

**GRK 1478 PhD-lectures
2nd round 2009**

Name	Date	Title
Dempwolff, F.	18. September 09	Expression of bacterial cytoskeletal elements
Förster, K.	18. September 09	Investigation of Membrane Protein Complexes by ESI-Mass
Cymer, F.	16. Oktober 09	CANCELLED!
Gerbeth, C.	16. Oktober 09	CANCELLED!
Glessner, U.	16. Oktober 09	CANCELLED!
Barthel, S.	30. Oktober 09	Two type I signal peptidases in <i>Synechocystis</i> sp. PCC6803
Werner, M.	30. Oktober 09	CANCELLED!
Cymer, F.	30. Oktober 09	Detergent binding explains anomalous SDS-PAGE migration of membrane proteins
Kaufenstein, M.	13. November 09	<i>E. coli</i> transports aggregated proteins to the poles by a specific and energy-dependent process
Kuhn, P.	13. November 09	White spots on a map: Defining the molecular contacts of FtsY
Labatzke, R.	13. November 09	Isolation and Crystallization of NuoEF from <i>Aquifex aeolicus</i>
Pawlik, G.	20. November 09	
Probst, K.	20. November 09	Microbial Drug Efflux Proteins
Seidel, J.	20. November 09	Crystal structure of the ATP-gated P2X4 ion channel in the closed state
Simon, L.	27. November 09	Dynamics of DNA binding proteins in <i>Bacillus subtilis</i>
Stroud, D.	27. November 09	How to find a new project: High-density epigenetic interaction maps
Vögtle, N-F.	27. November 09	Identification and Characterization of Icp55, a novel mitochondrial processing peptidase
Erhardt, H.	4. Dezember 09	<i>In vivo</i> localization of the OXPHOS complexes in <i>E. coli</i>
Gerbeth, C.	4. Dezember 09	Identification of novel mitochondrial kinases and phosphatases
Glessner, U.	4. Dezember 09	Localizing the Q-binding site in <i>E. coli</i> Complex I via spin-coupling experiments
Wenk, M.	11. Dezember 09	When an ATPase is not an ATPase: at Low Temperatures the
Wenk, M.	11. Dezember 09	C-Terminal Domain of the ABC Transporter CvaB is a GTPase
Werner, M.	18. Dezember 09	CANCELLED!
Härle, J.	18. Dezember 09	Protein production by an mRNA interferase
Dörner, K.	18. Dezember 09	CANCELLED!
Werner, M.	5. February 2010	FTIR Difference Spectroscopy: a powerful tool for investigating primary processes in proteins