



HEC 12 - PROGRAMME

Thursday, September 24

9:00 16:00 Exhibition installation
12:00 15:50 Registration

15:30 *Coffee break*

15:50 18:00 Thursday evening session 1

15:50 16:20
L1 h3
Phillipp Dautel
(*Halle-Wittenberg*)
Influence of RNA binding on the structure of Borna disease virus matrix protein

16:25 16:45
L2 h3
Tereza Skálová
(*Prague IMC*)
Two-domain laccase from *Streptomyces coelicolor*: a link between laccases and nitrite reduc

16:50 17:10
L3 h4
Tatiana Prudnikova
(*České Budějovice*)
Crystallisation and structural structure analysis of a novel haloalkane dehalogenase DbeA from *Bradyrhizobium elkanii* US

17:15 17:35
L4 h5
Eike Schulz
(*Göttingen*)
Crystal Structure of a Novel Intramolecular Chaperone Mediating Triple- α -Helix Folding

17:40 17:55
L5 h6
Marianna Biadene
(*Bruker*)
Ready, set, screen: X8 PROSPECTOR

18:00 19:00 Dinner

19:00 20:50 Thursday evening session 2

19:00 19:20
L6 h7
Harald Striegl
(*Berlin*)
Unusual armadillo fold in the human general vesicular transport factor p115

19:25 19:45
L7 h8
Madhumati Sevvana
(*Erlangen-Nürnberg*)

Structures of human and mouse apoM – serendipitous fatty acid in the binding pocket and refolding gone wrong!

18:50 20:50
L8 h8
Paul Emsley
(*Oxford, UK*)
Coot

21:00 22:00 HEC committee Meeting
21:00 Welcome party

Friday, September 25

8:00 9:00 Breakfast

9:00 10:40 Friday morning session 3

9:00 9:20
L9 h9
Daniela Dalm
(*Greifswald*)
Structural background of non-antibiotic affinity of minocycline to secretory phospholipase A2

9:25 9:45
L10 h9
Linda Schuldt
(*Hamburg*)
Stabilizing proteins for crystallization – how Thermofluor can help

9:50 10:10
L11 h10
Wojtek Bialek
(*Poznań*)
Atomic-resolution structure of reduced and oxidized cytochrome c6

10:15 10:35
L12 h11
Sven Dahms
(*Leibniz*)
Subdomains of the Amyloid Precursor Protein (APP) fall into Place: X-ray Structure of its N-terminal Heparin Binding Domain E1

10:40 *Coffee break*

11:00 12:10 Friday morning session 4

11:00 11:20
L13 h12
Thomas Monecke
(*Göttingen*)
Structural basis for CRM1 nuclear export complex assembly



Saturday, September 26		
11:25 11:45 L14 Pavel Mader (Prague IMG) Structural studies of human carbonic anhydrase isoenzymes	h13	
11:50 12:10 L15 Monika Sokolowska (Warsaw) The $\beta\beta$ -Me type II restriction endonuclease Hpy99I with target DNA	h13	
12:15 12:30 Stefan Kolek (Douglas Instruments)		h18
12:30 13:30 Lunch		
13:30 17:30 Free afternoon/Trip		h19
17:30 <i>Coffee break</i>		
18:00 20:010 Friday evening session 5		
18:00 18:15 L16 Mathias Grün (Jena Bioscience) Turning the bad into good: A rational approach to crystal quality improvement	h14	
18:20 18:40 L17 Karen Yates (Leipzig) Refolding of Ecto-Nucleotidases	h15	
18:45 19:05 L18 Elena Stsiapanava (České Budějovice) Structural changes of pathways in mutated DhaA proteins from Rhodococcus rhodochrous	h16	
19:10 19:30 L19 David Schwefel (Berlin, MDC) Structural insights into the GIMAP family	h17	
19:35 19:55 L20 Anja Lehweß-Litzmann (Göttingen) X-ray crystallographic snapshots of intermediates in transaldolase	h17	http://www.img.cas.cz/hec
20:00 Get-together		