

Course number: CHEM630/CRB 630
Course Title: Proteomics Approaches for Biologists
Spring 2018

Course Instructor:

Professor Ying Ge, Office, WIMR II 8551; Phone: 608- 265-4744. email: ge2@wisc.edu

Lecture time: Wed. and Fri. 8:50- 9:40 am (2 credits)

Office hours: 1:30-3:30 PM on Fridays or by appointment

Classroom: WIMR II 8th floor conference room (Room 8571)

Course overview:

Proteomics and metabolomics are playing an increasingly important role in biology and medicine. Many biology and Chembio labs are now starting to use proteomics and metabolomics in their research projects. This course is designed specifically for students in chemical/biological sciences who have interests to learn proteomics and metabolomics. It will integrate formal classroom lectures with one-on-one consultation. Lectures include the essential fundamentals and applications in mass spectrometry-based proteomics and metabolomics to address biochemical/biological/medical problems. Meanwhile, one-on-one consultation will be offered to respond to students' individual needs, including the design of proteomics/metabolomics experiments, troubleshooting, and proper interpretation of the results.

Topics to be covered in the course include:

- Introduction to mass spectrometry-based proteomics
- Proteomics for decoding signaling network
- Proteomics for structural biology
- Top-down proteomics
- Sample preparation for proteomics experiments
- Proteomics strategy for identification of kinase substrate
- Proteomics in clinical applications
- Mass spectrometry based metabolomics

The class will meet twice a week (Wed. and Fri. 50 min). Classroom sessions involve one of two formats: 1) traditional lectures interspersed with classroom discussion; 2) student presentation. The essential knowledge and practical application of proteomics and metabolomics will be taught during the traditional lectures. Towards the end of the course, the students will present a specific topic of interest on proteomics/metabolomics and discuss their specific proteomics/metabolomics needs in their thesis projects. The student can also choose to write a proposal on a specific proteomics/metabolomics subject which is not related to their thesis project. Along with the regular classroom instruction, students are offered ample one-on-one individual consultations with the course instructor to discuss their specific proteomics/metabolomics needs. Major goals of the course include accruing fundamental knowledge and understanding of proteomics/metabolomics, developing confidence in proteomic/metabolomic experimental design and data interpretation as well as critical thinking and oral presentation skills.

Grades: Grading will be based on the three course components; attendance and participation in the discussion (40%), two homework sets (30%), and final presentation on a proteomics/metabolomics related project or a proposal on a proteomics/metabolomics topic (30%). Grading scale for the course, A-F (92-100=A, 89-91=AB, 82-88% = B, 79-81% = BC, 72-78% = C, 69-71% = CD, 60-68% = D, Below 60% = F.)

Below is a tentative schedule of topics:

Week	Lecture Topics
1	Course Overview
2	The Nuts and Bolts of Mass Spectrometry-based Proteomics (Part I)
3	The Nuts and Bolts of Mass Spectrometry-based Proteomics (Part II) <i>Homework #1 handout</i>
4	Proteomics for Decoding Signaling Network (Part I)
5	Proteomics for Decoding Signaling Network (Part II) <i>Homework #1 Due</i>
6	Proteomics for Structural Biology (Part I)
7	Proteomics for Structural Biology (Part II) <i>Homework #2 handout</i>
8	Top-down Proteomics (Part I)
9	Top-down Proteomics (Part II)
10	Sample preparation for MS-based proteomics <i>Homework #2 handout</i>
11	Identification of Kinase Substrate by Mass Spectrometry- based Proteomics
12	Proteomics in Clinical Applications
13	Mass Spectrometry-based Metabolomics
14	Student Presentation
15	Student Presentation
16	Student Presentation