



6. Ulm Meeting – Biophysics of Amyloid Formation Virtual meeting

22 February 2022

09:00 **Marcus Fändrich** | Ulm University, Germany
Welcome Address

Chair Margaret Sunde

09:00 **Astrid O. Gräslund** | Stockholm University, Sweden
The amyloid beta peptide: biophysical studies of interactions, structures and aggregation

09:30 **Ruiqing Ni** | University of Zurich, Switzerland
High precision in vivo multi-scale imaging of Alzheimer's β -amyloid deposits

09:50 **David Klenerman** | University of Cambridge, UK
Tau aggregation and spreading in neurodegenerative disease

10:20 **Amberley Stephens** | University of Cambridge, UK
Decreased water mobility increases alpha-synuclein protein misfolding

10:40 **Wolfgang Hoyer** | Heinrich Heine University Düsseldorf, Germany
Inhibitor-substrate cooperativity in blocking of α -synuclein fibril

11:00 **Break**

Chair Karin Kühnel

11:30 **Emma Sparr** | Lund University, Sweden
 α -synuclein interactions with lipid membranes-cooperative binding and membrane deformation

11:50 **Nunilo Cremades** | University of Zaragoza, Spain
Heterogeneous vs homogeneous nucleation in alpha-synuclein amyloid formation

12:10 **John Collinge** | University College London, UK
Understanding prion propagation and neurotoxicity

12:40 **Itzel Condado-Morales** | Institute of Neuropathology, University Hospital Zurich, Switzerland
Scaling analysis reveals the mechanism and rates of prion replication in vivo

13:00 **Break**

Chair Roland Riek

14:00 **Qiuye Li** | Case Western Reserve University, USA
Cryo-EM structure of disease-related prion fibrils provides insights into seeding barriers

14:20 **Yann Fichou** | IECB / University Bordeaux, France
Mechanisms of tau protein aggregation

14:40 **Cláudio M. Gomes** | University of Lisbon, Portugal
The chaperone activity of S100B prevents tau aggregation and seeding

15:10 **Susanne Aileen Funke** | Hochschule für angewandte Wissenschaften Coburg, Germany
Potent Tau aggregation inhibitor D-peptides selected using mirror image phage display

15:30 **Michel Goedert** / MRC Laboratory of Molecular Biology, UK
Cryo-EM structures of amyloid filaments from human brain

16:00 **Break**

Chair David S. Eisenberg

16:30 **Robert Tycko** / National Institutes of Health, USA
Structures and structural variations in amyloid-beta fibrils from brain tissue

17:00 **Lewis E. Kay** / University of Toronto, Canada
Seeing the invisible by solution NMR provides insight into protein aggregation

17:30 **Byron Caughey** / National Institutes of Health, Hamilton, USA
Prions at high resolution

18:00 **End of today's sessions**

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Chair Yuji Goto

09:00 **Daniel Otzen** / Aarhus University, Denmark
Contrasting mechanisms of aggregation of pathological and functional amyloid

09:30 **Frederic Rousseau** / Switch Laboratory KU Leuven, Belgium
Heterotypic amyloid interactions and their effect on amyloid assembly

10:00 **Christofer Lendel** / KTH Royal Institute of Technology, Sweden
Food protein amyloid: structure, fibrillation mechanisms and cross-seeding

10:30 **Ehud Gazit** / Tel Aviv University, Israel
Metabolite and protein amyloid formation: Interplay and its pathological implications

11:00 **Break**

Chair Per Hammarström

11:30 **Aphrodite Kapurniotu** / Technical University Munich, Germany
Designed peptides as cross-amyloid inhibitors of amyloid self-assembly of IAPP and Abeta42

12:00 **Lucie Khemtouri** / University of Bordeaux, France
Modulation of IAPP fibril formation by intrinsic and extrinsic factors

12:20 **Nicolas Guthertz** / University of Leeds, UK
The effect of mutation on an aggregation-prone protein

12:40 **Kichitaro Nakajima** / Osaka University, Japan
Mechanism of supersaturation-limited onset of dialysis-related amyloidosis by β 2-m

13:00 **Break**

Chair Stefano Ricagno

14:00 **Giampaolo Merlini** / University of Pavia, Italy
What makes light chains amyloidogenic and alinical relevance

- 14:30** **Lynn Rademaker** / Ulm University, Germany
Cryo-EM structures of amyloid fibrils from AL amyloidosis
- 14:50** **Andrea Cavalli** / Institute for Research in Biomedicine, Switzerland
Antibody somatic mutations predict immunoglobulin light chain toxicity
- 15:20** **Georg J. Rottenaicher** / Technical University Munich, Germany
Insights into the mechanism of amyloid formation in systemic light chain amyloidosis
- 15:40** **Manuel Hitzenberger** / Technical University Munich, Germany
MD as a tool for the investigation of the mutation-driven fibrillation of antibody light chains

16:00 **Break**

Chair Sheena E. Radford

- 16:30** **Luis del Pozo-Yauner** / University of South Alabama, USA
The Mechanism of Amyloid Fibril Formation in the Lambda-6 Immunoglobulin Light Chains
- 17:00** **Marina Ramirez-Alvarado** / Mayo Clinic, USA
Biophysics of Light Chains from Bacteria and Human Cells: Not all light chains are equal
- 17:30** **Peter E. Wright** / The Scripps Research Institute, USA
Kinetics and mechanism of transthyretin misfolding and aggregation
- 18:00** **Marcus Fändrich** | Ulm University, Germany
Concluding remarks, farewell

18:00 **End of the meeting**

Organization: Marcus Fändrich & Astrid Albiez