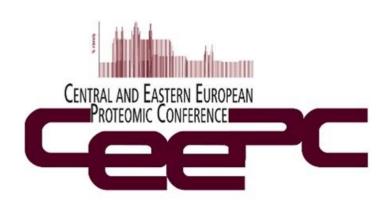
Scientific program



The 11th Central and Eastern European Proteomic Conference

27th-29th September 2017 in Košice, Slovakia





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Laboratory of biomedical microbiology and immunology

Department of microbiology and immunology

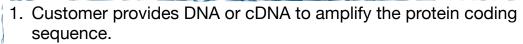








- Custom production of recombinant proteins in E.coli, Yeast (eukaryotic) and Leishmania (Eukaryotic) expression systems
- Pre-made recombinant proteins/antigens
- Custom protein array design

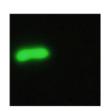


- 2. We give full FREE assistance in case codon optimization is necessary.
- We offer different tags as an option: (single 6XHis, Myc; dual tags – His and GFP, His and mCherry, His and Infra-red fluorescent protein, Myc and GFP).
- 4. Exclusion are applicable to highly toxic proteins and targets greater than 600 amino acids.
- 5. If the protein production fails because of high toxicity or high hydrophobicity (e.g. membrane proteins), any charge to the customer will be waived.
- 6. The timeline is approximately 4-6 weeks for *E. coli* expression system and 60-80 days for *Leishmania* expression system.
- 7. The endotoxin removal option is free (max purity should be <1 EU/ μ g of protein).
- 8. Protein must contain at least one tag. The customer should clearly state the preferred tag in order form.
- 9. Production of difficult proteins e.g. membrane proteins, toxic proteins.

Contact – lbmi@uvlf.sk or bhidemangesh@gmail.com Further information – http://lbmi.uvm.sk

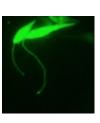
Custom protein price range - 300 - 600 Eur/protein/mg depending on the expression system used. Special price for order of more than 10 proteins.

Click to see price list of pre-made proteins - http://lbmi.uvm.sk/LBMI/ Premade_proteins_antigens.html.









Tuesday 26th September

17:00 - 19:30	Registration	
Wednesday 27th September		
8:00 - 9:00	Registration	
9:00 - 9:30	Opening ceremony	
Morning session		
Chair: Suresh Gadher		
09:30 - 10:10	Plenary - Chris W. Turck Pathway illumination for disease research - from Omics to Biosignatures	
10:10 - 10:35	Suresh Jivan Gadher The power of tears: tear proteogenomics using ProQuantum - High sensitivity immunoassay could revolutionize precision medicine in the clinic	
10:35 - 11:00	Lars Kristensen Next generation technology for reproducible and precise proteome Profiling	
11:00 - 11:30	Coffee break/ Poster viewing (poster presentation P#1 to P#23)	
Chair: Mario Cindrić		
11:30 - 12:10	Plenary - Alon Savidor Full-length de-novo protein sequence determination by Database Independent Protein Sequencing (DiPS)	
12:10 - 12:30	Jozef Hanes Highly sensitive immunological methods	
12:30 - 13:30	Lunch at Hilton Hotel / Posters and commercial booths viewing	

Afternoon session

Chair: Vladimir Havlíček

13:30 - 14:10 Plenary - Mario Cindrić

Automated sample preparation in proteomics, multidimensional chromatography, biotypization and beyond

14:10 - 14:35 Michal Bohac

Current challenges in proteomic research and latest solution using Bruker mass spectrometry

14:35 - 15:00 Jana Melničáková

Library preparation for NGS using NEBNext

15:00 - 15:30 Coffee break/ Poster viewing (poster presentation P#1 to P#23)

Chair: Mangesh Bhide

15:30 - 16:10 Plenary - Gustavo R. Makert

Vaccine development for the control of the ectoparasite Dermanyssus gallinae: a potential vector of pathogenic agents

16:10 - 16:30 Lucia Borszekova Pulzova

Design, construction and evaluation of multi-epitope chimeric proteins for diagnosis of Lyme borreliosis

17:00 Welcome dinner/Wine tasting, Departure to TOKAJ WINERY,

Meeting point – In front of Hotel Hilton. Travel to Tokaj Winery by bus.

Thursday 28th September

Morning session

Chair: Dasa Čižková

09:30 - 10:10 Plenary - Lee Wei Yang DynOmics: dynamics of structural proteome and beyond

10:10 - 10:30 Dasa Cizkova

Proteomic analyses along injured spinal cord axis: moving frontiers toward effective regeneration with smart scaffolds

10:30 - 10:50 Adriana Natalia Murgoci

Proteomic and morphological features of rat microglia derived exosomes

10:50 - 11:30 Coffee break/ Posters viewing (poster presentation P#24 to P#55)

Chair: Gabriel Mazzucchelli

11:30 - 12:10 Plenary - Vladimir Havlíček

Meet interesting abbreviations in clinical mass spectrometry: from compound classification by REIMS to multimodal and mass spectrometry imaging (MSI)

12:10 - 12:30 Dominika Luptáková

Membrane depolarization and aberrant lipid distributions in neonatal rat brain upon hypoxic-ischemic insult

12:30 - 12:50 László Drahos

Proteomic and phosphoproteomic analysis of the synaptosome in sleep deprivation

12:50 - 13:40 Lunch at Hilton Hotel/ Posters and commercial booth viewing

Afternoon session

Chair: Chris Turck

13:40 - 14:20 Plenary - Katrin Marcus

Tissue-based clinical proteomics in neurological/neuromuscular diseases

14:20 - 14:40 Santosh Jadhav

Tau protein as a perpetrator of synaptic impairment

14:40 - 15:05 Moustafa Khalf

Bio-Plex Technology (Overview and Applications) Stain Free, V3 Workflow & Fluorescence WB (toward a validated protein normalization)

15:05 - 15:40 Coffee break/ Posters viewing (poster presentation P#24 to P#55)

Chair: László Drahos

15:40 - 16:20 Plenary - Gabriel Mazzucchelli

MELD proteolysis: how to take advantage from a limited and multi-enzymatic digestion

16:20 - 16:40 Miroslava Némethová

Proteomic analysis of hippocampal CA1 region after ischemic tolerance induction in the rat brain

16:40 - 17:00 Laszlo Prokai

Neuroproteomics focusing on drug abuse in animal models

17:00 - 17:20 Katarína Bíliková

Potential application of honey bee proteins and antimicrobial peptides in human and veterinary medicine

Friday 29th September

Chair: Katrin Marcus

09:30 - 10:10	Plenary - David Eckersall Secretory proteomes and their roles in host defense against infection	
10:10 - 10:35	Dietrich Merkel 100 Proteomes A Week – The next step for industrialized quantitative proteomic	
10:35 - 10:55	Sabrina Bousbata Complement subversion by <i>Rhodnius prolixus</i> and its impact on <i>Trypanosoma cruzi</i> establishment on the insect's gut	
10:55 - 11:30	Coffee break	
Chair: Suresh Gadher		
11:30 - 11:50	Nicolas Guillemin Bioinformatics investigation of dog babesiosis from host differentially expressed proteins identified by proteomics	
11:50 - 12:10	Deepti Patel Molecular and structural differences in lumpsucker skin among various body sites	
12:10 - 12:30	Tomas Smolek Spreading of tau pathology in transgenic model of tauopathy	
12:30 - 12:50	Eva Klement Beyond the cell wall - extracellular phosphorylation	
12:50 - 13:20	Closing Ceremony	
13:20 - 14:00	After conference networking with coffee	

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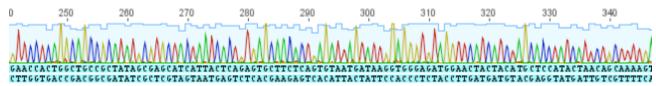
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CUSTOM SERVICES





DNA sequencing and genotyping



- → Minimal volume of sample: 10 ul
- → Quality of sample: we recommend to purify the PCR product from agarose gel or purification directly from PCR product, if there is any non-specific products in your PCR product. We are recommend to use Qiagen, Promega, Jena or similar.
- → Volume of primer: 10 ul of each primer with the concentration 10 pmol.
- → Maximal size of product: we can provide sequencing of max. 500 bp of each forward and reverse primer
- → Delivery: 2-7 days
- → Output format: sequence and chromatogram
- → Price: 9 Euro per sample (price is including sequencing with 1 primer and tax)

After sequencing, we are able to analyze your sequence and determine the gene, site of mutation, identify the pathogen and isolate form the sample. The price for genotyping is on request.

Contact - lbmi@uvlf.sk or bhidemangesh@gmail.com

Further information – http://lbmi.uvm.sk