu_i, \mu_j))$$

- where:
  - nc # of clusters
  - μ<sub>i</sub> centroid of cluster i
  - $\sigma_i$  average of distances between elements form cluster i and the centroid  $\mu_i$
  - $d(\mu_i, \mu_i)$  distance between centroid  $\mu_i$  and centroid  $\mu_i$

#### Dunn index

- Identifies the dense clusters and well separated
- $\square$   $D=d_{min}/d_{max}$
- where:
  - $d_{min}$  minimal distance between 2 elements from different clusters intracluster distance
  - $d_{max}$  maximal distance between 2 elements from the same cluster intercluster distance

- How the clusters are forming
  - Hierarchic clustering
  - Non-hierarchic (partitioned) clustering
  - Clustering based on data density
  - Clustering based on a grid

- How the clusters are forming
  - Hierarchic clustering
    - Creates a dendogram (taxonomic tree)
      - Creates the clusters (recursively)
      - k (# of clusters) is un-known
    - □ Aglomerativ clustering (bottom-up) → small clusters to large clusters
    - □ Divisiv clustering (top-down) → large clusters to small clusters
    - □ Eg.
      - Clustering ierarhic aglomerativ

- How the clusters are formed
  - Non-hierarchic
    - □ Partitional → determine a data separation → all the clusters in the same time
    - Optimises an objective function defined
      - Locally by using some features only
      - Globally by using all attributes

#### that can be:

- squared error sum of squared distances between data and the cluster's centroid → min
  - Ex. K-means
- Graph-based
  - Ex. Clustering based in minimum spanning tree
- Based on probabilistic models
  - Ex. Identify the data distribution → expectation maximisation
- Based on the nearest neighbour
- Required to fix k apriori → fix the initial clusters
  - Algorithm is run more times with different parameters and the most efficient version is selected
- Ex. K-means, ACO

- How the clusters are forming
  - Based on data densities
    - Data density and data connectivity
      - Cluster formation is based on data density from a given region
      - Cluster formation is based on data connectivity from a given region
    - Function of data density
      - Tries to model the data distribution
    - Advantage:
      - Modeling of clusters of any shape

- How the cluster are forming
  - Based on a grid
    - Is not a distinct approach
      - Can be hierarchic, partitional or density-based
    - Involves data space segmentation in regular areas
    - Objects are placed on a multi-dimensional grid
    - □ Eg. ACO

- How the algorithms work
  - Agglomerativ clustering
    - 1. Initially, each instance form a cluster
    - 2. Compute the distance between any 2 clusters
    - 3. Reunion the closest 2 clusters
    - Repeat steps 3 and 4 until a single cluster is obtained or other stop criterion is satisfied
  - Divisive clustering
    - Establish the number of clusters (k)
    - 2. Initialise the centre of each cluster
    - 3. Determine a data separation
    - 4. Re-compute the centre of each cluster
    - Repeat steps 3 and 4 until the partition is unchanged (algorithm converges)
- How the algorithm takes into account the attributes (features)
  - Monotonic attributes are taken into account one-by-one
  - Polytonic attributes are simultaneous taken into

### How the data belong to clusters

- Exact clustering (hard clustering)
  - $\square$  Each instance  $x_i$  has associated a label (class)  $c_j$
- Fuzzy clustering
  - □ Each instance has associated a membership degree (probability)  $f_{ij}$  to a given class  $c_j$  → an instance xi can belongs to more clusters
  - □ Asociază fiecarei intrări  $\mathbf{x}_i$  un grad (probabilitate) de apartenență  $f_{ij}$  la o anumită clasă  $c_j \rightarrow$  o instanță  $\mathbf{x}_i$  poate aparţine mai multor clusteri

### Unsupervised learning – algorithms

- Agglomerative hierarchical clustering
- K-means
- AMA
- Probabilistic models
- Nearest neighbour
- Fuzzy
- Artificial Neural Network
- Evolutionary algorithms
- ACO

- Agglomerative hierarchical clustering
- □ Consider a distance between 2 instances  $d(x_{i1}, x_{i2})$
- Form N clusters, each of them containing an instance
- Repeat
  - Determine the closest 2 clusters
  - Reunion the 2 clusters → a cluster
- Until a single cluster is obtained (that contains all the instances)

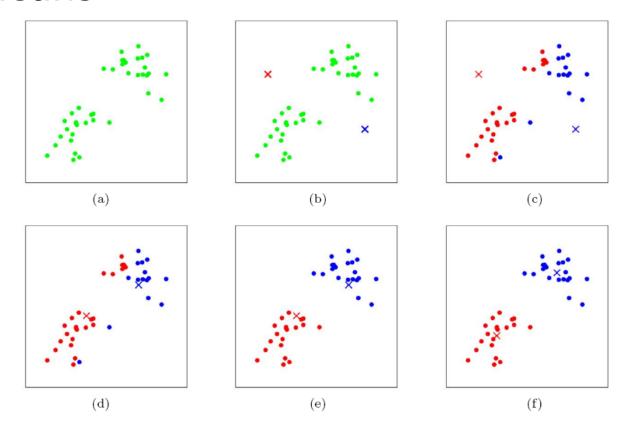
### Agglomerative hierarchical clustering

- $\square$  Distance between 2 clusters  $c_i$  and  $c_j$ :
  - Simple link → minimal distance between the objects of 2 clusters
  - Complete link → maximal distance between the objects of 2 clusters
  - Average link → mean of distances between the objects of 2 clusters
  - Average link over group → distance between the means (centroids) of 2 clusters

$$\square d(c_i, c_j) = \rho(\boldsymbol{\mu}_i, \boldsymbol{\mu}_j), \rho - distance, \boldsymbol{\mu}_j = 1/n_j \sum_{x_i \in c_j} \boldsymbol{x}_i$$

- K-means (Lloyd algorithm / Voronoi iteration)
- Suppose that k clusters will form
- Initialise k centroids  $\mu_1, \mu_2, ..., \mu_k$ 
  - A centroid  $\mu_j$  (i=1,2,...,k) is a vector of m values (m # of features)
- Repeat until convergence
  - Associated to each instance the nearest centroid  $\rightarrow$  for each instance  $\mathbf{x}_i$ , i = 1, 2, ..., N
    - $c_i = arg min_{j=1, 2, ..., k} || \mathbf{x}_i \mathbf{\mu}_j ||^2$
  - Re-compute the centroids by moving them in the mean of instances associated to it  $\rightarrow$  for each cluster  $c_j$ , j=1,2,...,k

### K-means



#### K-means

- Initialisation of k centroids  $\mu_1, \mu_2, ..., \mu_k$ 
  - With random values (in the definition domain of the problem)
  - With k instances of N (randomly selected)
- Does algorithm converge always?
  - Yes, because of distortion function J

$$\Box J(c, \mu) = \sum_{i=1,2,...,N} || \mathbf{x}_i - \boldsymbol{\mu}_{cj}||^2$$

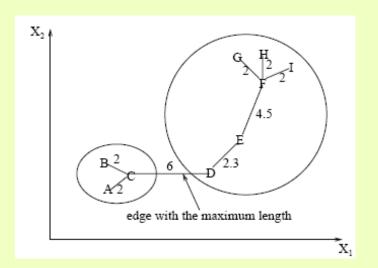
which is decreasing

- Converges in a local optima
- Finding the global optima → NP-difficult problem

### Unsupervised learning – algorithms Clusterizare bazată pe arborele minim de acoperire (AMA)

## Clustering based on minimum spanning tree

- Construct the minimum spanning tree of data
- Eliminate from the tree the longest edges and form clusters



### Învățare ne-supervizată – algoritmi Modele probabilistice

- http://www.gatsby.ucl.ac.uk/~zoubin/course0
- http://learning.eng.cam.ac.uk/zoubin/nipstut.

# Învățare ne-supervizată – algoritmi Cel mai apropiat vecin

- Se etichetează câteva dintre instanțe
- Se repetă până la etichetarea tuturor instanţelor
  - O instanţă ne-etichetată va fi inclusă în clusterul instanţei cele mai apropiate
    - dacă distanţa între instanţa neetichetată şi cea etichetată este mai mică decât un prag

## Învățare ne-supervizată – algoritmi Clusterizare fuzzy

- Se stabileşte o partiţionare fuzzy iniţială
  - Se construiește matricea gradelor de apartenență U, unde  $u_{ij}$  gradul de apartenență al instanței  $\boldsymbol{x_i}$  (i=1,2,...,N) la clusterul  $c_i$  (j=1,2,...,k)  $(u_{ij} \in [0,1])$ 
    - $\Box$  Cu cât  $u_{ij}$  e mai mare, cu atât e mai mare încrederea că instanța  $x_i$  face parte din clusterul  $c_i$
- Se stabileşte o funcţie obiectiv
  - $\square E^{2}(U) = \sum_{i=1,2,...,N} \sum_{j=1,2,...,k} u_{ij} || \mathbf{x}_{i} \boldsymbol{\mu}_{j} ||^{2},$ 
    - unde  $\mu_j = \sum_{i=1,2,...,N} u_{ij} \mathbf{x}_i$  centrul celui de-al j-lea fuzzy cluster
  - care se optimizează (min) prin re-atribuirea instanţelor (în clusteri noi)
- □ Clusering fuzzy → clusterizare hard (fixă)
  - impunerea unui prag funcţiei de apartenenţă u<sub>ij</sub>

### Învățare ne-supervizată – algoritmi Algoritmi evolutivi

- Algoritmi
  - Inspiraţi din natură (biologie)
  - Iterativi
  - Bazaţi pe
    - populaţii de potenţiale soluţii
    - căutare aleatoare ghidată de
      - Operaţii de selecţie naturală
      - Operaţii de încrucişare şi mutaţie
  - Care procesează în paralel mai multe soluţii
- Metafora evolutivă

Evoluţie naturală	Rezolvarea problemelor
Individ	Soluţie potenţială (candidat)
Populație	Mulţime de soluţii
Cromozom	Codarea (reprezentarea) unei soluţii
Genă	Parte a reprezentării
Fitness (măsură de adaptare)	Calitate
Încruवृद्धिक्षुं समृत्रं विद्यान	_
Mediu (SVM, K	-means) Spaţiul de căutare al problemei

## Învățare ne-supervizată – algoritmi Algoritmi evolutivi

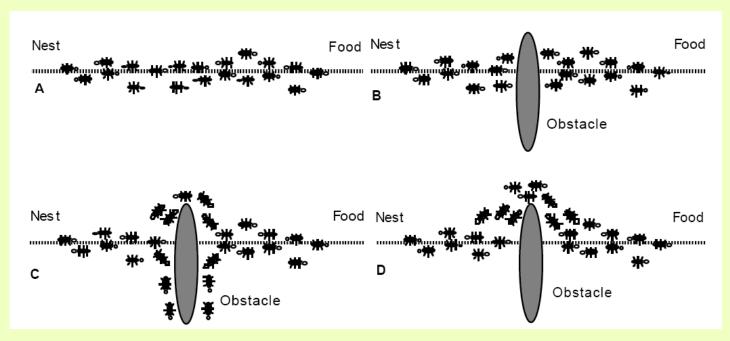
```
Initializare populație P(0)
Evaluare P(0)
g := 0; //generaţia
CâtTimp (not condiţie_stop) execută
   Repetă
                                                          -Încrucişare
        Selectează 2 părinți p1 și p2 din P(g
        Incrucişare(p1,p2) => 01 şi o2
                                              pentru
                                                perturbare
        Mutație(o1) => o1*
        Mutație(o2) => o2*
                                              Selecție
        Evaluare(o1*)
        Evaluare(o2*)
        adăugare o1* și o* în P(g+1)
                                                           Selecție de
   Până când P(g+1) este completă
                                                           supravieţuire
   g := g + 1
Sf CâtTimp
                                                                     Populație
                                               Populație
                                                                     (urmaşi)
                                               (părinţi)
```

### Învățare ne-supervizată – algoritmi Algoritmi evolutivi

- Reprezentare
  - Cromozomul = o partiţionare a datelor
    - □ Ex. 2 clusteri → cromozom = vector binar
    - □ Ex. K clusteri  $\rightarrow$  cromozom = vector cu valori din  $\{1,2,...,k\}$
- Fitness
  - Calitatea partiţionării
- Iniţializare
  - Aleatoare
- Încrucişare
  - Punct de tăietură
- Mutaţie
  - Schimbarea unui element din cromozom

### Învățare ne-supervizată – algoritmi ACO

- Preferinţa pentru drumuri cu nivel ridicat de feromon
- Pe drumurile scurte feromonul se înmulţeşte
- Furnicile comunică pe baza urmelor de feromon



### Învățare ne-supervizată – algoritmi ACO

- Algoritm de clusterizare bazat pe un grid
- Obiectele se plasează aleator pe acest grid, urmând ca furnicuţele să le grupeze în funcţie de asemănarea lor
- 2 reguli pentru furnicuţe
  - Furnica "ridică" un obiect-obstacol
    - Probabilitatea de a-l ridica e cu atât mai mare cu cât obiectul este mai izolat (în apropierea lui nu se află obiecte similare)
    - $p(ridica) = (k^+/(k^++f))^2$
  - Furnica "depune" un obiect (anterior ridicat) într-o locație nouă
    - Probabilitatea de a-l depune e cu atât mai mare cu cât în vecinătatea locului de plasare se afla mai multe obiecte asemănătoare
    - $p(depune) = (f/(k^-+f))^2$
  - $k^+$ ,  $k^-$  constante
  - *f* procentul de obiecte similare cu obiectul curent din memoria furnicuţei
- Furnicuţele
  - au memorie
    - reţin obiectele din vecinătatea poziţiei curente
  - se mişcă ortogonal (N, S, E, V) pe grid pe căsuţele neocupate de alte furnici

## Recapitulare



- Sisteme care învaţă singure (SIS)
  - Maşini cu suport vectorial (MSV)
    - Modele computaționale care
      - rezolvă (în special) probleme de învăţare supervizată
      - prin identificarea celui mai bun hyper-plan de separare a datelor
  - K-means
    - Modele computaţionale care
      - rezolvă probleme de clusterizare
        - → nu se cunosc etichetele claselor
      - prin
        - minimizarea diferenţelor între elementele aceleaşi calse
        - maximizarea diferenţelor între elementele claselor diferite

## Cursul următor

- A. Scurtă introducere în Inteligența Artificială (IA)
- B. Rezolvarea problemelor prin căutare
  - Definirea problemelor de căutare
  - Strategii de căutare
    - Strategii de căutare neinformate
    - Strategii de căutare informate
    - Strategii de căutare locale (Hill Climbing, Simulated Annealing, Tabu Search, Algoritmi evolutivi, PSO, ACO)
    - Strategii de căutare adversială

### c. Sisteme inteligente

- Sisteme bazate pe reguli în medii certe
- Sisteme bazate pe reguli în medii incerte (Bayes, factori de certitudine, Fuzzy)
- Sisteme care învaţă singure
  - Arbori de decizie
  - A. Rețele neuronale artificiale
  - в. Maşini cu suport vectorial
  - в. Algoritmi evolutivi
- B. Sisteme hibride

- Informaţiile prezentate au fost colectate din diferite surse de pe internet, precum şi din cursurile de inteligenţă artificială ţinute în anii anteriori de către:
  - Conf. Dr. Mihai Oltean www.cs.ubbcluj.ro/ ~moltean
  - Lect. Dr. Crina Groşan www.cs.ubbcluj.ro/~ cgrosan
  - Prof. Dr. Horia F. Pop www.cs.ubbcluj.ro/~hfpop