

## Seminar

# Bayerischer Forschungsverbund Interaktion von humanen Gehirnzellen "ForInter"

### 14. - 15. 10. 2019

# Abtei Frauenwörth, Fraueninsel

### 83256 Frauenchiemsee

#### 14. 10. 2019

10.45 - 11.15 am	Welcome and Introduction
11.15 – 12.00 am	Short reports from the Task forces
12.00 - 1.00 pm	Lunch
1.00 - 3.00 pm	<b>Poster Blitz:</b> Short presentations of the projects (3 slides, 3 minutes/poster) followed by the <b>Postersession</b>
3.00 - 3.30 pm	Coffee Break
3.30 – 5.45 pm	<b>Bioethics Workshop</b> David Hamburger/Hannes Wolff, Universität Passau
	<ol> <li>Relevance of accompanying legal research (D. Hamburger)         <ul> <li>Why is accompanying legal research necessary?</li> <li>Why do legal questions arise? What legal questions need to be answered?</li> </ul> </li> </ol>
	<ol> <li>The constitutional basis. Exemplified on the freedom of research (H. Wolff)</li> <li>An introduction to the systematics of constitutional law.</li> </ol>
	<ol> <li>Giving a practical example: Interpreting an EU-directive (D. Hamburger)</li> <li>A presentation of the relevance of accompanying legal research in ForInter.</li> </ol>

	<ul> <li>Legal methods exemplified by an explanation of the interpretation of an EU-directive.</li> </ul>
	<ul> <li>4. The relationship of Ethics and the Law (H. Wolff) <ul> <li>Which ethical questions might arise?</li> <li>What relationship do Ethics and the Law have? What influence do ethics have on the Law and vice versa. What impact does this have on accompanying ethical and legal research.</li> </ul> </li> </ul>
6.00 pm	Dinner
	Get to together
15. 10. 2019	
9.00 – 10.30 am	Bioinformatics Workshop Maren Büttner, TU München
-	<ul> <li>Topics:</li> <li>General overview on single-cell RNAsequencing (experimental aspects, borrow from 10X genomics)</li> <li>Brief overview on experimental design questions</li> <li>Brief introduction on the costs of single-cell RNAsequencing</li> <li>Overview on computational models for single-cell RNAsequencing</li> <li>Introduction to the analysis pipeline</li> <li>Recent developments in the field</li> </ul>
	<ul> <li>Specific topics could be:</li> <li>How to deal with other type of omics (bulk RNAseq, ATACseq,)</li> <li>Ambient RNA, doublet</li> <li>frozen tissue and nucleus RNA-sequencing</li> <li>spatial transcriptomics</li> </ul>
10.30 – 11.00 am	Coffee Break
11.00 – 11.45 am	<b>Q&amp;A session:</b> Specific questions from the participants to the Bioinformatics (please send your questions, expectations in advance)
11.45 – 12.00 am	Closing remarks, summary and outlook
12.00 –  1.00 pm	Lunch