







Structure : EUR LIFE Code de l'UE : SMUSVPI Lieu d'enseignement : Valrose

Niveau du cours : M1/M2 Semestre : Impair Langue : Anglais

Professors :

- Edoardo SARTI (Edoardo.SARTI@univ-cotedazur.fr)
- Ercan SECKIN (ercan.SECKIN@univ-cotedazur.fr)

Description:

The purpose of this course is to provide the student with the mathematical framework and technical toolkit needed in order to understand advanced topics in machine learning and be able to adapt and refine reference methodologies to meet their scientific needs. Given its brevity, the course does not attempt to provide an exhaustive overview of the wide variety of ML sub-disciplines, but to give the student the tools to autonomously delve deeper in these subjects without losing track of their scientific aim.

Warning : if you are looking for a discursive and exhaustive overview of Machine Learning and/or Deep Learning techniques, this course is <u>not</u> for you : EFELIA offers other resources covering the aforementioned subjects without delving too much into the mathematical background.

Upon successful completion of this course, students will be awarded **3 ECTS**.

This course was co-developed by the EUR LIFE and the EFELIA team (École Française de l'Intelligence Artificielle).

Public :

Students in the first or second year of one of the following Masters : Life Sciences Major : Bioinformatics and Computational Biology, MemBioMed, ImAGEin MSc BOOST: BOOST - Biocontrol for Plant Health

Prerequisites :

The course assumes a good expertise in Python programming (up and included the use of classes and objects), as well as a good understanding of multivariate calculus, linear algebra, and probability. Topics such as – but not limited to - partial derivatives, matrix diagonalization, expectation values will be used with little to no introduction. A self-evaluation test will be available for the student to decide whether or not take this course.

Learning outcomes :

At the end of this course, learners will be able to...

- Clearly explain both the logic and the mathematical framework behind some of the most renown ML methodologies
- Detail the mathematical derivation of some ML algorithms
- Use off-the-shelf implementations of ML algorithms in diverse biomedical contexts
- Correct, modify, adapt, refine existing ML algorithms to meet specific scientific needs
- Analyze, preprocess, and transform data for ensuring an effective and correct use of a target ML method

Teaching methods :

- 1h30 Theory (blackboard + slides)
- 0h30 Exercises (on paper)
- 2h00 Practicals (on machine)

Organisation :

- Présentiel :
 - o 12 heures (CM)
 - 20 heures (TP)

Programme :

| Session | Date | Duration | Professor | Topics |
|---------|------|----------|-------------------------------|---|
| 1 | | | Edoardo Sarti Ercan Seçkin | Complexity and other core concepts ML [1h] - What is Machine Learning? - Essential vocabulary: supervised/unsupervised, regression/classification, etc. - How to approach the study of Machine Learning Towards DL [0.5h] - Bias, variance, and their tradeoff law - Double descent (might not exist) |
| 2 | | | Edoardo Sarti Ercan Seçkin | Linearity is not what it seems ML [1h] - Linear regression, revisited - Polynomial regression Towards DL [0.5h] - Regularizations |
| 3 | | | Edoardo Sarti Ercan Seçkin | Building Neurons ML [1h] - Logistic regression, revisited - Strategies for more than two labels - Separability and linear trick Towards DL [0.5h] - Perceptrons and dense layers |

| | | | SVMs and kernels |
|---|--|-------------------------------|---|
| 4 | | Edoardo Sarti Ercan Seçkin | ML [1h] - Support vector machines - Soft margin Towards DL [0.5h] - Kernel trick |
| | | | Generating with Bayes |
| 5 | | Edoardo Sarti Ercan Seçkin | ML [1h] - Introduction to Bayesian probability - Naïve Bayes and LDA - Recursion: conjugate PDFs Towards DL [0.5h] - Generative Bayesian models |
| | | | The curse of dimensionality |
| 6 | | Edoardo Sarti Ercan Seçkin | ML [1h] - High-dimensional data - Linear dimension reduction: PCA - Dimension reduction: t-SNE, UMAP, etc. Towards DL [0.5h] |
| | | | - Embeddings |
| 7 | | Edoardo Sarti Ercan Seçkin | Clustering ML [1h] - Structured data - K-means - DB scan Towards DL [0.5h] - Expectation-maximization |
| | | | Trees, forests, and crowds |
| 8 | | Edoardo Sarti Ercan Seçkin | ML [1h] - Decision trees - Random forests |

| | | Towards DL [0.5h] - Bootstrap aggregation |
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Assessment : Theoretical questions (50%, 1h) and practical exercise (50%, 2h). More information will be provided at the beginning of the course

Equipment :

• A personal computer is recommended but not required

Bibliography :

- **Mehta et al. 2018** *High-bias, low-variance introduction to Machine Learning for physicists* **[REVIEW]**
- Goodfellow et al. 2016 Deep Learning [TEXTBOOK]
- **Murphy 2012** Machine Learning : a probabilistic perspective **[TEXTBOOK]**
- **Bishop 2006** Pattern recognition and Machine Learning [TEXTBOOK]
- **Tripathy et al. 2021** Unsupervised learning approaches for dimensionality reduction and data visualization [**TEXTBOOK**]
- Friedman et al. 2001 The elements of statistical learning [TEXTBOOK]

Resources for success :

- <u>TUT'TOP</u> : peer tutoring on methodological, social, administrative or logistical issues.
- <u>écri+</u> : to improve your written French.
- <u>Centre de ressources en langues</u> : to improve your foreign language skills (French or other).
- METODA : to improve your documentary research skills.
- <u>S'orienter / Se réorienter</u> : to be advised by the university's career counsellors.
- <u>Centre de santé et aide sociale</u> : to look after your physical and mental health, and to seek support in the event of social hardship.
- <u>Cellule Handicap</u> : support for students with disabilities.
- <u>Plateforme de signalement</u> : to report acts of violence, harassment or discrimination (sexual and gender-based violence, LGBTphobia, racism, xenophobia, etc.) you have witnessed or experienced at the university, and to get support.

Important : This syllabus has no binding value. Its content may change during the course of the year.

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