

<b>Modulcode</b>	<b>biol258</b>
<b>number online registration</b>	
<b>Module number</b>	<b>MM10</b>
<b>Module name</b>	<b>Computational and Comparative Genomics</b>
<b>Module name - german</b>	
<b>Program of Study</b>	MSc Compulsory Module AgriGenomics MSc Biology
<b>Offered</b>	Once a year, winter semester
<b>Module coordinator</b>	Prof. Dr. Tal Dagan
<b>Module advisors</b>	Prof. Dr. Tal Dagan
<b>Courses and teachers</b>	<b>Lecture:</b> Computational and Comparative Genomics: (Prof. Dr. Dagan, Dr. Ke Xiau, Dr. Georg Hemmrich, Dr. David Ellinghaus, Dr. Giddy Landan) <b>Exercises:</b> Computational and Comparative Genomics: Prof. Dr. Dagan, Dr. Ke Xiau, Dr. Georg Hemmrich, Dr. David Ellinghaus, Dr. Giddy Landan
<b>Prerequisites</b>	Advanced understanding of genetics, molecular biology, animal and plant breeding as well as plant nutrition and phytopathology
<b>Language</b>	English
<b>Module capacity on campus students</b>	40 (20 places are allocated to the AgriGenomics program) enrollment from November 1 to November 30 via OLAT
<b>Module capacity off campus students</b>	0
<b>Course types (classroom/ total workload)</b>	Lecture (30 h/90 h), exercise (90 h/90 h)
<b>Schedule</b>	block course (UNIVIS)
<b>Grading</b>	Written test 100 % (K)
<b>ID-card</b>	Required for exams
<b>European Credit Points</b>	6
<b>Module Objectives</b>	The module is aimed at teaching basic methods for the analysis of genomic data. This includes an overview of the theory and practice of computational methods for the identification and characterization of genetic elements from DNA sequence data. The course focuses on approaches for extracting the maximum amount of information from protein and DNA sequence similarity through sequence database searches, statistical analysis, and multiple sequence alignment.
<b>Contents</b>	Genomic data mining, sequence comparison, phylogenetic trees, protein domain prediction, genome sequencing and assembly, genome annotation, identification of genomic structural variants, transcriptomics.
<b>Taught Skills</b>	Basic knowledge in the analysis of genomic and transcriptomic data.
<b>Course materials</b>	Praktikumsskript, Manuals, Videos