



BB2510 Proteomik 6,0 hp

Proteomics

När kurs inte längre ges har student möjlighet att examineras under ytterligare två läsår.

Fastställande

Kursplan för BB2510 gäller från och med HT14

Betygsskala

A, B, C, D, E, FX, F

Utbildningsnivå

Avancerad nivå

Huvudområden

Bioteknik

Särskild behörighet

At least 150 credits from grades 1, 2 and 3 of which at least 100 credits from years 1 and 2, and bachelor's work must be completed. The 150 credits should include a minimum of 20 credits within the fields of Mathematics, Numerical Analysis and Computer Sciences, 5 of these must be within the fields of Numerical Analysis and Computer Sciences, 20 credits of Chemistry, possibly including courses in Chemical Measuring Techniques and 20 credits of Biotechnology or Molecular Biology.

Knowledge corresponding to the following courses, or similar, is required.

BB1010 Introduction to Biotechnology

BB1020 Cell biology with Immunology

BB1080 Biochemistry, Theory

BB1110 Gene Technology and Molecular Biology

BB1130 Analysis and Purification of Biomolecules

Undervisningspråk

Undervisningspråk anges i kurstillfällesinformationen i kurs- och programkatalogen.

Lärandemål

The aim of this course is to provide the students with an introduction into methodologies in the field of proteomics. Therein, examples of current trends for global protein analysis will be presented from different perspectives in order to demonstrate complexity and terminologies of this field of research. The course will provide an overview of typical proteomics applications used today and present their use for different applications.

After having completed this course, the students shall be able to

- describe and discuss the possibilities offered by proteomics technologies
- judge and compare the various proteomics approaches
- suggest and reason for suitable strategies for specified projects.
- critically evaluate results from proteomic studies.
- participate in scientific discussions regarding proteomics technologies
- speculate and argue about coming trends of proteomics technologies.

Kursinnehåll

The course is focused on different methods, technologies and strategies currently applied in the field of proteomics. General concepts as well as an emphasis on biomarker discovery will be presented in order to demonstrate proteomic application, which span different areas in life science and biotechnology.

Kursupplägg

The lectures will cover the different disciplines within proteomics by providing insights from researchers within the field as well as the perspectives of PhD students working with proteomics. Background on classical proteomics, such as 2D-gel electrophoresis or mass spectrometry, will be complemented by recent developments and strategies built on affinity proteomics, the high-throughput generation of binding reagents, data analysis as well as structural and interaction approaches.

Kurslitteratur

- Principles of Proteomics by R.M Twyman, Garland Science, ISBN: 9780815344728 (2013).
- Review articles and
- Handouts from presentations are distributed at each lecture.

Examination

- TEN1 - Skriftlig tentamen, 5,0 hp, betygsskala: A, B, C, D, E, FX, F
- ÖVN1 - Övning, 1,0 hp, betygsskala: P, F

Examinator beslutar, baserat på rekommendation från KTH:s handläggare av stöd till studenter med funktionsnedsättning, om eventuell anpassad examination för studenter med dokumenterad, varaktig funktionsnedsättning.

Examinator får medge annan examinationsform vid omexamination av enstaka studenter.

The examination consists an oral and a written part. The oral exam will in form of a panel discussion where groups of students will prepare arguments for a discussion on a given proteomic subject. One group will argue for the possibilities and advantages for a certain method or strategy and another group will then argue against this with drawbacks and limitations. It is important to see this part as not only an examination, but also as a learning activity to argue for your scientific perspective and to collect information from the lectures and other sources.

The written examination will mostly consists of questions, where the students are expected to describe and discuss the obtained knowledge about proteomics according to the aims of the course. The students will also be able to express your scientific opinion regarding various aspects of proteomics technologies. There is no right or wrong in the provided opinion, the students you will be able to demonstrate their knowledge about proteomics based on their argumentation.

Övriga krav för slutbetyg

A written examination (TEN1, 5 hp, grading scale A-F).

Participation in panel discussion (ÖVN1, 1 hp, Pass/Fail)

Etiskt förhållningssätt

- Vid grupparbete har alla i gruppen ansvar för gruppens arbete.
- Vid examination ska varje student ärligt redovisa hjälp som erhållits och källor som använts.
- Vid muntlig examination ska varje student kunna redogöra för hela uppgiften och hela lösningen.