

SYLLABUS

INTEGRATIVE BIOINFORMATICS

ACADEMIC YEAR 2024-2025

1. Information regarding the programme

1.1 Higher education institution	Babeş-Bolyai University
1.2 Faculty	Faculty of Biology and Geology
1.3 Department	Doctoral School of Integrative Biology
1.4 Field of study	Biology
1.5 Study cycle	Doctorate, 4 years
1.6 Study programme / Qualification	PhD in Biology/ Biologist

2. Information regarding the discipline

2.1 Name of the discipline (en) / (ro)		Integrative Bioinformatics Bioinformatică integrativă					
2.2 Course coordinator		Cojocaru Vlad, PhD <i>Habil.</i>, CS I Banciu Horia Leonard, PhD <i>Habil.</i>, Prof.					
2.3 Seminar coordinator		Cojocaru Vlad, PhD <i>Habil.</i>, CS I Banciu Horia Leonard, PhD <i>Habil.</i>, Prof.					
2.4. Year of study	1	2.5 Semester	1	2.6. Type of evaluation	E	2.7 Type of discipline	O
2.5. Code of the discipline		BDR1106					

E – exam; O – optional

3. Total estimated time (hours/semester of didactic activities)

3.1 Hours per week	4	3.2. Out of which: Lectures	1	3.3 Seminars / Laboratory classes	3
3.4 Total hours in the curriculum	48	3.5. Out of which: Lectures	12	3.6 Seminars / Laboratory classes	36
Allocation of study time:					hrs
Study supported by textbooks, other course materials, recommended bibliography and personal student notes					40
Additional learning activities in the library, on specialized online platforms and in the field					30
Preparation of seminars / laboratory classes, topics, papers, portfolios and essays					30
Tutoring					24
Examinations					3
Other activities: -					40
3.7 Individual study (total hours)	127				
3.8 Total hours per semester	175				
3.9 Number of credits	7				

4. Prerequisites (if necessary)

4.1. Curriculum	<ul style="list-style-type: none"> Genetics, Biochemistry, Biophysics, Cell and molecular biology
4.2. Competencies	<ul style="list-style-type: none"> Average computer skills

5. Conditions (if necessary)

5.1. for the course	<ul style="list-style-type: none"> Beamer, blackboard; Internet connection; Online meeting platform
5.2. for the seminar /lab activities	<ul style="list-style-type: none"> Attendance of at least 4 course/lecture activities. Attendance of all (100%) practical/ seminar classes, Computers, specific development environment

6. Specific competencies acquired

Professional competencies	<ul style="list-style-type: none"> • Ability to use Linux and command line interfaces in life sciences; • Development of the ability to generate, integrate, and analyse sequence data for life sciences; • The ability to use bioinformatics databases, prediction, analysis and visualization tools to model and predict protein structures and protein-nucleic acid complexes; • The ability to use basic methods for molecular dynamics simulations • The ability to use bioinformatics databases, prediction, analysis and visualization tools to infer the diversity and functionality of microbial communities in natural ecosystems; • Development of the capacity for analysis, synthesis and communication of specialized scientific information
Transversal competencies	<ul style="list-style-type: none"> • Acquiring the necessary information to complete a doctoral thesis in Biology field in which generation, processing and analysis of high-throughput data is central. • Carrying out a research project with all that implies the use of specific concepts, the selection and application of study methods, the interpretation of data, to the communication of results.

7. Objectives of the discipline (outcome of the acquired competencies)

7.1 General objective of the discipline	<ul style="list-style-type: none"> • Gaining knowledge on the structure and dynamics of proteins, nucleic acids and other biological macromolecules, as well as diversity and ecological roles of microbial communities using databases and dedicated prediction, analysis and visualization methods.
7.2 Specific objective of the discipline	<ul style="list-style-type: none"> • Developing the capacity to analyze and interpret structures of biomolecules and to link the structures with biological function; • Developing the skills to follow an inter-disciplinary approach to study biomolecules; • Understanding the principles of generation and processing of high-throughput sequence data by NGS; • Use of the main computational strategies for the prediction, modelling, analysis and visualization of community diversity and its ecological functions. • Developing the skills to summarize and disseminate scientific literature in structural and environmental bioinformatics.

8. Content

8.1 Course	Teaching methods	Remarks
Structural Bioinformatics (part I) Introduction to Linux	<ul style="list-style-type: none"> • Interactive exposure • Presentation • Explanation • Practical examples • Case-study discussions 	Hybrid teaching: up to 40% onsite and up to 60% online classes
Structural Bioinformatics (part II) Visualization and analysis of biomolecular structures (e.g. proteins, nucleic acids)		
Structural Bioinformatics (part III) Molecular modelling (prediction of biomolecular structures, structure-based design)		
Structural Bioinformatics (part IV) Molecular docking, molecular dynamics simulations, free		

energy calculations		
Environmental Bioinformatics Introduction to genomic sequencing technologies. Quality control and read alignment. Genome assembly and comparison of genome assembly methods. Genome visualization tools. Dedicated genomic databases. Genome annotation. Taxonomic prediction of genomes	<ul style="list-style-type: none"> • Interactive exposure • Presentation • Explanation • Practical examples • Case-study discussions 	
Bibliography 1. Keith J.M., Bioinformatics. Vol. 1: Data, sequence analysis, and evolution. New York : Humana Press, 2017.. In: Bioinformatics, vol. Vol. 1 2. Keith J.M., Bioinformatics. Vol. 2: Structure, function, and applications. New York : Humana Press, 2017.. In: Bioinformatics, vol. Vol. 2, 3. Leach, A.R. Molecular modelling: principles and applications. Pearson education.2001. 4. Pevzner P., Bioinformatics for biologists. Cambridge ; New York : Cambridge University Press, 2013 5. Stryer L., Biochemistry. New York : W. H. Freeman and Company, 1995\ 6. Schlick T., Molecular modeling and simulation : an interdisciplinary guide. New York, Springer, 2010. 7. Xiong J., Essential bioinformatics. New York : Cambridge University Press, 2006 References (1-2, 4-5, 7) are available in printed format at the libraries of the Faculty of Biology and Geology. Reference (3) is available upon request from the class tutor. Reference (6) is available in printed format at the library of the Faculty of Chemistry and Chemical Engineering.		
8.2 Seminar / laboratory	Teaching methods	Remarks
Study cases and exercises in structural bioinformatics (Modeling of three-dimensional structures of biomacromolecules; Comparison of 3D structures; Modelling and visualization of molecular dynamics; Data validation, integration and comparison)	Interactive explanations Explanation Conversation Case study	Hybrid teaching: up to 50% onsite and up to 50% online classes
Case studies and exercises in environmental bioinformatics: analysis of model genomes of microorganisms using online bioinformatics tools		
Presentation of a relevant scientific article	Active learning (student-led seminar)	
Preparation of a short individual project on structural or environmental bioinformatics	Evaluation	2 hours
Bibliography Collection of research articles available in digital format at the libraries of the Faculty of Biology and Geology and `Lucian Blaga` Central University Library, Cluj-Napoca.		

9. Corroborating the content of the discipline with the expectations of the epistemic community, professional associations and representative employers within the field of the program

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| <ul style="list-style-type: none"> • The course promotes the gaining of theoretical knowledge and practical skills required for teamwork in the field of research and development in academic and research entities, but also in R&D units in private companies; • The course is present in the curriculum of similar specializations at USA and European Universities. |
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10. Evaluation

Type of activity	10.1, Evaluation criteria	10.2, Evaluation methods	10.3 Share in the grade/qualification (%)
10.3. Course	Knowledge of concepts and methods from the topics of the course	Written exam (combined test)	40%
10.4. Seminar/lab activities	Evaluation of -presentation of a research article; - individual project on structural bioinformatics topic	Oral colloquium	20% 40%
10.5. Minimum performance standards			
Each student should obtain a minimum qualification of "SUFFICIENT" (from the following list of available qualifications: INSUFFICIENT / SUFFICIENT / GOOD / VERY GOOD) at the written exam and the oral colloquium. To obtain the qualification "SUFFICIENT", the student must demonstrate the mastery of the basic concepts described during the course and practicum classes.			

Date
29.07.2024

Signature of course coordinator
COJOCARU Vlad, PhD *Habil.*, CS I

Signature of seminar coordinator
COJOCARU Vlad, PhD *Habil.*, CS I

BANCIU Horia, PhD *Habil.*, Prof.

BANCIU Horia, PhD *Habil.*, Prof.

Date of approval
29.07.2024

Signature of the head of department
Prof. dr. *Habil.* PAP Peter Laszlo