Identificat Number MN-B-SM (C 4) 1	tion Course a) Lect b) Prac c) Sem Modul Studer	ctical/Lab ninar e Objectives	Credit Points 12 CP	18 h 90 h	rm of ing act Time	Offe Sum	ered Every nmer term, half Private Sto	only	ner term	Duration 7 weeks				
Number MN-B-SM (C 4) 1 2	Course a) Lect b) Prac c) Serr Modul Studer	360 h e Types tures ctical/Lab hinar e Objectives	Points	2 nd ter studyi Conta 18 h 90 h	rm of ing act Time	Sum	nmer term, half Private Stu	summ only						
(C 4) 1 2	Course a) Lect b) Prac c) Sem Modul Studer	e Types tures ctical/Lab hinar e Objectives	12 CP	studyi Conta 18 h 90 h	act Time		half Private Sto	only		7 weeks				
2	a) Lect b) Prac c) Sem Modul Studer	tures otical/Lab ninar e Objectives		18 h 90 h				udy		7 weeks				
2	b) Prac c) Sem Modul Studen	ctical/Lab ninar e Objectives		90 h					Planned	Group Size				
2	c) Sem Modul Studer	ninar e Objectives					36 h		max. 6					
2	Modul Studer	e Objectives				156 h			max. 6					
	Studer	•		24 h		36 h		max. 6						
			Module Objectives and Skills to be Acquired											
		Students who successfully completed this module												
	have gained in-depth knowledge in advanced sequence analysis and its application to real-life problems in the experimental biosciences.													
	have acquired skills to identify bioinformatically tractable problems relevant for experimental projects and are familiar with the selection and application of appropriate bioinformatical tools.													
	 have learned how large-scale databases (genomics, transcriptomics, proteomics etc.) can be mined for information useful to solve biological problems, and how bioinformatical tools can be used to extract and process data obtained from these repositories. 													
	• Have learned how to write a scientific publication (or similar text), using the bioinformatical results obtained in this course as an example.													
	have learned how to present research results in oral and written form and to critically discuss scientific publications related to the topic of the module on a professional level.													
	are able to transfer skills acquired in this module to other fields of biology.													
3	Module Content													
	Methods of biological sequence analysis, predicting the function, structure and other properties of protein based on its sequence.													
	• Public databases containing genomics, transcriptomics, proteomics and other large-scale data.													
		•	multiple data		•					and function				
	 Protein evolution, analysis of conservation patterns and their application to structure and function prediction. 													
	Assessment and validation of bioinformatical predictions													

Applied Bioinformatics – From Sequence Analysis to Publication (MN-B-SM [C 4]) continued

4	Teaching Methods
	The module will be centered around a topical biological problem, suitable for being addressed by bioinformatical methods. Guided by the instructors, the students will be working in small groups on various aspects of the problems, covering a wide range of bioinformatical methods and databases. Twice weekly, the results obtained by the groups will be presented, discussed and analyzed in the plenum. Also, new tasks will be assigned. In the last two weeks, the focus will be on documenting the line of reasoning and the results as scientific figures, tables and text. In a joint effort, the results and analyses will be converted into a manuscript-style document – suitable for publication on BioRxiv.
	Teaching forms: Lectures (biological background, selected methods) Practical/Lab & Guidance to independent research (supervised project work performed in small groups) Seminar/Oral presentation techniques (Presentation of group results) Protocol/Written presentation techniques (Generation of a manuscript-style document)
5	Prerequisites (for the Module)
	Enrollment in the Master's degree course "Biological Sciences".
	Additional academic requirements
	Previous attendance of the lecture module "Computational Biology (C)".
	Previous attendance of the seminar module "Computational Biology (C)" is highly desirable, but lack of attendance can be compensated by some familiarity with the LINUX operating system and the application of sequence analysis methods. Programming skills are not required for this module.
6	Type of Examination
	The final examination consists of two parts: oral presentation (20-30 min; 50 % of the total module mark), written report (50 % of the total module mark)
7	Credits Awarded
	Regular and active participation; Each examination part at least "sufficient" (see appendix of the examination regulations for details)
8	Compatibility with other Curricula
	None.
9	Proportion of Final Grade
	In the Master's degree course "Biological Sciences": 15 % of the overall grade (see also appendix of the examination regulations)
10	Module Coordinator
	Prof. Dr. Kay Hofmann, phone 470-1701, e-mail: kay.hofmann@uni-koeln.de

Applied Bioinformatics – From Sequence Analysis to Publication (MN-B-SM [C 4]) continued

11	Further Information							
	Subject module of the Master's degree course "Biological Sciences", Specialization: (C) Computational Biology							
	Participating faculty: Prof. Dr. Kay Hofmann, Dr. Karsten Klopffleisch.							
	Literature: Information about textbooks and other reading material will be given on the ILIAS representation of the course (https://www.ilias.uni-koeln.de/ilias/ goto_uk_cat_2815610.html)							
	General time schedule: Week 1-5 (MonFri.): Lectures, practical/lab and seminar presentation of group results (topic and date will be arranged individually); Week 6 (MonFri.): Seminar talks and finalization of the result document; Week 7 (MonFri): Preparation and finalisation of written report.							
	Note: The module does not contain hands-on laboratory work. The module contains computer-based practicals/research as a main component.							
	Introduction to the module: May 16, 2022 at 10:00 a.m., Center for Molecular Biosciences (COMB), Seminarroom 0.46 (ground floor) or online (in this case, further information/link will be sent to your Smail-Account); for preparation to the module before this introduction see ILIAS link under literature.							
	Oral examination: More details will be given at the beginning of the module.							