WIESŁAW ABRAMOWICZ, M.S.

Oncological Molecular Biologist & Computational Scientist

RESEARCH EXPERIENCE

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

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

- 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 (preprocessing, 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. [manuscript in preparation]

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

"Genome-wide methods in cancer genetics". BASTION. Advisor: Prof. Marzanna Cechowska-Pasko, Ph.D.

- 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). [manuscript in preparation]



DRIVING LICENCE

Currently 30/08/2006 AM/B₁/B

Smooth driving, dynamic driving, no collisions.

▲ Download this resume or cover letter

Example of microarray analysis

CONTACT INFO

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CAbout Me

M WARZEM

SKILLS





Mt Markdown, LTEX

HPC (openMPI, Rmpi). HTML5, CSS, JS, YAML, 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 (urology inflammations)

LANGUAGES

Polish: Native

English: Proficient (C1)



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, 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 (rmarkdown, bookdown, etc.);
- Pathway Enrichment Analysis (GO, KEGG, Reactome, WikiPathways, HumanCyc, clusterProfiler, rSEA);
- Multivariate Analysis (PCA, tSNE, UMAP, SONG, NMF, Ricci flow);

- · Comparative Analysis (Venn diagram);
- · Isobolograms;
- · use of data sets (ncbi, google patents, etc.);
- use of FDA resources (CDER, CDRH, NCTR mitochondial toxicity, MAQC/SEQC);
- · numerical validation awarness;
- · Linux command line;
- · basics of MySQL;
- basics of Python (PyScripter, PyMol, PyLasso, Autodock/Vina);
- basics of machine Learning (Random forest, CatBoost, LightGBM, XGBoost, H2O);
- · basics of neural Networks (TensorFlow, Keras, etc.);
- · Agilent's GeneSpring GX;
- · CLC Main Workbench;
- · more information in each description of course;

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

2017 | 2011

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 anti-inflammation, 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)

- RMongo, RPostresSQL;
- · dplyr, httr, rhdf5, xml, RMySQL, foreign, rmongodb, · 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

O PI

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, research at the academic level (30h); 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
- · Professional practice (30h);
- · OHS (4h).



M.Sc. in Medical Analitics, PQF 7

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

PL

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

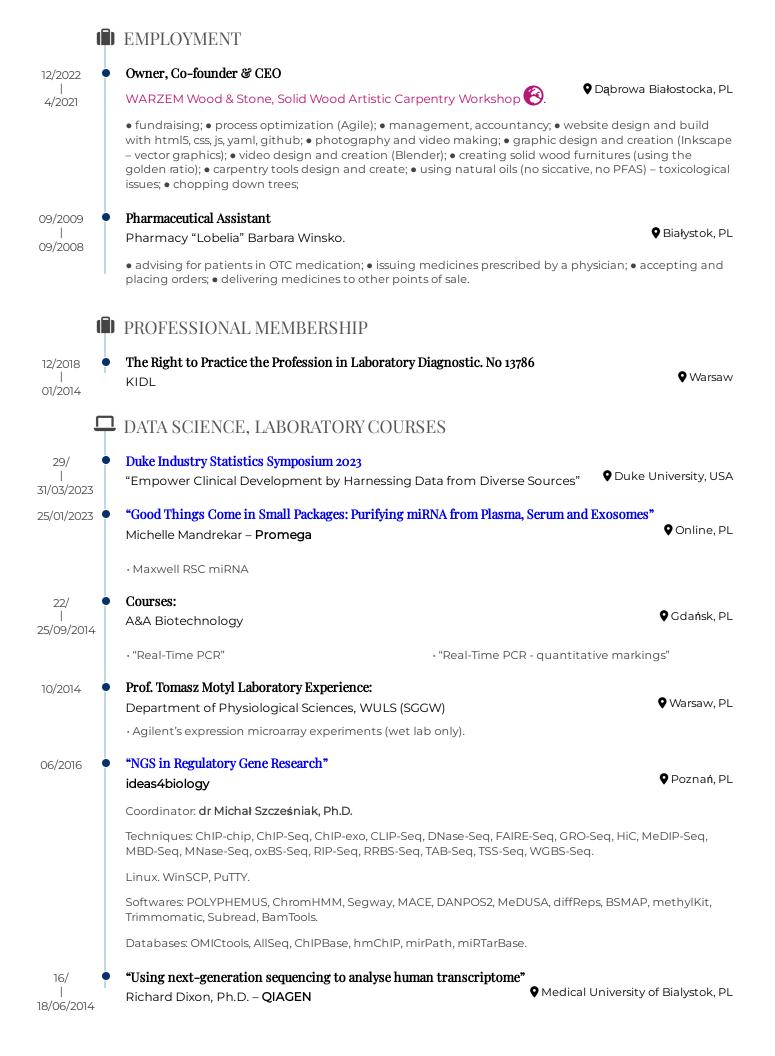
- · GPA: 4,30 (transcript of master degree);
- · isolation and work with human blood cells;
- · Advisor: Prof. dr hab. Ewa Jabłońska. Ph.D.
- · Co-Advisor: dr Kamil Grubczak, Ph.D.
- · experience in **biochemistry** and **immunology** research (see Wet lab, Student membership & Publications sections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

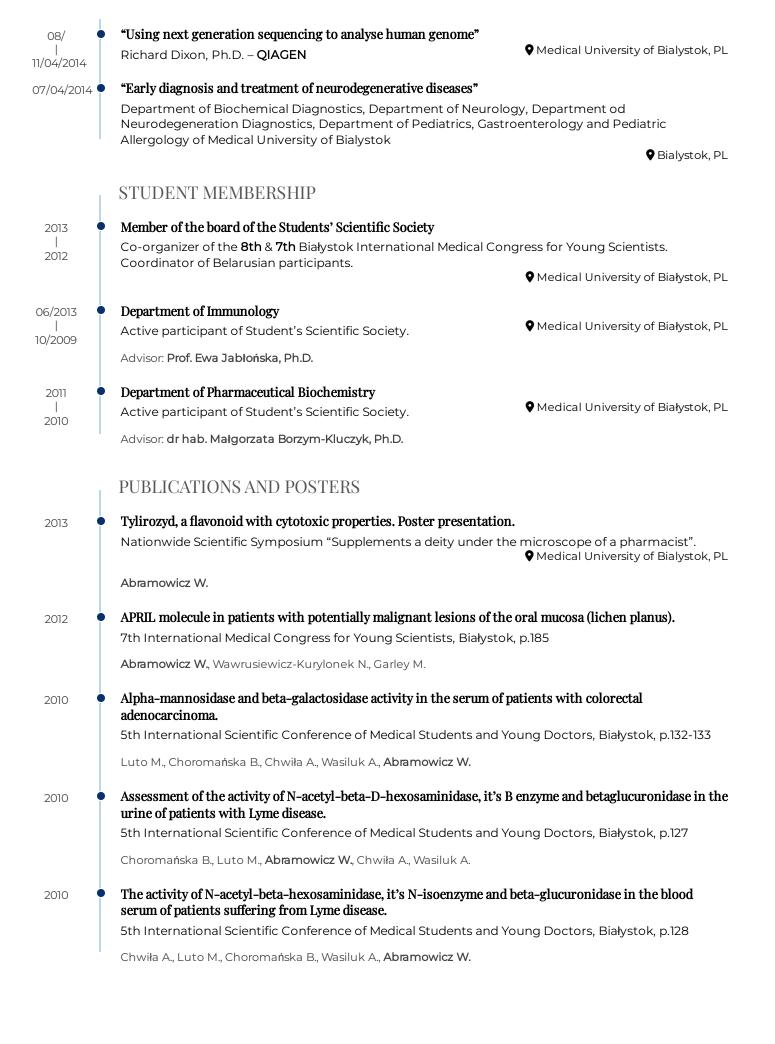


Pharmaceutical Assistant, PQF 4

Vocational School No. 1 of Health Care

Białystok, PL





2010

Alanine and aspartic aminotransferase activity in blood serum of men after acute and chronic ethyl alcohol poisoning.

5th International Scientific Conference of Medical Students and Young Doctors, Białystok, p.137-138 Wasiluk A., **Abramowicz W.**, Luto M., Choromańska B., Chwiła A.