

Monday, 2 November

- 10:00 - 10:10 Welcome note: Jiří Vondrášek - ELIXIR CZ Director
- 10:10 - 10:40 Lukáš Levák (Director of Department for Research and Development) - TBA
- 10:40 - 11:00 Jan Hrušák (ESFRI Chair) - TBA
- 11:00 - 11:30 Keynote 1: Corinne Martin (ELIXIR Hub) - Towards an assessment of ELIXIR's performance and impact
- 11:30 - 12:00 Keynote 2: Jerry Lanfear (ELIXIR Hub) - TBA
- 12:00 - 13:00 Lunchtime
- 13:00 - 14:15 Jiří Vondrášek (ELIXIR CZ Director)
ELIXIR CZ Strategy + Scientific Programme
- Structural bioinformatics
 - Chemical biology
 - Genomics
 - Bioinformatic data access and management
 - Precision medicine
- 14:15 - 15:00 Keynote 3: Bengt Persson (ELIXIR Sweden) - Human data - Federated EGA and the European 1+Million Genome Project
- 15:00 - 15:30 Coffee break
- 15:30 - 16:30 RIs fight the SARS-CoV-2/Covid-19 pandemic
- Jan Konvalinka (UOCHB)
 - Ludek Matyska (CEITEC MU)
 - Martin Modrak (MBU)
- 16:30 - 18:00 ELIXIR CZ international collaboration
ELIXIR Staff Exchange
- Miroslav Kratochvíl (UOCHB)
 - Pavel Ostašov (ZCU)
- Commissioned Services/Implementation study
- TBA (MU)
 - TBA (BTU)
 - Martin Hubalek (UOCHB)
- 18:00 - 19:30 Poster Session

Tuesday, 3 November

- 9:00 - 9:45 Keynote 4 : Martin Čech (Institute of Organic Chemistry and Biochemistry of the CAS) - Accessible, reproducible and transparent biomedical analyses with Galaxy
- 9:45 - 10:15 ELIXIR CZ Services/Platforms
- Karel Berka (PU) - Data Platform
 - Radka Svobodová (CEITEC MU) - Tools & Services Platform
 - TBA - Compute Platform
 - Jakub Galgonek (UOCHB) - Interoperability Platform
 - Vojtěch Spiwok (VSCHT) - Training Platform
- 10:15 - 10:45 Internal projects ELIXIR CZ
- Jan Pačes (IMG) - AmtDB (ancient mtDNA database) upgrade and bio.tools integration
 - Karel Berka (MU/PU) - Tools overview and validation platform for 3DBioinfo community
 - Radka Svobodová (MU/PU) - Visualization of Biomacromolecular Quality Criteria via LiteMol suite
- 10:45 - 11:00 Coffee break
- 11:00 - 12:30 Flash talks
- Bohdan Schneider (BTU) - Structural bioinformatics of nucleic acids at the Institute of Biotechnology of the Czech Academy of Sciences
 - Robert Pergl (FIT CVUT) - Data Stewardship Wizard helps fight COVID-19
 - David Hoksza (UK) - Structural bioinformatics tools at Charles University
 - Marek Schwarz (MBU) - rboAnalyzer webserver
 - Miloš Prokýšek (JCU) - UniCatDB
- 12:30 - 13:30 Lunchtime
- 13:30 - 14:30 Significant achievements of ELIXIR CZ
- David Bednar (FNUSA – ICRC) - EnzymeMiner: In silico mining of novel enzymes in sequence databases
 - David Sehnal (CEITEC MU) - Mol*: high performance macromolecular data delivery and visualization toolkit for the web
 - Petr Baldrian (MBU) - GlobalFungi, a global atlas of fungi derived from high-throughput-sequencing metabarcoding studies and a live database of fungal records
- 14:30 Farewell