





<u>Updates from the Centre of Molecular Structure</u>

Hottest information: We are part of a new Horizon 2020 Infrastructure MOSBRI



The European Union's Horizon 2020 INFRAIA programme has awarded a 5 M€ infrastructure grant to MOSBRI (Molecular-Scale Biophysics Research Infrastructure), a consortium of 13 academic centres of excellence and 2 industrial partners from 11 different European countries, coordinated by Institut Pasteur (Paris, France). One of the partners is the Institute of Biotechnology of the Czech Academy of Sciences with its facilities: Biophysical techniques and Diffraction techniques.

The MOSBRI project will enable to create a **research infrastructure** combining the distinct instrumentation and expertise of the individual partner laboratories, thus allowing to tackle an unusually wide variety of life science research questions. It will also **disseminate its knowhow** through an ambitious programme of training **workshops and meetings**, particularly suited to early career researchers and others new to the field. You can find more information at https://www.mosbri.eu.

CF Protein Production facility

The Centre of Molecular Structure has started a pilot operation of a **new Protein Production facility**. This initiative follows the preliminary interest from potential users and will result in a platform covering three main steps to **produce purified proteins of interest**. We offer:

- 1) The **cloning services** comprising both traditional restriction enzymes system and restriction-free (seamless cloning) methodology.
- 2) The **expression of proteins** in bacterium *Escherichia coli* as our standard expression system. We are in the process of establishing the baculovirus FlexiBAC expression system.



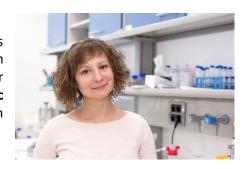


3) The **purification of proteins** using various types of chromatography. The most common are affinity (NiNTA, StrepTactin) and size exclusion chromatography.

Our current team consists of **Pavel Mikulecký** and **Tereza Nepokojová**. They are ready to help you with both routine large-scale protein production and demanding small-scale development of methods in semi high-throughput format.

CF Biophysical Techniques

At the start of the year the Biophysical Techniques facility welcomed its **new member Olga Dzmitruk**. With her MSc degree in physics and PhD in biophysics Olga is an expert in characterization of biomolecules and their interactions by biophysical techniques. She has **10 years of scientific research experience**. Olga has joined the CMS team with the focus on providing user support amongst others with the **FTIR technique**.













CF Diffraction Techniques

Our single crystal diffractometer D8 Venture received a new batch of upgrades. The most significant one was an upgrade of the detector to **Photon III (Bruker AXS)**. The detector hosts a bigger detection area, which allows faster high-resolution data acquisition, and higher sensitivity. The second item is an **automatic goniometer head (Bruker AXS)** which helps with high precision centering of the samples. Older accessories of the diffractometer, such as the in-situ stage and crystal dehumidifier, have registered increased popularity with our users and are bringing some interesting results.

CF Structural Mass Spectrometry



A new mass spectrometer timsTOF Pro from Bruker Daltonics was installed in the Structural mass spectrometry core facility. The mass spectrometer is mostly used for high throughput shotgun proteomics, hydrogen-deuterium exchange and covalent labelling experiments, and native mass spectrometry with ion mobility separation. The high sensitivity and sequencing speed allow identification of more peptides and their post-translational modifications.

A big congratulation is in order to **Pavla Vaňková** from the Structural Mass Spectrometry facility who successfully achieved her **PhD**! She has expertise in techniques of hydrogen/deuterium exchange MS and chemical protein crosslinking MS.



We are looking forward to you in CMS,

Tatsiana Charnavets (Biophysical Techniques), Olga Dzmitruk (Biophysical Techniques), Jiří Pavlíček (Crystallization of Proteins and Nucleic Acids), Petr Pompach (Structural Mass Spectrometry), Pavla Vaňková (Structural Mass Spectrometry), Jan Stránský (Diffraction Techniques), Michal Strnad (IT), Lubica Škultétyová (technician), Magdalena Schneiderová (admin), Jan Dohnálek (CMS Head)

You can realize your projects at CMS via an online application at <u>ciisb.org/open-access/proposal-submission</u>.

Find out more information at ciisb.org, ibt.cas.cz/core-facility/CMS/ or contact magdalena.schneiderova@ibt.cas.cz, jan.dohnalek@ibt.cas.cz



