# **Scientific Program**

## Thursday, May 3, 2012

19:30 Welcome drink, vila Lanna

### Friday, May 4, 2012

8:30 – 8:40 Invitation by Zdeněk Havlas, IOCB director

#### **Session 1. Proteins in Membranes**

Chairman Andreas Plueckthun

<b>1.1</b> 08:40 – 09:20	Daniel Mueller (ETH Zurich, CH) Reversible un- and refolding of single transmembrane beta-barrel proteins into membrane bilayers
<b>1.2</b> 09:20 – 10:00	Gunnar Von Heijne (Univeristy of Stockholm, SE) Energetics and dynamics of translocon-mediated membrane-protein insertion
<b>1.3</b> 10:00 – 10:40	Kvido Strisovsky (IOCB AS CR, Prague, CZ) Proteolysis in lipid membrane: substrate specificity and mechanism of rhomboids
10:40 - 11:00	Coffee Break
<b>1.4</b> 11:00 – 11:40	Anne Ulrich (Universitat Karlsruhe, DE) Charge zippers: a novel motif for folding and self-assembly of membrane proteins
<b>1.5</b> 11:40 – 12:20	Valerie Daggett (University of Washington, Seattle, USA) Proteins in different environments: the effect of osmolytes, lipids, and neighboring proteins on protein unfolding

12:30 - 14:00 Lunch

#### Session 2. Proteins in osmolytes, denaturants and salts

Chairman Valerie Daggett

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<b>2.1</b> 14:00 – 14:40	George Rose (Johns Hopkins University, Baltimore, USA) Protein domains – a thermodynamic definition
<b>2.2</b> 14:40 – 15:20	George Makhatadze I (Rensselaer Polytechnic Institute, USA)  Protein folding and aggregation: linking computation and experiment
<b>2.3</b> 15:20 – 16:00	Jim Warwicker (Manchester Interdisciplinary Biocentre, UK) Continuum electrostatics models for biology
16:00 - 16:20	Coffee Break
<b>2.4</b> 16:20 – 17:00	Vladimir Uversky (University of South Florida, Tampa, USA) Intrinsically disordered proteins in salted water and thick soup
<b>2.5</b> 17:00 – 17:40	Stefano Gianni (University of Roma, IT) Nearly identical, yet very different: addressing the folding of proteins with



increasing degree of sequence identities but different structure and function