Monday, September 29th 2014

11.00 11.10	Walaama Claudia Stainam
11:00 – 11:10 11:10 – 11:55	Welcome – Claudia Steinem
11.10 - 11.55	Tobias Baumgart Curvature coupling of peripheral membrane proteins
11:55 – 12:20	Markus Müller
11.00 - 12.20	Coarse-grained simulation of changes of membrane topology
12:20 - 12:45	Michael Meinecke
12.20 12.10	Identification of a mitochondrial membrane morphology affecting
	protein necessary for cristae junction formation
12:45 – 13:45	Lunch
13:45 – 14:30	Bruno Antonny
	Atomic view of the response of phospholipids to membrane
	curvature and the advantage of polyunsaturated acyl chains for
	membrane fission
14:30 – 14:55	Claudia Steinem
	Pore-spanning membranes: Monitoring protein-mediated
14:55 – 15:20	membrane fusion and fission processes Tim Salditt
14.55 - 15.20	Membrane fusion in model bilayers stacks: in search for the
	magic mixture
15:20 – 15:45	Andreas Janshoff
10.20 10.10	Interaction forces in membrane fusion - how colloidal beads and
	coiled coil model peptides can help to quantify them
15:45 – 17:00	Coffee Break and Poster Session
17:00 – 17:45	Daniel Nietlispach
	Functional studies of 7TM receptors: insight from solution NMR
17.15 10.10	spectroscopy
17:45 – 18:10	Markus Zweckstetter
	Atomic views of the structure and dynamics of membrane
18:10 – 18:35	proteins: VDAC and TSPO Rasmus Linser
10.10 - 10.55	Characterization of protein function by proton-detected solid-state
	NMR spectroscopy
18:40 – 19:00	Cornelia Pfordt
	Guided tour of the Paulinerkirche
19:00	Conference Dinner – Location: Bullerjahn

Tuesday, September 30th 2014

09:00 – 09:45	Christoph Thiele Click-chemistry based lipid tracing to study lipid metabolism and localisation
09:45 - 10:10	Daniel B. Werz
09.45 - 10.10	Gb ₃ glycosphingolipids differing in their fatty acid chain: Chemical synthesis and biophysical behavior in solid-supported membranes
10:10 – 10:35	Igor Tkach Distances and orientations in peptides studied with 94 and 263 GHz EPR and PELDOR
10:35 – 11:15	Coffee Break
11:15 – 12:00	Eric Moulines Hidden Markov models and non linear filters: a state of the art and beyond
12:00 - 12:25	Axel Munk
	Statistical analysis of highly fluctuating signals: from electrophysiology to protein dynamics
12:25 - 12:50	Jörg Enderlein
	Diffusion in lipid bilayers
	Lunch
12:50 – 14:00	Lunch
12:50 – 14:00 14:00 – 14:45	Siewert Jan Marrink
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Wednesday, October 1th 2014

09:00 - 09:45	Margitta Dathe Antimicrobial peptides: The secrets of the Lord of the Rings
09:45 – 10:10	Jochen Hub Peptide-membrane interactions from potential of mean force calculations
10:15 – 10:45	Coffee Break
10:15 – 10:45 10:45 – 11:30	Coffee Break Sandro Keller An unusual membrane-protein topology for sensing bilayer thickness and triggering bacterial biofilm formation