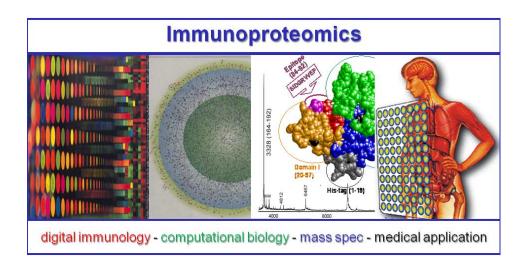
6th Proteome Forum Rostock - 2010



2nd to 4th of December, 2010

University of Rostock

Medical Faculty

BMFZ

Organizers:

M.O. Glocker, Proteome Center Rostock, University of Rostock H.-J. Thiesen, Institute of Immunology, University of Rostock

Thursday, 2nd of December 2010

16:00 – 20:00 Faculty Research Seminar and Excellence Research Project "Mecklenburg-Western Pommerania"

Chair: M.O. Glocker, Rostock

16:00 – 17:00	J. Lindquist, Magdeburg Systems Biology of T-cell Activation
17:00 – 17:30	HJ. Thiesen, Rostock, Coordinator Epitope Mapping of human IVIg (challenge 1 of DREAM 5)
17:30 – 18:00	M.O. Glocker, Rostock Epitope Mapping on Topoisomerase IIa with Sera from Colon Cancer Patients
18:00 – 18:30	M. Linnebacher, Rostock B-lymphocytes Infiltrating Colorectal Carcinoma show no Evidence of Malignant Transformation but Produce Tumor- Recognizing IgG
18:30 – 19:00	M. Lustrek and G. Füllen, Rostock Epitope Prediction and Analysis Based on Peptide Array Data
19:00 – 20:00	Y. Li, Shanghai The Essential Role of Protein Phosphorylation in the Evolution of Vertebrates
20:30 – 22:00	Speakers Dinner

Friday, 3rd of December 2010

10:00 - 13:15 MedSYS - meeting of the BMBF-consortium

Chair: H.-J.. Thiesen, Rostock

10:00 – 10:45	H. Juhl, Hamburg Challenges to use Signaling Pathway Activity as Predictors for Drug Response
10:45 – 11:15	HJ. Thiesen, Rostock Immune Responses to Tumors
11:15 – 11:45	M. Walzer, Tübingen High-throughput identification of tumor-associated HLA class I ligands
11:45 – 12:15	M. Schmitt, Rostock Peptide Vaccination of Leukemia Patients
12:15 – 12:45	F. Buelens, Göttingen Methods for the calculation of free energy differences in macromolecular complexes
12:45 – 13:15	G. Eschenburg, Berlin DNA vaccination in TRAMP mice
13:15 – 14:30	Lunch Break

Friday, 3rd of December 2010

14:30 – 18:30	Systems Pathology and Method Development / Deep Sequencing
	Chair: HJ. Thiesen, Rostock
	Systems Pathology and Method Development
14:30 – 15:00	HJ. Thiesen, Rostock Systems Pathology
15:00 – 15:30	J. Heise, Halle Refraction-2D: Novel Technology for High Performance Multipex 2D Analysis
15:30 – 16:00	B. Frerich, Rostock Trends in Diagnostic and Treatment of Oral Cancer and other Head and Neck Tumors
16:00 – 16:30	Break
16:30 – 17:00	E. Dülsner, Berlin Small Molecules Capture Compounds: Towards a Functional Reduction of Proteome Complexity
17:00 – 17:30	M. Maruschke, Rostock In Search of Prognostic Parameters for Novel Therapeutic Approaches in Renal Cancer
	Deep Sequencing
17:30 – 18:00	Y. Li, Shanghai Identification of Causative Variations based on the Technology of Next Generation Sequencing
18:00 – 18:30	HJ. Thiesen, D. Koczan, M. Kreutzer, Y. Li, G. Füllen Discussion on Computational Infrastructure for Deep Sequencing
19:30 – 21:00	Dinner

Saturday, 4th of December 2010

10:00 - 13:30 Structured Ph.D. curriculum - Participants' Camp

Chair: M.O. Glocker, Rostock

10:00 – 10:45	M.O. Glocker, Proteome Center Rostock Welcome and Introduction
10:45 – 11:15	NN / R. Jaster, Gastroenterolgy, Rostock Role of ROS in Pancreatitis and Pancreatic Ageing
11:15 – 11:45	T. Eickner / M.O. Glocker, Proteome Center Rostock Affinity Mass Spectrometry of Low Abundant Endogenous Proteins: How to differentiate a Target from its Background
11:45 – 12:15	M. Wölter / M.O. Glocker, Proteome Center Rostock Proteome Signatures by Mass Spectrometric Blood Profiling of Patients
12:15 – 12:45	M. Ludwig / J. Li, Heart Surgery, Rostock The c-kit+(AT2R+) Cell Population and their Potential in Cardiac Repair after Ischemic Injury
12:45 – 13:15	C. Röwer / M.O. Glocker, Proteome Center Rostock Quantitative Proteomics and Pathology - Merging Two Worlds
13:15 – 13:30	M.O. Glocker, Proteome Center Rostock Summary and Farewell