



UNIVERSITÀ
DEGLI STUDI DI MILANO-BICOCCA

SYLLABUS DEL CORSO

Modelli Statistici II

2526-2-F8203B042-F8203B013M

Learning objectives

The course falls within the learning areas of statistical sciences, computer science, and social sciences. The course aims to provide students with preparation regarding analytical and inferential procedures concerning: non-parametric bootstrap, multivariate Gaussian distribution, generalized linear models for count data, and univariate and multivariate Gaussian mixture models, as well as predictive models. The course aims to develop a critical understanding of the model assumptions underlying the theory through empirical applications with real and simulated data. The student also acquires skills related to conducting reproducible and replicable research. In addition, they develop written communication skills, as they are required to produce texts accompanying the results of the analyses carried out.

Knowledge and understanding

This teaching enables the student to:

- Analyze data with advanced univariate and multivariate statistical models developed for both categorical and continuous response variables; implement simulations independently.
- Develop simulation methods.
- Use the semantics of R software, including through the RMarkdown environment, to learn a replicable and reproducible research method. The generated documents include code, results, and comments on the code and performed analyses.
- Interpret the results of the analyses rigorously, developing expressive and summarizing skills, including for dissemination purposes aimed at a non-academic audience. In this way, the student develops independent judgment and refines their communication skills.

Ability to apply knowledge and understanding

The course allows the student to:

- Develop statistical inference using modern bootstrap techniques.

- Estimate, select, and interpret the statistical model-based clustering techniques especially considering finite mixtures of distributions for heterogeneous populations;
- Apply theoretical knowledge to the analysis of data collected in various fields, including epidemiology, medicine, biology, genetics, and public health;
- Conceptualize, and estimate through the maximum likelihood method models with latent variables
- Implement code with the open source software R.

The course enables students to acquire solid theoretical foundations and to develop applications through a "problem-solving" approach. The course pertains to data science, which is now essential for the target job contexts (biostatistics/statistics/demography and related) of graduates in Biostatistics.

Contents

In the first part of the course, the main probability distributions used to simulate realizations from random variables are reviewed. The resampling procedure known as bootstrap is presented to obtain precision measures in a non-parametric context for some estimators of interest.

In the second part of the course, after introducing the multivariate Gaussian distribution, the Expectation-Maximization (EM) algorithm is illustrated in each E and M step as a method for imputing missing data using maximum likelihood estimates of the parameters of a generalized linear model. After introducing Gaussian mixture models, the steps of the EM algorithm for the maximum likelihood estimation of the parameters of these models and latent variable models with discrete distribution are described. Theoretical lessons are accompanied by practical exercises conducted with many different data. The course provides skills in using the semantics of the R software, also utilizing the RMarkdown library through the knitr package to integrate code, analysis results, and comments. At the end of the course, thanks to the provided materials (the instructor's handouts accompanied by an extensive bibliography, code for the R and SAS software, and the RMarkdown interface), the student is able to independently continue deepening their understanding of the subject.

Detailed program

The first part of the course deals with simulation methods and linear congruential methods to generate pseudo-random numbers. Graphical tools for testing the series are illustrated. Simulation of random numbers from specific distributions is considered such the exponential, binomial, and Gaussian distributions.

Resampling methods, such as the jackknife and bootstrap, are introduced in the second part of the course. The bootstrap is applied for bias adjustment and the estimation of dispersion. Bootstrap confidence intervals based on the percentile method and the bias-corrected accelerated bootstrap method are explained.

The autoregressive Poisson model for count data and a similar model based on the negative binomial distribution to account for overdispersion are introduced. These models are applied to monitor endangered species.

Among the optimization methods, the Expectation-Maximization algorithm is considered and explained first as a tool to impute missing values through a generalized linear model and then as a tool to maximize the log-likelihood function for incomplete data problems. Finite mixture models are introduced both for continuous and categorical data, and a particular focus is given to the mixture of Gaussian distributions and latent variable models for categorical data.

Some time is devoted to explaining the theory by imparting the flavor of the empirical applications using data collected from different fields arising in epidemiology, pharmacoepidemiology, medicine and biology, and ecology and environmental sciences. They are developed within the statistical software R, RStudio with the RMarkdown interface. The main R packages used are bootstrap, dplyr, MASS, MultiLCIRT, tscount, mclust e skimr.

The student is encouraged to develop reproducible documents in which he/she critically comments on the code

and the analysis results, also through cooperative learning. Weekly exercises are assigned, and students are encouraged to write reports in which they comment on the code and provide the reader with an explanation of the analysis procedure performed, along with a critical description of the results obtained. Students are invited to work on the assigned exercises in groups to promote cooperative learning. During the course, the solutions to the assigned exercises are discussed.

Prerequisites

For an easier understanding of the course content, it is recommended to know Probability and Statistical Inference notions. The student should also know the basic semantics of the programming language in the R environment

Teaching methods

Lectures are scheduled, and the theory lessons are accompanied by practical exercises that allow students to learn through problem solving by analyzing real and simulated data. The lessons take place in the computer lab. Weekly review exercises related to the covered syllabus are assigned. During the course, with the use of R in the RStudio environment and the RMarkdown interface, students learn to produce reproducible documents that include code, explanations, and comments on the analysis results. They are encouraged to collaborate with each other in solving applied problems in order to promote cooperative learning.

The number of scheduled lecture hours is 30, while interactive teaching hours amount to 17. In the second part of the three-hour lessons, students are actively engaged in an interactive way. The exercises are conducted interactively and in person in the computer lab. Asynchronous video recordings of both the lectures and the exercises are made available on the e-learning platform.

Assessment methods

The following methods of verifying learning apply to both students attending and non-attending lectures in presence. The examination is written with open questions and an optional oral part is possible. There are no intermediate tests, but with the submission of certain exercises, the student can earn two bonus points, which will contribute to the final grade. The written exam has a maximum total duration of two hours and takes place in the computer lab. During the exam, open theory questions must be answered, and exercises must be solved based on the theoretical topics covered during the course. The theory questions assess the understanding of the theoretical concepts taught during the course. The empirical analyses are conducted using the R environment, Rstudio, and RMarkdown allowing verification of the ability to understand the problem and resolve it by applying advanced statistical models to real or simulated data.

The exam also aims to promote students' ability to effectively plan and manage the time needed to write their report. During the exam, the use of study materials and R code developed during the course and personally by the student is allowed. Each part of each exercise is worth approximately 3 points. The student passes the exam with a score of no less than 18 out of 30.

Textbooks and Reading Materials

The teaching material consists mainly of handouts prepared by the teacher. These cover theory, applications, exercise and solutions developed with R software. All the files are available on the course page of the university's e-learning platform. In addition, the teacher publishes at the end of each lesson: the slides, the calculation programs, the exercises, the datasets, and the solutions to the exercises. Previous exam texts are also published on the same page.

The main references are listed in the bibliography of the handouts, some of which are as follows and are available in the university library, also in ebook format:

Bartolucci, F., Farcomeni, A., Pennoni, F. (2013). *Latent Markov Models for longitudinal data*, Chapman and Hall/CRC, Boca Raton.

Bishop, Y. M., Fienberg, S. E., Holland, P. W. (2007). *Discrete multivariate analysis: theory and practice*. Springer Science & Business Media, New York.

Blitzstein, J. K., Hwang, J. (2014). *Introduction to probability*, Chapman & Hall/CRC.

Gentle, J. E., Hardle W., Mori Y. (2004). *Handbook of computational statistics*. Springer-Berlin. Lange, K. (2010). *Numerical analysis for statisticians*, 2nd Edition, Springer, New York.

Pennoni, F. (2025). *Dispensa di Modelli Statistici II, parte di teoria e applicazioni con R*. Dipartimento di Statistica e Metodi Quantitativi, Università degli Studi di Milano-Bicocca.

R Core Team (2022). *R: A Language and Environment for Statistical Computing*. R Foundation for Statistical Computing, Vienna, Austria. <https://www.R-project.org/>.

Semester

Semester I, cycle I, September-November 2025

Teaching language

The course is provided in Italian. Erasmus students can use the handouts material in English and ask the teacher to carry out the exam in English.

Sustainable Development Goals

GOOD HEALTH AND WELL-BEING | REDUCED INEQUALITIES | CLIMATE ACTION
