

Anno Accademico 2021/2022	
COMPUTATIONAL LEARNING IN BIOMEDICINE	
Enrollment year	2021/2022
Academic year	2021/2022
Regulations	DM270
Academic discipline	ING-INF/06 (ELECTRONIC AND INFORMATION BIOENGINEERING)
Department	DEPARTMENT OF ELECTRICAL, COMPUTER AND BIOMEDICAL ENGINEERING
Course	BIOENGINEERING
Curriculum	Cellule, tessuti e dispositivi
Year of study	1°
Period	2nd semester (07/03/2022 - 17/06/2022)
ECTS	9
Lesson hours	80 lesson hours
Language	Italian
Activity type	WRITTEN TEST
Teacher	BELLAZZI RICCARDO - 6 ECTS RAMAT STEFANO - 3 ECTS
Prerequisites	Basic knowledge of statistics and probability theory. Basic knowledge of informatics and statistical software tools
Learning outcomes	The course aims to provide students with methodological skills and techniques to: * use in biomedical applications a large class of algorithms that are able to learn decision rules from data and automatically improve their performance based on experience. In the first part of the course, basic methods of machine learning will be introduced. At the end of this part, the student should be able to: * soundly apply machine learning approaches to learn decision rules from

data * use machine learning software tools and statistical packages The

course will include both lectures and practical hands-on computer

In the second part of the course a specific focus will be given to two

lessons.

widely used methodologies in the field of Artificial Intelligence: neural networks and deep learning on the one hand and genetic algorithms on the other. The former represent a computational learning tool for both static and dynamic recognition and classification tasks, the latter are an extremely versatile stochastic-based optimization method. At the end of the course, students should be able to implement the main "shallow" and "deep" network architectures for classification and approximation, as well as generational and steady state genetic algorithms in the Matlab environment.

Course contents

Part 1.

Learning decision rules - supervised learning

Introduction: Machine Learning and Data Mining in the biomedical sciences.

Areas of application of automatic methods for classification: diagnosis, prognosis, research

The basic concepts: examples, instances, attributes, and representation of decision rules

Decision Trees: learning techniques for pruning

Bayesian methods: Naive Bayes discriminant analysis

Regression models: linear model, logistic regression, neural networks, support-vector machines

Method and k-nearest distance measures

Random forests, Boosting

Techniques of feature selection. Information gain and Relief Evaluation of learning algorithms and problems of evaluation in the biomedical field

Training and Testing. Accuracy, calibration, sensitivity and specificity, precision and recall, F measure

Methods for performance evaluation. Cross Validation, Bootstrap and ROC curves.

Unsupervised learning

Association Rules

Clustering methods: K-means, K-medoids, hierarchical clustering, self-organizing maps

Evaluation of the results of the clustering methods

Applications of data mining in bio-medicine: diagnosis, prognosis, classification, functional genomics

Practical Activities

The CRISP methodology for data mining in bio-medicine.

Hands-on with computer programs: Orange, Python and Matlab for the solution of classification problems.

Part 2.

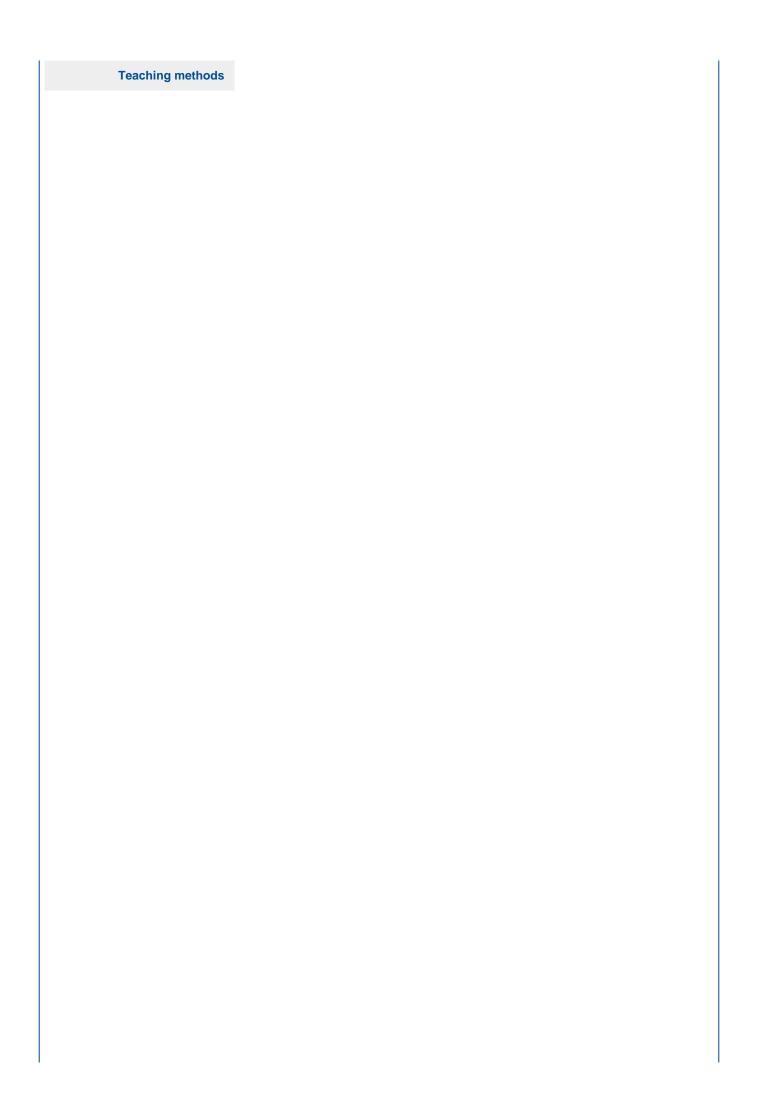
Introduction to neural networks.

The perceptron and adaline, networks based on a single neuron for classification and linear approximation.

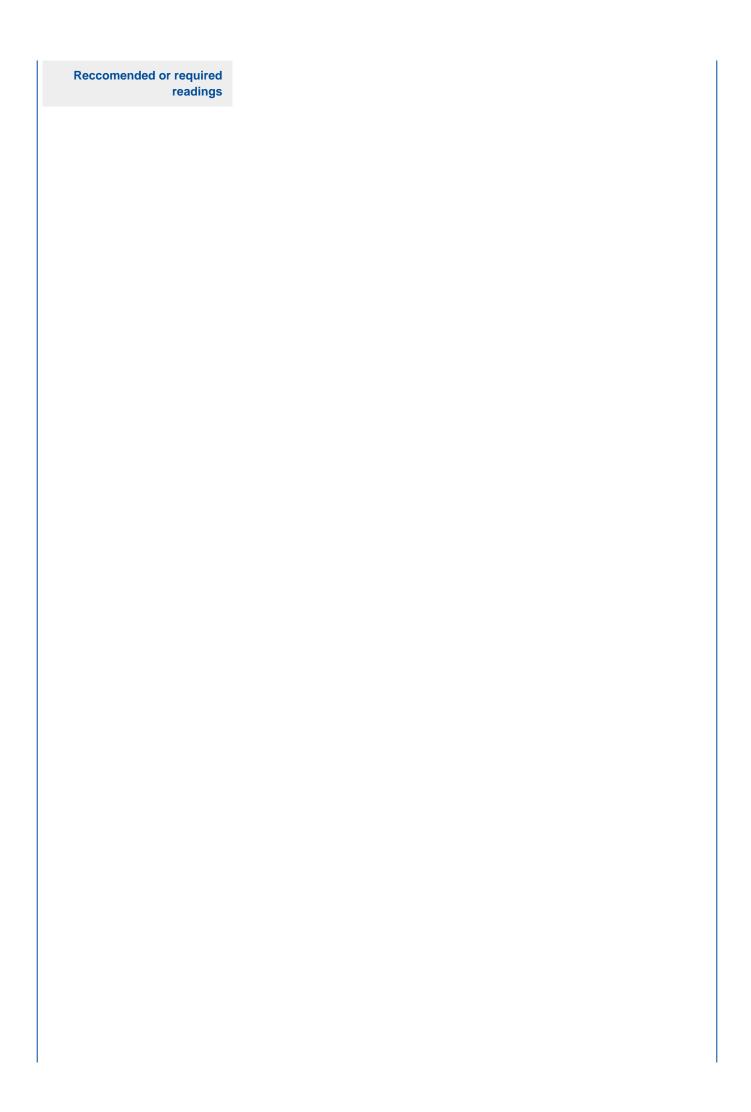
Multilayer perception and radial basis function networks.

Self organizing maps for unsupervised clustering.

Dynamic networks: the Hopfield network, the Elman network and its evolutions, the state-space model network. Recurring networks and Long Short Term Memory network. Convolutional networks, autoencoders, generative networks.



Teaching about theory, exercises and computer classes

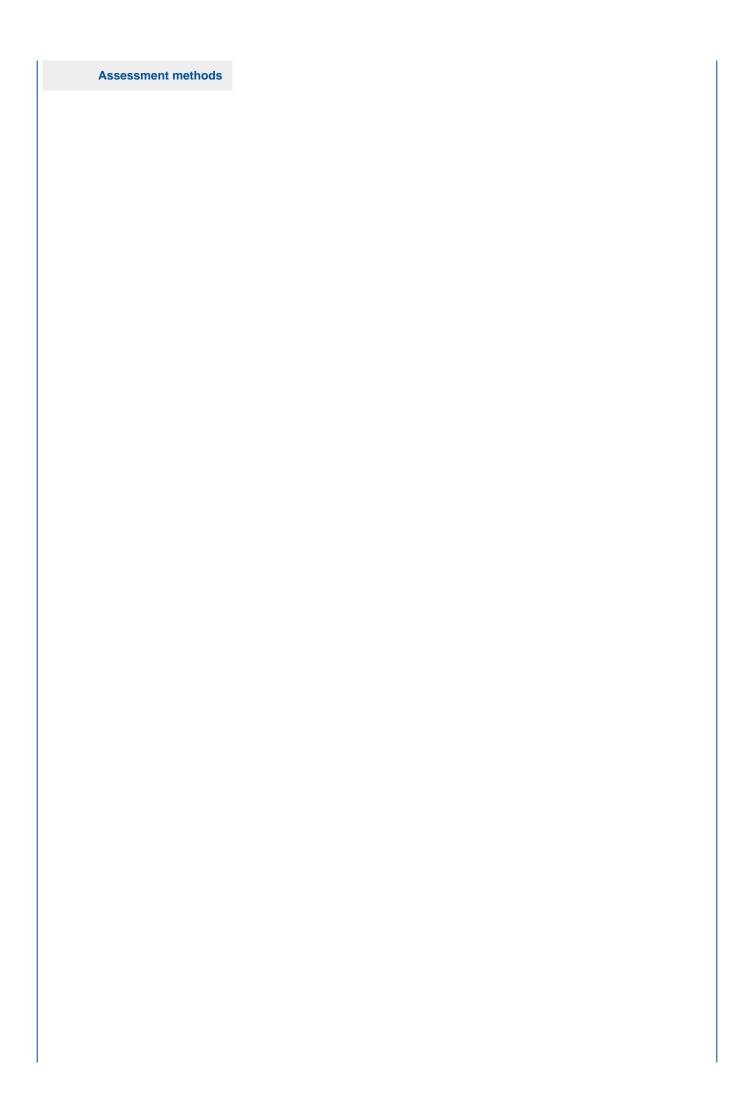


- T. Mitchell. Machine Learning. Mc Graw Hill..
- P. Tan, M. Steinbach, V. Kumar. Introduction to data mining. Addison Wesley..
- I. Witten, E. Frank. Data mining. Morgan Kaufmann.

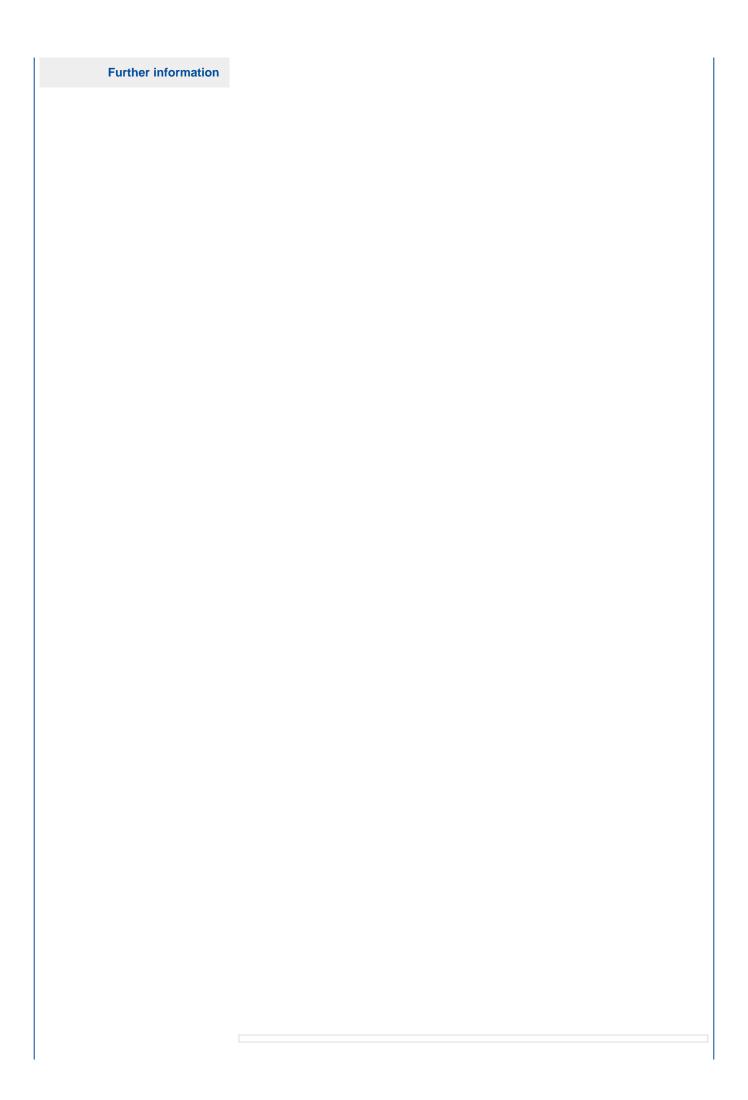
Riccardo Bellazzi. Course Slides.

- S. Haykin, Neural Networks and Learning Machines, Prentice Hall, 3rd Ed., 2009
- D.E. Goldberg, Genetic Algorithms in search, optimization and machine learning, Addison Wesley, 1989

Stefano Ramat. Course Slides.



Written test and discussion about two essays on data analysis problems, one on machine learning methods and one on neural networks and deep learning



Sustainable development goals - Agenda 2030

\$lbl legenda sviluppo sostenibile