

<b>Level:</b> master				
<b>Course title:</b> Rational drug design (IB-503)				
<b>Status:</b> elective				
<b>ECTS:</b> 7				
<b>Requirements:</b> none				
<b>Learning objectives</b> Acquaintance with modern methods for the rational design of new biologically active molecules, potential drugs.				
<b>Learning outcomes</b> Upon successful completion of this course, the student is able to use the selected “Open Source” software for the rational drug design.				
<b>Syllabus</b> <i>Theoretical instruction</i> Chemical and biochemical databases of interest for the development of new drugs. Molecular recognition as the basis for rational drug design. Analysis of protein-ligand interactions. The process of drug design when the structure of the target protein is unknown and pharmacophore modelling. Examples of molecular mimicry of known drugs and biomolecules. Drug design when the structure of the target protein is known (structure based design). Molecular docking and virtual screening. Rational design of HIV-1 protease inhibitors.  <i>Practical instruction</i> Identification and visualization of pharmacophore and ligand receptor interactions in Accelrys Discovery Studio Visualizer. Molecular docking using AutoDock and AutoDock Vina.				
<b>Weekly teaching load</b>				<b>Other:</b>
Lectures: 2	Exercises: 2	Other forms of teaching: 1	Student research:	