

<b>Module number</b>	<b>MM2</b>
<b>Module name</b> <b>Program of study</b>	<b>Organization and analysis of eukaryotic genomes</b> MSc Mandatory Module
<b>Offered</b>	Once a year, winter semester
<b>Module coordinator</b>	Prof. Dr. Christian Jung
<b>Module advisor</b>	Prof. Dr. Christian Jung
<b>Courses and teachers</b>	<b>Lectures:</b> Organization of the eucaryotic genome (Prof. Dr. C. Jung with Dr. Carlos Molina) Genome analysis I, structural genome analysis (Prof. Dr. G. Thaller by Dr. J. Tetens) Genome analysis II, functional genome analysis (Prof. Dr. D. Cai)
<b>Prerequisites</b>	Fundamental knowledge in molecular biology, molecular genetics and gene technology
<b>Language</b>	English
<b>Module capacity on campus students</b>	20
<b>Module capacity off campus students</b>	-
<b>Course types (classroom/ total workload)</b>	Lecture (15 h/45 h), lecture (22,5 h/67,5 h), lecture (22,5 h/67,5 h)
<b>Schedule</b>	
<b>Grading</b>	Oral examination: 100% (C. Jung, J. Tetens)
<b>ID-card</b>	Required for exams
<b>European Credit Points</b>	6
<b>Module Objectives</b>	The students understand the structure and evolution of plant and animal genomes. They know the major components of complex eukaryotic genomes. They learn the relevant techniques for structural and functional analysis of plant and animal genomes. They understand how to sequence genomes and to analyze complex genomic sequences.
<b>Contents</b>	structure and evolution of plant and animal genomes, techniques for analyzing eucaryotic genomes, mapping, gene identification, genome sequencing, sequence analysis
<b>Taught skills</b>	Methodical responsibility, key qualifications
<b>Course materials</b>	Textbooks, lecture notes, internet