

## SYLLABUS

### 1. Information regarding the programme

1.1 Higher education institution	<b>Babeş Bolyai University</b>
1.2 Faculty	<b>Faculty of Biology and Geology</b>
1.3 Department	<b>Department of Molecular Biology and Biotechnology</b>
1.4 Field of study	<b>Biology</b>
1.5 Study cycle	<b>Master</b>
1.6 Study programme / Qualification	<b>Bioinformatics applied in life sciences</b>

### 2. Information regarding the discipline

2.1 Name of the discipline (en) (ro)	<b>Command line scripting</b> <b>Codarea liniilor de comandă</b>						
2.2 Course coordinator	<b>Dr. Dorin Mirel Popescu</b> <b>Andrei Alexsson</b>						
2.3 Seminar coordinator	<b>Dr. Dorin Mirel Popescu</b> <b>Andrei Alexsson</b>						
2.4. Year of study	<b>1</b>	2.5 Semester	<b>I</b>	2.6. Type of evaluation	<b>E</b>	2.7 Type of discipline	<b>Elective</b>
2.8 Code of the discipline	<b>BME1116</b>						

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

3.1 Hours per week	<b>4</b>	Of which: 3.2 course	<b>2</b>	3.3 seminar/laboratory	<b>2</b>
3.4 Total hours in the curriculum	<b>56</b>	Of which: 3.5 course	<b>28</b>	3.6 seminar/laboratory	<b>28</b>
Time allotment:					hours
Learning using manual, course support, bibliography, course notes					34
Additional documentation (in libraries, on electronic platforms, field documentation)					30
Preparation for seminars/labs, homework, papers, portfolios and essays					20
Tutorship					10
Evaluations					4
Other activities: .....					-
3.7 Total individual study hours	<b>98</b>				
3.8 Total hours per semester	<b>154</b>				
3.9 Number of ECTS credits	<b>6</b>				

### 4. Prerequisites (if necessary)

4.1. curriculum	<ul style="list-style-type: none"> <li>• Basic algebra calculation skills</li> </ul>
4.2. competencies	<ul style="list-style-type: none"> <li>• Average programming skills</li> </ul>

### 5. Conditions (if necessary)

5.1. for the course	<ul style="list-style-type: none"> <li>• Beamer</li> <li>• Online meeting platform</li> </ul>
5.2. for the seminar /lab activities	<ul style="list-style-type: none"> <li>• Attendance of a minimum of 90% of practical/ seminar classes,</li> <li>• Computers, specific development environment</li> </ul>

### 6. Specific competencies acquired

<b>Professional competencies</b>	<ul style="list-style-type: none"> <li>• Ability to navigate and use the command line in a Linux environment for accessing and analyzing biological data efficiently.</li> <li>• Skills in executing commands, managing files, and automating tasks using scripts.</li> <li>• Ability to manipulate biological data files for preparing data for further analysis and visualization of large-scale genomic datasets.</li> <li>• Skills in writing and executing BASH scripts to automate data processing tasks and to increase workflow efficiency through improved reproducibility and scalability in data analysis.</li> </ul>
<b>Transversal competencies</b>	<ul style="list-style-type: none"> <li>• Capacity to analyze complex biological problems, break them down into smaller components, and devise effective solutions through means of logical reasoning, creativity, and adaptability</li> <li>• Ability to work effectively in multidisciplinary teams, share and document scripts and workflows, and communicate findings clearly to both technical and non-technical audience.</li> </ul>

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

7.1 General objective of the discipline	<ul style="list-style-type: none"> <li>• To train students on practical skills for efficiently working with biological data by introducing them to Linux and command-line basics, teaching scripting and automation, and demonstrating how to manipulate biological data files using study cases</li> </ul>
7.2 Specific objective of the discipline	<ul style="list-style-type: none"> <li>• To familiarize students with the Linux operating system, show them how to navigate the command line, execute basic and advanced commands, and understand the file system structure</li> <li>• To train students in writing and executing simple Bash scripts</li> <li>• To provide hands-on exercises using Linux commands to handle biological data files including data export into other environments for further analysis.</li> <li>• To apply command-line skills on real biological data examples and specific research contexts</li> </ul>

## 8. Content

8.1 Course	Teaching methods	Remarks
Introduction to Bash and the Command Line (Basic Shell Commands)	<ul style="list-style-type: none"> <li>• Presentation</li> <li>• Explanation</li> <li>• Conceptualized-learning</li> <li>• Active learning through case-study discussions</li> </ul>	
Variables, Arrays and Parameters		
Shell Scripting Basics - creating and running shell scripts		
Control Structures (conditional statements and looping)		
File Permissions and Ownership		
Functions in Bash		
Working with Text Files (awk, vim, sed)		
Regular expressions		
Input and output redirection, Error handling and debugging		
Managing processes (ps, top, kill, bg, fg, etc)		
Integrating bash with other programming languages		
Running bioinformatics tools in bash		
Bibliography <ol style="list-style-type: none"> <li>1. Bash Reference Manual, freely available at <a href="https://www.gnu.org/software/bash/manual/bash.html">https://www.gnu.org/software/bash/manual/bash.html</a></li> <li>2. Linux Handbook, <a href="https://linuxhandbook.com/">https://linuxhandbook.com/</a></li> <li>3. The Bash Guide, freely available at <a href="https://guide.bash.academy/">https://guide.bash.academy/</a></li> </ol>		

8.2 Seminar / laboratory	Teaching methods	Remarks
Real-world applications and case studies	<ul style="list-style-type: none"> <li>• Interactive discussion</li> <li>• Explanation</li> <li>• Conversation</li> <li>• Project-based learning</li> <li>• Practical demonstration</li> </ul>	
Team-work or individual assignments focused on creating scripts for various bioinformatic applications (data retrieval from databases, alignments, etc.)		
Students project presentations, discussions and conclusions		
Final evaluation		
Bibliography <ol style="list-style-type: none"> <li>1. Printed hands-on provided by the trainers</li> <li>2. Bash Shell Scripting Practice Exercises, available at: <a href="https://linuxhandbook.com/practice/bash/">https://linuxhandbook.com/practice/bash/</a></li> <li>3. w3resource's Bash Scripting Exercises, available at: <a href="https://www.w3resource.com/bash-script-exercises/index.php">https://www.w3resource.com/bash-script-exercises/index.php</a></li> </ol>		

**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**

<ul style="list-style-type: none"> <li>• The course is present in the curriculum of many universities in the world.</li> <li>• The course enables the developing fundamental skills in efficient handling and analysis of large-scale biological data, automating repetitive tasks and enhancing reproducibility through scripting. .</li> </ul>
--

**10. Evaluation**

Type of activity	10.1 Evaluation criteria	10.2 Evaluation methods	10.3 Share in the grade (%)
10.1. Course	Know concepts and methods from the topics of the course	Written exam	50%
10.2. Seminar/lab activities	Apply tools and concepts of scripting in real-life problems	Solving an individual assignment and presenting/explaining the solutions of the assigned project.	50%

10.3. Minimum performance standards
Each student should obtain a minimal mark of 5 (five) for the written exam and for the individual assignment. To obtain the minimum grade 5, the student must demonstrate the mastery of the basic concepts described during the course and seminars/practical classes, respectively.

Date                                      Signature of course coordinator                                      Signature of seminar coordinator

22.07.2024

**Dr. Dorin Mirel Popescu**

**Dr. Dorin Mirel Popescu**

**Andrei Alexsson**

**Andrei Alexsson**

Date of approval

23.07.2024

Signature of the head of department  
**Assoc. Prof. Beatrice Kelemen, PhD**