## WIESŁAW ABRAMOWICZ, M.S.

Laboratory Diagnostician, Oncological Molecular Biologist & Computational Scientist (genomics, epigenomics, transcriptomics, epitranscriptomics, proteomics, metabolomics)

Achivements: cost effective sellecting personalized transcriptomics targets by microarray for on- and off-label treatment of tumors

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Example of microarray analysis

## ☐ RESEARCH EXPERIENCE

Transcriptomic profiling of a human kidney cancer cell line, Caki-2, after treatment of Ixazomib (on-label), Ursolic acid and jointly using microarray technology. (pre-clinical)

Department of Pharmaceutical Biochemistry, Medical University of Białystok.

9

- · 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. [manuscript in preparation]

8/2017 - 10/2013

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).



### DRIVING LICENCE



#### AM/B<sub>1</sub>/B

Smooth driving, dynamic driving, no collisions.



#### PROFESSIONAL MEMBERSHIP

Currently ( | 01/2014 

#### CONTACT INFO

☑immunol22@gmail.com

+48 798 298 280

**P**About Me

WARZEM Wood & Stone -

– UI/UX/ Graphic Designer, Carpenter.

#### **SKILLS**

A Linux, R Studio,



Markdown, LATEX,

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 (urology inflammations)

#### LANGUAGES

English: Proficient (C1)

Polish: Native



#### 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 (rmarkdown, bookdown, etc.);
- Pathway Enrichment Analysis (GO, KEGG, Reactome, WikiPathways, HumanCyc, clusterProfiler, rSEA);
- · Functional Annotation;
- · GATK, STAR, DESeq2;
- 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);
- · Gene Expression Omnibus;
- · 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);
- 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 · Statistical modeling (30h; prof. Tomasz Burzykowski, research at the academic level (30h);
  - · 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

#### **III** EMPLOYMENT Owner, Co-founder & CEO 12/2022 Oppositor Dalaman Dala WARZEM Wood & Stone, Solid Wood Artistic Carpentry Workshop 4/2021 • 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; Pharmaceutical Assistant 09/2009 Białystok, PL Pharmacy "Lobelia" Barbara Winsko. 09/2008 • advising for patients in OTC medication; • issuing medicines prescribed by a physician; • accepting and placing orders; • delivering medicines to other points of sale. 🖵 DATA SCIENCE, LABORATORY COURSES **Duke Industry Statistics Symposium 2023** 29/ Duke University, USA "Empower Clinical Development by Harnessing Data from Diverse Sources" 31/03/2023 "Good Things Come in Small Packages: Purifying miRNA from Plasma, Serum and Exosomes" 25/01/2023 Online, PL Michelle Mandrekar - Promega · Maxwell RSC miRNA Courses: 22/ Odańsk. PL A&A Biotechnology 25/09/2014 · "Real-Time PCR" · "Real-Time PCR - quantitative markings" Prof. Tomasz Motyl Laboratory Experience: 10/2014 Warsaw, PL Department of Physiological Sciences, WULS (SGGW) · Agilent's expression microarray experiments (wet lab only). "NGS in Regulatory Gene Research" 06/2016 Poznań, PL ideas4biology Coordinator: dr Michał Szcześniak, Ph.D. Techniques: ChIP-seq, ChIP-seq, ChIP-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.

"Using next-generation sequencing to analyse human transcriptome"

Richard Dixon, Ph.D. - QIAGEN

16/

08/

11/04/2014

18/06/2014

₱ Medical University of Bialystok, PL

"Using next generation sequencing to analyse human genome" Richard Dixon, Ph.D. – QIAGEN

Medical University of Bialystok, PL

07/04/2014		"Early diagnosis and treatment of neurodegenerative diseases"		
07/04/2014		Department of Biochemical Diagnostics, Department of Neuro Neurodegeneration Diagnostics, Department of Pediatrics, Gas Allergology of Medical University of Bialystok		
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	ī	STUDENT MEMBERSHIP		
2013		Member of the board of the Students' Scientific Society		
 2012		Co-organizer of the <b>8th</b> & <b>7th</b> Białystok International Medical Co Coordinator of Belarusian participants.	ongress for Young Scientists.	
		Coordinator of Belardslarr participants.	<b>♥</b> Medical University of Białystok, PL	
06/2013		Department of Immunology		
10/2009		Active participant of Student's Scientific Society.	◆ Medical University of Białystok, PL	
.,		Advisor: <b>Prof. Ewa Jabłońska, Ph.D.</b>		
2011		Department of Pharmaceutical Biochemistry		
 2010		Active participant of Student's Scientific Society.	◆ Medical University of Białystok, PL	
	I	Advisor: <b>dr hab. Małgorzata Borzym-Kluczyk, Ph.D.</b>		
		PUBLICATIONS AND POSTERS		
2013		Tylirozyd, a flavonoid with cytotoxic properties. Poster presentation.		
		Nationwide Scientific Symposium "Supplements a deity under t	the microscope of a pharmacist".  • Medical University of Bialystok, PL	
		Abramowicz W.		
2012		APRIL molecule in patients with potentially malignant lesions of the oral mucosa (lichen planus).		
		7th International Medical Congress for Young Scientists, Białysto	ok, p.185	
		Abramowicz W., Wawrusiewicz-Kurylonek N., Garley M.		
2010		Alpha-mannosidase and beta-galactosidase activity in the serum of patients with colorectal adenocarcinoma.		
		5th International Scientific Conference of Medical Students and	Young Doctors, Białystok, p.132-133	
		Luto M., Choromańska B., Chwiła A., Wasiluk A., <b>Abramowicz W.</b>		
2010		Assessment of the activity of N-acetyl-beta-D-hexosaminidase, it's urine of patients with Lyme disease.	B enzyme and betaglucuronidase in the	
		5th International Scientific Conference of Medical Students and	Young Doctors, Białystok, p.127	
		Choromańska B., Luto M., <b>Abramowicz W.</b> , Chwiła A., Wasiluk A.		
2010		The activity of N-acetyl-beta-hexosaminidase, it's N-isoenzyme an serum of patients suffering from Lyme disease.	d beta-glucuronidase in the blood	
		5th International Scientific Conference of Medical Students and	Young Doctors, Białystok, p.128	
		Chwiła A., Luto M., Choromańska B., Wasiluk A., <b>Abramowicz W</b> .		
2010		Alanine and aspartic aminotransferase activity in blood serum of n alcohol poisoning.	notransferase activity in blood serum of men after acute and chronic ethyl	
		5th International Scientific Conference of Medical Students and	Young Doctors, Białystok, p.137-138	
	1	Wasiluk A., <b>Abramowicz W.</b> , Luto M., Choromańska B., Chwiła A.	Last updated on 2023-	