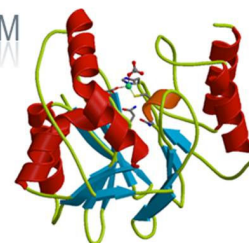




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The **BELGIAN BIOPHYSICAL SOCIETY**  
Further supported by the Graduate School  
“Structure and Function of Biological  
Macromolecules, Bioinformatics and Modelling”  
**ORGANIZES THE 14<sup>th</sup> ONE-DAY SYMPOSIUM ON**  
**PROTEIN FOLDING AND STABILITY**

**ON FRIDAY THE 2<sup>nd</sup> OF SEPTEMBER 2016**

**AT THE UNIVERSITY OF LIEGE**  
**AMPHITHEATRE 204, “Amphithéâtres de l’Europe”**  
**DOMAINE UNIVERSITAIRE DU SART TILMAN**  
**BATIMENT B4**

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09:45-10:00 WELCOME

**Chair : Carlo van Mierlo** (University of Wageningen, The Netherlands)

10:00-10:50 **John Christodoulou** (University College London, U.K.)

*“Protein Folding on the Ribosome”*

10:50-11:20 COFFEE BREAK

11:20-12:10 **Eric Lorent** (COMPLIX Alphabody Therapeutics, Belgium)

*“Alphabodies, a de novo designed protein with a uniquely stable fold”*

12:10-13:00 **Christian Johannessen** (Dept. Chem., University of Antwerp, Belgium)

*“Towards a standardized characterization of solution phase protein structure using Raman optical activity”*

13:00-14.00 LUNCH

14:00 –14:50 **Ruth Kellner** (Department of Biochemistry, University of Zurich, Switzerland)

*“Single-Molecule Fluorescence Spectroscopy of the Interaction of Proteins with the DnaK-DnaJ-Chaperone System”*

14:50-15:40 **Salvador Ventura** (Universitat Autònoma de Barcelona, Spain)

*“Repositioning tolcapone as a potent inhibitor of transthyretin amyloidogenesis”*

15:45 General assembly of the BBS and RECEPTION

**Contact:** Prof André Matagne ([amatagne@ulg.ac.be](mailto:amatagne@ulg.ac.be)). **Registration is required via [www.biophysics.be](http://www.biophysics.be)** . Admission is free for members; non-members of the Biophysical Society (<http://www.biophysics.be>) should pay their membership (10 EUR to be paid to the account of the Belgian Biophysical Society, IBAN BE73 0000 1463 3660, BIC BPOTBE1, P/A Yves Engelborghs, Celestijnenlaan 200 G-box 2403, B3001 Leuven).