

| COURSE RECORD | |
|------------------------------------|---|
| Code | BENG619 |
| Name | Proteomics and Metabolomics |
| Hour per week | 3+0 (Theory + Practice) |
| Credit | 3 |
| ECTS | 10 |
| Level/Year | Graduate |
| Semester | Spring |
| Туре | Elective |
| Location | |
| Prerequisites | |
| Special Conditions | |
| Coordinator(s) | |
| Webpage | |
| Content | Introduction to proteome and proteomics technology. General workflow for bottom-up and top-down proteomic approaches. Exploration of differential protein expression, post-translational modifications and protein-protein interactions (PPI). Introduction to metabolome and metabolomics. Metabolite identification, pathway identification and pathway integration. |
| Objectives | Learn principles of the most common proteomics techniques Understand the mass spectrometry based proteomics workflow Learn experimental design, sample preparation and enrichment of techniques. Design their own research project by proteomics and metabolomics techniques. |
| Learning Outcomes | LO1 Students completing this course will be able to learn modern strategies of proteomics and metabolomics. LO2 Students completing this course will be able to evaluate strengths and limitations of mass spectrometry based proteomics workflows. LO3 During this class students will be able to participate in group discussions LO4 Student will be able to practice writing process of a research proposal. |
| Requirements | You need to read assigned research articles before class and participate in- group discussion. You will need write and present a relevant research proposal at the end of the semester. |
| Reading List | Mass Spectrometry for the Novice, John Greaves and John Roboz, CRC Press 2013. Introducing Proteomics: From Concepts to Sample Separation, Mass Spectrometry and Data Analysis, Josip Lovric, Wiley 2011. |
| Ethical Rules and Course Policy | |

| LEARNING ACTIVITIES | Please | use this | one as a | reference | for | vour | course |
|---------------------|--------|----------|----------|-----------|-----|------|--------|
| LEARNING ACTIVITIES | reuse, | use uns | one us u | rejerence | 101 | vour | course |

| Activities | Number | Weight (%) | | | | |
|---------------|--------|------------|--|--|--|--|
| Lecture | 26 | 50% | | | | |
| Group Works | 13 | 25% | | | | |
| Presentations | 1 | 25% | | | | |
| Site Visits | 0 | 0 | | | | |
| | Tota | l 100 | | | | |

| Weight (%) |
|------------|
| |

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| Quizzes | 0 % |
|---|------------|
| Weekly Assignments | 10% |
| Group Project Assignments & Presentations | 20% |
| Attendance/Participation | 10% |
| Midterm | 20% |
| Final Exam/Submission | 40% |
| | Total 100% |

For a detailed description of grading policy and scale, please refer to the website https://goo.gl/HbPM2y section 28.



COURSE LOAD Please, use this one as a reference for your course

| Activity | Duration (hour) | Quantity | Work Load (hour) |
|---------------------------|---------------------------|--------------------|----------------------------|
| In class activities | 3 | 14 | 42 |
| Lab (Computer Lab) | 3 | 3 | 9 |
| Group work | 1 | 13 | 13 |
| Research (web, library) | 5 | 13 | 65 |
| Required Readings | 8 | 13 | 104 |
| Pre-work for Presentation | 10 | 3 | 30 |
| Lab reports | 0 | 0 | 0 |
| | | General Sum | 263 |

ECTS: 10 (Work Load/25-30)

CONTRIBUTION TO PROGRAMME OUTCOMES*

| | P01 | P02 | P03 | P04 | P05 | P06 | P07 | P08 | P09 | P010 | P011 | P012 | P013 | P014 |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|------|------|------|------|
| L01 | 4 | 5 | 3 | 2 | 1 | 0 | 0 | 0 | | | | | | |
| L02 | 4 | 4 | 5 | 1 | 1 | 0 | 0 | 3 | | | | | | |
| L03 | 4 | 5 | 4 | 3 | 5 | 5 | 3 | 1 | | | | | | |
| L04 | 4 | 5 | 5 | 3 | 5 | 5 | 2 | 3 | | | | | | |

* Contribution Level: 0: None, 1: Very Low, 2: Low, 3: Medium, 4: High, 5: Very High

WEEKLY SCHEDULE

| W | Торіс | Outcomes |
|----|--|----------------|
| 1 | Introduction to proteomics | L01 |
| | Activity: Group discussion | |
| 2 | Experimental strategies in proteomics | L01, L02, L04 |
| | Activity: Group discussion | |
| 3 | Basic principles of mass spectrometry based proteomics | LO1, LO2, LO4 |
| | Activity: Group discussion | |
| 4 | Interpretation of mass spectrometry data | LO1, LO2, LO4 |
| | Activity: Data analysis | |
| 5 | Qualitative proteomics: Protein/peptide identification | LO1, LO2, LO4 |
| | Activity: Data analysis | |
| 6 | Quantitative proteomics I | L01, L02, L04 |
| | Activity: Data analysis | |
| 7 | Quantitative proteomics II | L01, L02, L04 |
| | Activity: Group discussion | |
| 8 | Analysis of post translational modifications | LO1, LO2, LO3, |
| | Activity: Group discussion | LO4 |
| 9 | Top-Down proteomics | L01, L02, L03, |
| | Activity: Group discussion | LO4 |
| 10 | Proteomics for structural biology | L01, L02, L03, |
| | Activity: Group discussion | LO4 |
| 11 | Imaging Mass spectrometry | L01, L02, L03, |
| | Activity: Group discussion | LO4 |
| 12 | Interaction proteomics | LO1, LO2, LO3, |
| | Activity: Group discussion | LO4 |
| 13 | Student Presentations | L01, L02, L03, |
| | Activity: Group discussion | LO4 |
| 14 | Student Presentations | L02, L04 |
| | Activity: Group discussion | |

Prepared by Date