











XXV IUCr Congress satellite workshop

Methods of Structural Biology



Joining CCP4/CCP-EM online workshop and CMS workshop

Dates: August 9 – 10, 2021 (CCP4/CCP-EM), August 12 – 13, 2021 (CMS)

Venue: Online + Institute of Biotechnology of the Czech Academy of Sciences, Biocev Centre, Prumyslova 595, Vestec near Prague

Aim: The CMS workshop will provide training and hands on experience in selected structural biology techniques available at the Centre of Molecular Structure (CMS) at IBT. Introductory talks will be accompanied by more advanced insights into biophysics, protein crystallization, X-ray diffraction, small angle X-ray scattering and structural mass spectrometry. Sessions at instruments will focus on training in selected techniques.

The CCP4/CCP-EM online workshop will focus on theory and practice in computational approaches in bio-crystallography and Cryo-EM single particle analysis.

Format: The CCP4/CCP-EM workshop will be organized online by the CCP4 and CCP-EM staff. The CMS workshop, organized by the CMS staff, will be on site. According to the epidemiological situation, we will confirm this format or announce changes. There is a one-day break – August 11 – for travel of those who want to participate in both workshops.

Participation fee: CCP4/CCP-EM – free.

CMS workshop - 150 EUR (meals and course materials; does not cover accommodation and dinner on 11 Aug)

To apply: Please fill out the preliminary registration form for both parts at: https://forms.biocev.org/index.php/891685?newtest=Y&lang=en You can register to one or both workshops. In the case of high interest, preference will be given to those registered to both workshops. You will receive confirmation (and payment details if applicable) before the end of June.

Application deadline: The deadline for preliminary registration is June 17, 2021. The payment deadline is July 31, 2021.

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cssb.structbio.org

www.ccp4.ac.uk

www.ccpem.ac.uk











Preliminary programme (Prague, CEST time):

CCP4/CCP-EM online workshop

Monday, August 9

09:00-09:45 CCP4 Cloud Setup and General Introduction

09:45-10:45 Data processing and reduction

10:45-11:00 Break

11:00-12:00 Experimental Phasing and Density Modification

12:00-13:00 Lunch break

13:00-13:45 Molecular Replacement

13:45-14:45 Model Building and Refinement

14:45-15:00 Break

15:00-15:30 Automated Model Building

15:30-16:00 Validation and Deposition

16:00-16:30 Break

16:30-17:30 Introduction to CryoEM (talk)

17:30-18:00 CCP-EM (practical setup)

Tuesday, August 10

09:00-10:30 Single particle reconstruction (demo + short practical)

10:30-10:45 Coffee Break

10:45-11:15 Reconstructed volumes - interpretation, sharpening, noise reduction (practical)

11:15-12:00 De novo model building (practical)

12:00-13:00 Lunch

13:00-14:00 Model refinement and validation (practical)

14:00-15:00 Wrap-up / free time

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CCP4/CCP-EM online workshop

Registration

No fee is required for the CCP4/CCP-EM workshop. A webpage with details and registration information will be available soon. For now, please register your interest through the preliminary registration page.

Technical information/organizational

Participants will require access to a Linux or MacOS machine with CCP4 and CCP-EM installed. Access to Zoom and Slack will also be required.

Contact information for the CCP4/CCP-EM part ccp4 -at- stfc.ac.uk











CMS workshop

Wednesday, August 11

Travel, get together dinner (paid individually)

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Thursday, August 12

8:30 Arrival. corree. Instructions, meeting othe	8:30	Arrival, coffee, instructions, meeting others
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8:50-9:00	Welcome

Biophysics, introduction (Tatsiana Charnavets) 9:00-9:30

9:30-10:00 Biophysics, Advanced methods (TBA)

10:00-10:30 Basics of protein crystallisation (Jiří Pavlíček)

10:30-11:00 Crystallization of membrane proteins using lipidic systems (Martin Caffrey)

11:00-11:30 Coffee break

11:30-12:00 Basics of X-ray diffraction in structural biology (Jan Stránský)

12:00-12:30 Protein crystallography at XFELs (Arwen Pearson)

12:30-13:00 Application talk – TBA

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13:00-14:00 Lunch (local canteen)

st. Vitus 2 Practicals at instruments

14:00-15:15 Biophysical methods (Tatsiana Charnavets)

15:15-15:30 short break w. coffee/tea

15:30-16:30 Protein crystallisation (Jiří Pavlíček)

16:30-16:45 short break w. coffee/tea

16:45-18:00 X-ray diffraction (Jan Stránský)

19:00 Dinner







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Friday, August 13

8:30-9:00	Introduction to Small angle X-ray scattering		
	and its application in-house (Jan Stránský)		

9:00-9:30	Advanced SAXS modeling techniques for biologic		
	macromolecules in solution (Dima Molodenskyi)		

9.30-10.30	Mass Spectrometry	(Julien Marcoux -	Petr Pomr	nach)
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10:30-11:00 Coffee break

11:00-12:00 Bioinformatics (TBA)

12:00-12:30 Application talk (TBA)

12:30-13:30 Lunch (local canteen)





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Practicals at instruments/computers

13:30-14:45 Laboratory Small angle X-ray scattering (Jan Stránský)

14:45-15:00 short break w. coffee/tea

15:00-16:15 Native Mass Spectrometry (Petr Pompach)

16:15-16:30 short break w. coffee/tea

16:30-17:45 Bioinformatics (TBC)

19:00 Dinner



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Confirmed speakers/tutors:

Martin Caffrey (Trinity College, Dublin); Julien Marcoux (CNRS, Toulouse); Dima Molodenskyi (EMBL, Hamburg); Arwen Pearson (Universität Hamburg); Tatsiana Charnavets (IBT); Jiří Pavlíček (IBT); Petr Pompach (IBT); Jan Stránský (IBT)

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