WIESŁAW ABRAMOWICZ, M.S.

Laboratory Diagnostician, Oncological Computational Scientist (personalised diagnostic, nutri-, epi-, transcriptomics, epi-, genomics, proteomics, metabolomics)

Achivements:

- * Improving the functionality of the clinical laboratory IT system;
- * Cost effective sellecting personalized transcriptomics targets by microarray for on- and off-label treatment of tumors like in WINTHER clinical trial (NCT01856296):
- *The only person in Poland who performs the test properly on BD™ Bruker MALDI Biotyper;



EMPLOYMENT

Currently 05/2025

Implementation Specialist

MedLAN

Palaystok, PL

- · IT technical support provided to
- · implementation of IT applications;
- · creating and testing database software for web applications;
- · implementation of the needs of contractors from the area of pathology, laboratory clinical genetics in Poland;
- · use for HL7, P1 protocols, medical laws, implementation of recommendations of scientific societies (e.g. PTPat, PTGC);
- · support for implementation and development IT departments in the field of laboratory medicine.

12/2024 10/2024

Laboratory Diagnostician

Clinical genetic laboratory

Q Łódź. PL

- · improving the functionality of the laboratory;
- · quality control of each stage of laboratory work in accordance with ISO15189;
- · EMON:
- · GLP knowledge;
- · capacity calculation of flow cells; supervision of the execution of the library and patient sample preparation;
- · operation of five devices for NIPT (liquid biopsy - cfDNA), WES, WGS (DNBSEQ-G400 Sequencer, DNBSEQ-T7 Sequencer; SBS -Salus Nimbo, Pro, EVO)[up to 1.8 Gbp. 5.8 Gbp and 3 Gbpl:
- · cooperation with Professors;
- · superb contact with the vendors (Salus, MGI, Altium Group-service and genetic representative);
- · assistance in solving clients needs.

04/2024 11/2023

Laboratory Diagnostician – volunteering

Białystok Oncology Center – Laboratory Diagnostics Facility

Białystok, PL

- · InfoMedica;
- · hematology (Sysmex XN-1000, XN-550, ACL350);
- · biochemistry (Cobas 6000, e411, u411, u701, Integra 400 Plus, ABL90
- · microbiology (Vitek 2 Compact, BD™ Bruker MALDI Biotyper - the
- only person in Poland who performs the test properly, according to the manufacturer's
- · blood components distribution;
- · improving the functionality of the clinical laboratory IT system.

CONTACT INFO

✓ immunol22@gmail.com

+48 798 298 280

CAbout Me

MARZEM Wood & Stone -

- UI/UX/ Graphic Designer, Carpenter.

SKILLS

A Linux, R BStudio.

PGit. GitHub

Mlacktriangledown, $L\!\!\!/T_E\!X,$

HPC (openMPI, Rmpi). HTML5, CSS, JS, YAML, Google Analytics, Inkscape, GIMP, Blender.

GAMP5, GxP, CAPA, DoE, Agile oriented.

EXPERIENCE

Biostatistics (glm + Bayes)

Immunology, Oncology (4 years)

NGS, Agilent's Microarray, qPCR [wet lab, in silico] {kidney oncology}

Laboratory Diagnostics (Sysmex, Cobas 6000, e411, Integra, LIS improvements)

LANGUAGES

English: Proficient (C1)

Polish: Native

12/2022 4/2021

Owner, Co-founder & CEO

WARZEM Wood & Stone, Solid Wood Artistic Carpentry Workshop 🕙



- fundraising;
- · process optimization (Agile);
- · management, accountancy;
- · website design and build with HTML5, CSS, JS, YAML, Github, RStudio;
- · Google Analytics;
- · photography and video making;
- · graphic design and creation (Inkscape vector graphics);
- · video design and creation (Blender);
- · creating solid wood furnitures (using the golden ratio);
- · carpentry tools design and create;
- · using natural oils (no siccative, no PFAS) toxicological issues;
- · chopping down trees;

09/2009 09/2008

Pharmaceutical Assistant

Pharmacy "Lobelia" Barbara Winsko.

Białystok, PL

- · advising for patients in OTC medication;
- · issuing medicines prescribed by a physician;
- · accepting and placing orders;
- · delivering medicines to other points of sale.



PROFESSIONAL MEMBERSHIP

Currently 01/2014

The Right to Practice the Profession in Laboratory Diagnostic. No 13786 **KIDL**

Warsaw



RESEARCH EXPERIENCE

8/2017 10/2013 Transcriptomic profiling of a human kidney cancer cell line, Caki-2, after treatment of Ixazomib (onlabel), Ursolic acid and jointly using microarray technology. (pre-clinical)

Department of Pharmaceutical Biochemistry, Medical University of Białystok.

- · Budget management (successful negotiations with vendors within budget);
- · Design of Experiments (DoE);
- · Project manager and executive (wet lab, in silico analyses);
- · Performed wet laboratory experiments (Agilent's gene expression and miRNA microarrays of indirect transcriptome form total RNA (mRNA, IncRNA, lincRNA, sno/snRNA, ncRNA, miscRNA, miRNA), quality controls (pos/neg/spike-in), MTT drug testing (proteasome inhibitor, terpen), total RNA/DNA isolation, purification, quality, quantity (Bioanalyzer, NanoDrop), gene validation (qPCR, primer design, analysis))
- · and in silico analyses using open-source programming/biostatistical R/RStudio with packages from cran, bioconductor, github repos (pre-processing, numerical validation awarness, glm, Bayesian, multivariate, pathway enrichment analysis).

On time delivering wet lab experiments (I take care of the details and I am very precise, successful troubleshooting), performing and precaution of misleading computational analysis.

12/2015 10/2014 DNA sequencing of clear cell renal cell carcinoma and healthy kidney area. (clinical case)

"Genome-wide methods in cancer genetics". BASTION.

Department of Pharmaceutical Biochemistry, Medical University of Białystok.

- · project co-executive (wet lab, in silico analyses);
- · DNA sequencing (1000 genes) of human cancer and normal kidney using KAPA library. [with co-operation of Prof. Rafał Płoski, MD, PhD (Medical University of Warsaw).



Biostatistic experience

Interests: pre-processing (raw signal, transformations), manifolds/topology, glm, gee, mixed models

- exploratory and confirmatory analyses;
- · GLMs & Bayesian, mixed (limma, mmrm, lme4);
- · R, RStudio;
- · R packages (GO.db, pathview, RSubread, Gviz, biomaRt, plotly, ggplot, D3.js, igvR, etc.);
- · R programming (visualizations, office documentation, pdf, rtf);
- · data mining, regular expression, rest api;
- · pre-processing (log, lowess, loess, quantile) of transcriptomics data;
- · Reproducible tools (git, github, gitlab) and reporting Autodock/Vina); (rmarkdown, bookdown, etc.);
- · Pathway Enrichment Analysis (GO, KEGG, Reactome, CatBoost, LightGBM, XGBoost, H2O); WikiPathways, HumanCyc, clusterProfiler, rSEA);
- Functional Annotation;
- · GATK, STAR, DESeq2;
- · Multivariate Analysis (PCA, tSNE, UMAP, SONG, NMF, · more information in each description of course;

- · Comparative Analysis (Venn diagram);
- · Isobolograms:
- \cdot use of data sets (ncbi, google patents, etc.);
- · use of FDA resources (CDER, CDRH, NCTR mitochondial toxicity, MAQC/SEQC);
- · Gene Expression Omnibus;
- · numerical validation awarness;
- · Linux command line;
- · basics of MySQL;
- · basics of Python (PyScripter, PyMol, PyLasso,
- · basics of machine Learning (Random forest,
- · basics of neural Networks (TensorFlow, Keras, etc.);
- · Agilent's GeneSpring GX;
- · CLC Main Workbench;

HPC experience

High-Performance and Parallel Computing

· I built cluster of two personal computers with openMPI and Rmpi (an interface for R & RStudio) on Ubuntu (SSH connected).



WET LAB



Laboratory Experience

Research interests: Oncology, Immunology, EpiOmics, Transcriptomic, Proteasomes, Nutritranscriptomics, Watson-Crick/Hoogsteen base paring.

Medical University of Białystok, PL

- · BIO-RAD CFX Connect Real-Time PCR (qPCR);
- · Agilent's microarray (two-color gene expression, miRNA) + QC (positive, negative, spike-in controls);
- · NanoDrop 2000 (spectrophotometry);
- MTT tests (using proteasome inhibitor {Ixazomib} on-label FDA approved in multiple myeloma, terpen {Ursolic acid} - derivative, from e.g. apples, with antiinflammation, anti-cancer properties);
- · DNA, RNA extraction from cultured cells (Qiagen, Promega, A&A Biotechnology); RNase, DNase free environment:
- · Agilent 2100 Bioanalyzer (DNA/RNA microcapillary electrophoresis, flow cytometry) - lab-on-chip;
- · DNA, RNA gel electrophoresis;
- · Western blot with SNAP i.d.;
- PMN, PBMC cells isolation, neutrophils isolation with anti-CD16 mAb MicroBeads and magnetic separator Midi MACS;
- · cells counting chambers;
- setting up cell cultures;
- · optical microscope;
- · flow cytometry.



Currently | 01/2016

Data Science Specialization

coursera.org - online

♀ Johns Hopkins University, USA

"Exploratory Data Analysis" (11/2018)

- · lattice.
- · ggplot2.

- Clustering.
- · Dimension Reduction.

"Getting and Cleaning Data" (02/2016)

- · dplyr, httr, rhdf5, xml, RMySQL, foreign, rmongodb, RMongo, RPostresSQL;
- Regular Expressions (REGEX), R programming Data Cleaning, Data Manipulation.

"R programming" (01/2016)

· Data Analysis, R programming, Debugging, RStudio.

"The Data Scientist's Toolbox" (01/2016)

· Version Control (Git, GitHub).

· R Markdown.

08/2017 | 10/2013

Ph.D. Studies in Pharmaceutical Sciences

The Center for Innovative Research 2012-2017, Faculty of Medicine with the Division of Dentistry and Division of Medical Education in English, Faculty of Pharmacy and Division of Laboratory Medicine, Medical University of Białystok

₽PL

Research area: Transcriptomic profiling of cancer cell line after drugs treatment using Agilent microarrays and biostatistic open-source tools.

Advisor: Prof. dr hab. Marzanna Cechowska-Pasko, Ph.D.

All subjects passed (GPA 3,85):

- Principles of genetic analysis (6h);
- · Learning techniques of molecular biology (10h);
- Introduction to genomics (6h);
- · Epigenomics and transcriptomics (50h);
- Learning methods in structural and functional genomics (10h);
- · Basic Statistical refreshment (30h; prof. Tomasz Burzykowski, UHasselt);
- Statistical modeling (30h; prof. Tomasz Burzykowski, UHasselt);
- Design of experiments (18h; prof. Tomasz Burzykowski, UHasselt);

- · Statistics for omics (36h; prof. Ziv Shkedy, UHasselt);
- Metabolomics (16h);
- Facultative advanced courses in protein analysis/ proteomics/ metabolomics (50h);
- · Protein analysis and proteomics (32h);
- · Immunology (16h);
- · Facultative advanced courses in immunology (15h);
- Teaching the presentation and evaluation of research at the academic level (30h);
- · Professional practice (30h);
- · OHS (4h).



M.Sc. in Medical Analitics, POF 7

Faculty of Pharmacy and Division of Laboratory Medicine, Medical University of Białystok

PPL

Master's thesis: Expression of the APRIL particle in neutrophils of patients with potentially malignant lesions of the oral mucosa.

- · GPA: 4,30 (transcript of master degree);
- · isolation and work with human blood cells;
- · Advisor: Prof. dr hab. Ewa Jabłońska, Ph.D.
- · Co-Advisor: dr Kamil Grubczak, Ph.D.
- experience in biochemistry and immunology research (see Wet lab, Student membership & Publications sections in this resume);
- Dean's leave (10/2008 09/2009)

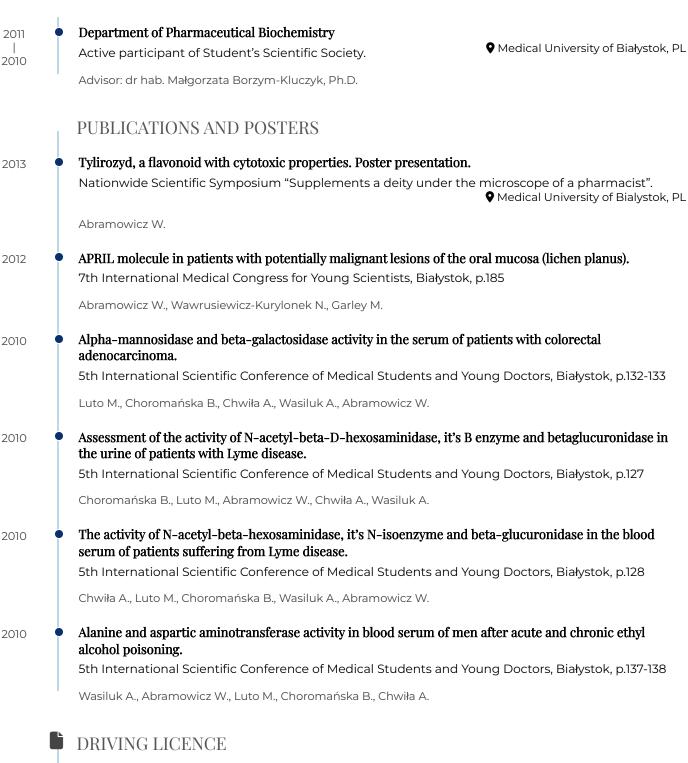


Pharmaceutical Assistant, PQF 4

Vocational School No. 1 of Health Care

🗣 Białystok, PL

-	DATA SCIENCE, LABORATORY COURSES	
29/ 31/03/2023	Duke Industry Statistics Symposium 2023 "Empower Clinical Development by Harnessing Data from Diverse Sources"	♥ Duke University, USA
25/01/2023	"Good Things Come in Small Packages: Purifying miRNA from Plasma, Serum and Exosomes" Michelle Mandrekar − Promega Online, PL	
	· Maxwell RSC miRNA	
22/ 25/09/2014	Courses: A&A Biotechnology	♥ Gdańsk, PL
	• "Real-Time PCR" • "Real-Time PCR - quantitative markings"	
10/2014	Prof. Tomasz Motyl Laboratory Experience: Department of Physiological Sciences, WULS (SGGW) Agilent's expression microarray experiments (wet lab only).	♥ Warsaw, PL
06/2016	"NGS in Regulatory Gene Research" ideas4biology	♀ Poznań, PL
	Coordinator: dr Michał Szcześniak, Ph.D. Techniques: ChIP-chip, ChIP-Seq, ChIP-exo, CLIP-Seq, DNase-Seq, FAIRE-Seq, GRO-Seq, HiC, MeDIP-Seq, MBD-Seq, MNase-Seq, oxBS-Seq, RIP-Seq, RRBS-Seq, TAB-Seq, TSS-Seq, WGBS-Seq. Linux. WinSCP, PuTTY.	
	Softwares: POLYPHEMUS, ChromHMM, Segway, MACE, DANPOS2, MeDUSA, diffReps, BSMAP, methylKit, Trimmomatic, Subread, BamTools.	
	Databases: OMICtools, AllSeq, ChIPBase, hmChIP, mirPath, miRTarBase.	
16/	"Using next-generation sequencing to analyse human transcriptome"	
l 18/06/2014	Richard Dixon, Ph.D. – QIAGEN	al University of Bialystok, PL
08/ 11/04/2014	"Using next generation sequencing to analyse human genome" Richard Dixon, Ph.D. – QIAGEN	al University of Bialystok, PL
07/04/2014	"Early diagnosis and treatment of neurodegenerative diseases" Department of Biochemical Diagnostics, Department of Neurology, Department od Neurodegeneration Diagnostics, Department of Pediatrics, Gastroenterology and Pediatric Allergology of Medical University of Bialystok Bialystok, PL	
1	STUDENT MEMBERSHIP	
2013	Member of the board of the Students' Scientific Society	
 2012	Co-organizer of the 8th & 7th Białystok International Medical Congress for Young Scientists.	
	Coordinator of Belarusian participants. • Medic	al University of Białystok, PL
06/2013	Department of Immunology	
10/2009	Active participant of Student's Scientific Society. Advisor: Prof. Ewa Jabłońska, Ph.D.	al University of Białystok, PL



AM/B₁/B

Currently 30/08/2006

Smooth driving, dynamic driving, no collisions.