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JARMILA STANKOVÁ

ABOUT ME

I am a research assistant and Ph.D. candidate at the Institute of Molecular and Translational Medicine in Olomouc, Czech Republic. Since 2016 I have been participating in the research program Chemical Biology and Experimental Therapeutics, mainly focusing on new methods for molecular target identification, such as microscopy and proteomics methods. I joined the multi-omics group in 2020 and currently working on the proteomics part of the EATRIS-Plus project. I am a former rugby 7s player and a big sports lover.

PERSONALITY TRAITS

Honesty - Dedication - Empathy - Creativity - Fun

WORK EXPERIENCE

RESEARCH ASSISTANT | INSTITUTE OF MOLECULAR AND TRANSLATIONAL MEDICINE, PALACKY UNIVERSITY 2016-NOW

<u>Cell analysis</u>: fluorescent confocal microscopy, live cells imaging, flow cytometry

Proteomics analysis: HPLC-MS, proteomic profiling, SILAC analysis, thermophoresis

Data analysis: Columbus, ImageJ, Proteome Discoverer, MaxQuant, Skyline, Spectronaut, Perseus

Robotic platform operation: HighRes Bio with a wide range of instrumentation

<u>Lab manager</u>: SOP preparation, public tender preparation, instrumentation service handling (RM and PM) <u>Teaching</u>: bachelor and diploma students

Results presentation skills and experience with working in an international team

Organization of "Proteomic Group Meetings"

MEDICAL LABORATORY PROFESSIONAL | OLOMOUC UNIVERSITY HOSPITAL |2020-2022

Covid-19 diagnostics, PCR testing, BSL 2-3 safety workflow (handling patient samples in BSL 2-3 safety condition)

EDUCATION

MASTER OF SCIENCE | FACULTY OF SCIENCE, PALACKY UNIVERSITY OLOMOUC, CZECH REPUBLIC | 2016 Experimental biologist – Medical and chemical biology / analytical chemistry – Thesis: Proteomic profile of CCRF-CEM cell line treated by 5-fluorouracil.





TRAINING AND PROFESSIONAL PRACTISE

ADVANCED PRACTICAL PROTEOMICS | EUROPEAN PROTEOMICS ASSOCIATION | VIENNA, AUSTRIA | 2018

Responsible teacher – Karl Mechtler, Subject of training – TMT Quantification, Cross-linking (XL-MS), Targeted proteomics by PRM, Proteome Bioinformatics

CHEMICAL PROTEOMICS | SCIENCE FOR LIFE LABORATORY, KAROLINSKA INSTITUTET | STOCKHOLM, SWEDEN | 2017

Responsible teacher – Massimiliano Gaetani and Roman Zubarev, Subject of training – Thermal proteom profiling (TPP), Functional identification of target by expression Proteomics (FITExP), Elucidation of the interaction interface and mapping of the binding site of a drug with its target protein by Hydrogen/Deuterium (H/D) exchange mass spectrometry (HDX MS)

MULTIPLEX PROTEINS ANALYSIS | PALACKY UNIVERSITY | OLOMOUC, CZECH REPUBLIC | 2013

Responsible teacher - Milan Raška, Subject of training - multiplex ELISA, imuno-PCR, Luminex

MEDICAL LABORATORY PROFESSIONAL | INTELMED S.R.O., OLOMOUC | JULY 2014

Laboratory work - FISH, immunohistochemistry, Comparative Genomic Hybridization - CGH, real-time PCR

MEDICAL LABORATORY PROFESSIONAL | CLINICAL BIOCHEMISTRY AND HEMATOLOGY S.R.O. | JULY 2013

Laboratory work - blood analysis and clinical testing for biochemistry and hematology

GRADUATED STUDENTS

ELIŠKA KŘESŤANOVÁ | DEFENSE IS PLANNED FOR 2024

Diploma thesis: Development of DIA method useful for plasma proteomic profiling of large sample cohort

ELIŠKA HLADÍKOVÁ | 2020

Bachelor thesis: Reporter lentiviral systems for subcellular localization

KATEŘINA JEČMEŇOVÁ | 2018

Bachelor thesis: Spectroscopic characteristics of used drugs and their use in the identification of molecular targets

AWARDS

Dean's award for the best student research paper (2021)





PUBLICATION LIST – IMPACT FACTOR JOURNALS ONLY

(7) D. BARUCIC*, S. KAUSHIK, J. KYBIC, **J. STANKOVÁ**, P. DŽUBÁK, M. HAJDÚCH, Characterization of drug effects on cell cultures from phase-contrast microscopy images, Computers in Biology and Medicine, 2022, 151, 106171, 0010-4825, **IF: 6.698**, PMID: 36306582.

(6) D. KODR*, **J. STANKOVÁ***, M. RUMLOVA, P. DŽUBÁK, J. ŘEHULKA, T. ZIMMERMANN, I. KRIZOVA, S. GURSKÁ, M. HAJDÚCH, P. DRAŠAR, M. JURÁŠEK, Betulinic Acid Decorated with Polar Groups and Blue Emitting BODIPY Dye: Synthesis, Cytotoxicity, Cell-Cycle Analysis and Anti-HIV Profiling, Biomedicines, 2021, 9, 1104, 2227-9059, **IF: 6.081**, PMID: 34572290.

(5) M. PORUBSKÝ*, K. VYCHODILOVÁ, D. MILICEVIC, M. BUDESINKY, **J. STANKOVÁ**, P. DŽUBÁK, M. HAJDÚCH, J. HLAVÁČ, Cytotoxicity of Amino-BODIPY Modulated via Conjugation with 2-Phenyl-3-Hydroxy-4(1H)-Quinolinones, ChemistryOpen, 2021, 10, 1104-1110, 2191-1363, **IF: 2.911**, PMID: 34427046.

(4) M. PORUBSKÝ*, S. GURSKÁ, **J. STANKOVÁ**, M. HAJDÚCH, P. DŽUBÁK, J. HLAVÁČ, AminoBODIPY Conjugates for Targeted Drug Delivery Systems and Real-Time Monitoring of Drug Release, Molecular Pharmaceutics, 2021, 18, 2385-2396, 1543-8384, **IF: 3.500**, PMID: 33961440.

(3) M. PORUBSKÝ*, S. GURSKÁ, **J. STANKOVÁ**, M. HAJDÚCH, P. DŽUBÁK, J. HLAVÁČ, Amino-BODIPY as the ratiometric fluorescent sensor for monitoring drug release or "power supply" selector for molecular electronics, RSC Advances, 2019, 9, 25075-25083, 2046-2069, **IF: 3.119**.

(2) S. KRAJČOVIČOVÁ*, **J. STANKOVÁ**, P. DŽUBÁK, M. HAJDÚCH, M. SOURAL, M. URBAN, A Synthetic Approach for the Rapid Preparation of BODIPY Conjugates and their use in Imaging of Cellular Drug Uptake and Distribution, Chemistry- A European Journal, 2018, 24, 4957-4966, 0947-6539, **IF: 5.317**, PMID: 29411907.

(1) T. OŽDIAN*, D. HOLUB, Z. MACEČKOVÁ, L. VARANASI, G. RYLOVÁ, J. ŘEHULKA, J. VÁCLAVKOVÁ, H. SLAVÍK, P. MOUDRÝ, P. ZNOJEK, **J. STANKOVÁ**, J. DE SANCTIS, M. HAJDÚCH, P. DŽUBÁK, Proteomic profiling reveals DNA damage, nucleolar and ribosomal stress are the main responses to oxaliplatin treatment in cancer cells, Journal of Proteomics, 2017, 162, 73-85, 1874-3919, **IF: 3.867**, PMID: 28478306.