

## Scientific and Methods Module “Molecular Modelling”

<b>Module</b>	<b>HIGH THROUGHPUT SCREENING IN DRUG DISCOVERY</b>
<b>Aims</b>	Teach theoretical and practical aspects of drug discovery with a focus on computer-aided drug discovery and high throughput screening.
<b>Basics</b>	Biochemistry, amino acid structure, peptide bond, secondary structure, tertiary structure, small molecule binding, central dogma of molecular biology
<b>Contents</b>	The class will give an overview of drug discovery with a focus on computer-aided drug discovery and high throughput screening. Students will further learn the overall drug discovery process through the discussion of case studies.
<b>Methods</b>	High-throughput Screening in Drug Discovery, Experimental and Computational
<b>Type</b>	10 day block course, 05 – 16 December 2016
<b>Work load</b>	<b>MODULE: HIGH THROUGHPUT SCREENING IN DRUG DISCOVERY</b> 30 hours presence / 60 hours self-study Each 3-hour session includes a lecture presented by one of the two instructors. Some sessions include a 90min discussion of a case study prepared and presented by one student.
<b>Examination</b>	Written Exam
<b>Credit points</b>	1
<b>Responsible scientists</b>	Prof. Dr. David Weaver (Vanderbilt University) Prof. Dr. Jens Meiler (Vanderbilt University) Prof. Dr. Annette Beck-Sickinger (Leipzig University)
<b>Literature</b>	Scientific papers will be assigned in class
<b>Venue</b>	Universität Leipzig, Fakultät für Biowissenschaften, Pharmazie und Psychologie, Institut für Biochemie, Brüderstraße 34, 04107 Leipzig

## Instructors:

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## Schedule

HIGH THROUGHPUT SCREENING IN DRUG DISCOVERY						
2016	Dec 05	Dec 06	Dec 07	Dec 08	Dec 09	
LECTURES	09	<b>Topic</b> Course Overview and Introduction to the Drug Discovery Pipeline	<b>Topic</b> An Introduction to High-throughput Screening	<b>Topic</b> HTS Assay Technologies: Strengths, Limitations, and Future Directions	<b>Topic</b> Technology-enabled Hits-to-Leads and Lead Optimization	<b>Topic</b> Practical uses of HTS for academic and industrial probe discovery and development
	– 12	Jens Meiler	Charles David Weaver	Charles David Weaver	Charles David Weaver	Charles David Weaver
HIGH THROUGHPUT SCREENING IN DRUG DISCOVERY						
2016	Dec 12	Dec 13	Dec 14	Dec 15	Dec 16	
LECTURES	09	<b>Topic</b> Protein-Ligand-Docking	<b>Topic</b> Structure-based Virtual Screening & Drug Design	<b>Topic</b> Constitution, Configuration, & Conformation & Structure Generators	<b>Topic</b> QSAR, Ligand-based Virtual Screening, and Pharmacophore mapping	<b>Topic</b> Overflow, Conclusion, Exam
	– 12	Jens Meiler	Jens Meiler	Jens Meiler	Jens Meiler	Jens Meiler