

DISCRETE MATHEMATICS

| Course Workload | | Assessment form (examination/ graded test/ |
|-----------------|-------|--|
| ECTS | Hours | ungraded test) |
| 3 | 108 | Exam |

The course will introduce students to the basic concepts of Boolean logic, combinatorics, dynamic programming and algorithms on graphs, which are the basis for solving many bioinformatics problems.

Course structure:

- 1. INTRODUCTION TO SET THEORY, BOOLEAN FUNCTIONS
- 1.1. Introduction to set theory: definitions, set operations and properties.
- 1.2. Boolean functions, boolean logic and proof methods.
- 2. COMBINATORICS
- 2.1. Combinations, permutations, pigeonhole principle.
- 2.2. Combinatorial generation.
- 3. ASYMPTOTIC ANALYSIS AND SORTING ALGORITHMS
- 3.1. Asymptotic analysis of algorithms.
- 3.2. Sorting algorithms.
- 4. DYNAMIC PROGRAMMING
- 4.1. Definitions, subtask optimization, basic problems.
- 4.2. Dynamic programming in bioinformatics: local and global alignments.
- 4.3. RNA's secondary structure prediction.
- 5. GRAPH THEORY
- 5.1. Definitions, DFS, BFS.
- 5.2. Algorithms, de Bruijn graph.