

ANNUAL REPORT

ARMENIAN BIOINFORMATICS INSTITUTE

The Armenian Bioinformatics Institute (ABI) is a non-profit private foundation, launched in February of 2021 to support bioinformatics capacity building and the development of precision medicine and modern biotechnologies. This report summarizes the third year of our activities.

Prepared by:

Araz Chiloyan (content)
Karine Shahgaldyan (content)
Sona Melikjanyan (design)

29 December 2023

ARMENIAN BIOINFORMATICS INSTITUTE Scientific Educational Foundation



CONTENT

23 **INTRODUCTION CONFERENCES** & EVENTS 27 **INFRASTRUCTURE PEOPLE** 31 **CELEBRATING EDUCATION INDIVIDUAL ACHIEVEMENTS** 33 **RESEARCH FUNDING**

INTRODUCTION

Empowering Armenia's Bioinformatics Landscape: A Year of Growth and Impact

Bioinformatics and genomics have crucial importance for the development of life sciences, biotechnologies and medicine in Armenia. In 2021, we conducted an assessment that underscored the escalating demand for bioinformatics specialists and the insufficiency of efforts to meet the growing demands of the human workforce. Fast-forward to 2023, and we've witnessed a significant surge in the requirement for bioinformaticians within both local and global pharmaceutical and biotechnology companies, as well as the clinical sector, particularly in the realm of genetic testing initiatives.

Addressing this burgeoning demand for human resources necessitates a strategic scaling effort. This motivated the establishment of the Armenian Bioinformatics Institute (ABI) as a scientific educational foundation in 2021. Our mission is to train bioinformatics specialists not only with technical expertise but also possessing leadership skills and values that will empower them to drive transformative changes in the biotechnology and biomedical sectors in Armenia.

The year 2023 marked several milestones for ABI. Through a concerted fundraising initiative, we successfully secured a high-performance compute node, backed by generous donations, infrastructure support and research grants, amounting to a substantial budget of nearly 60,000 USD. Currently, this node caters to the needs of approximately

30 researchers at ABI and the Institute of Molecular Biology National Academy of Sciences (IMB NAS) of the Republic of Armenia (RA), with plans for further expansion to support numerous research, educational and clinical projects across the country. Its primary function is to provide a secure and efficient platform for storing, sharing, and analyzing genomics datasets.

In 2023, we published several findings in the human and plant genomics and microbiome fields. We applied machine learning and other bioinformatics methodologies to explore the molecular pathomechanisms and treatment response heterogeneity of cancer subtypes. In another study, we identified new genetic determinants for adaptation of grapevine varieties to changing climate conditions. Finally, as a collaborator in a major study in Europe, we contributed to the development of a novel and rapid testing approach for antibiotic susceptibility.

This year, our partnerships with two prominent global biotechnology companies, based in the US and the UK, expanded within the realm of cancer precision medicine. Furthermore, a third collaboration, focusing on microbiome research, commenced towards the end of the year. These collaborations lay the foundation for the evolution of a modern and vibrant life sciences landscape in Armenia.

The invaluable support from our community has been pivotal in achieving goals we set for ourselves for 2023. Simultaneously, we

have made substantial progress towards self-sustainability, with grants and industry collaborations accounting for a significant portion of our expenses, surpassing the 300,000 USD mark in 2023.

Looking ahead, our objective for the coming years remains the development of sustainable human resource potential in the life sciences. For ABI as an organization, this entails a strategic balance between expanding research teams and the addition of senior staff to enhance efficiency and facilitate scaling.



ARMENIAN BIOINFORMATICS INSTITUTE

2023 in Numbers



Research Groups/Labs

2



Research Grants

4



Industry Collaboration

2



Papers

9



Research Team

22



Alumni

8



Courses

4



Students

65

INFRASTRUCTURE

Compute Node Acquisition

Genome bioinformatics involves handling large volumes of genomic sequencing data. This requires computational resources and infrastructure with substantial storage capacity, memory, and CPU cores. In 2022 we initiated a fundraising campaign to acquire a dedicated highperformance compute node that would not only serve ABI but also support other research, educational and clinical projects. With the backing of several grant funding agencies, as well as approximately 100 donor organizations and individuals via the ReArmenia platform, ABI successfully acquired a new compute node in 2023 and established a state-of-the-art genome bioinformatics computing infrastructure. Major organizations supporting this initiative included the Higher Education and Science Committee of the Republic of Armenia (through a development grant), FAST the Foundation for Armenian Science and Technology (ADVANCE research grant), Flagship Pioneering Inc., SASTIC - Strategic Armenian Science & Technology Investment Community- (infrastructure support grant), SmartClick (donation), and many more.



Our new compute node is a 2x AMD EPYC 7702 (dual-socket 64 core) system with 2TB of memory and 240TB of storage, meeting the distinct space and memory requirements needed by our researchers at ABI. The node was acquired with the technical support of Alpha Engineering. In addition to infrastructure development, the campaign's success has allowed for the recruitment of a dedicated system administrator who has set up the node and data center and ensures its operational maintenance.

The server is meticulously maintained with over thirty individuals having access from both ABI and the IMB NAS RA. Our commitment is to not only maintain the current computing environment but

also to continuously improve and expand accessibility. We are actively working to ensure the server's optimal functionality, regularly updating and uploading all necessary tools and software for research staff. To further enhance our support infrastructure, we are in the process of expanding the system administration team. Our overarching goal is to extend access beyond ABI and IMB NAS RA, reaching educational and research institutes, as well as eventually clinical centers. This initiative aligns with our commitment to fostering widespread accessibility to cutting-edge bioinformatics resources, ultimately benefiting the broader scientific and medical communities.

EDUCATION

Courses

Algorithms for Computational Biology and Advanced Algorithms for Computational Biology

Both of these two-month long courses, tailored for bioinformatics students and researchers, provided a comprehensive exploration of informatics skills, particularly in the realm of computational biology with an emphasis on sequence alignment methods. Topics encompassed fundamental algorithms, data structures, and concepts. Led by Dr. Pesho Ivanov, a seasoned professional with expertise in combinatorial algorithms for biological problem-solving, the course included weekly coding challenges and bonus research tasks for interested participants. Pesho, who holds a background in applied mathematics and informatics, drew on his experiences as a teacher and programmer, showcasing projects in single-cell RNA-seq, T-cell receptors, phylogenetic tree intersection, and sequence assembly scaffolding. Pesho holds a PhD from ETH Zurich and is currently a postdoctoral researcher at the Pennsylvania State University in the United States.

Led by



Pesho Ivanov, PhD Pennsylvania State University, USA

Dates 28.02.2023-28.04.2023

Led by



Aleksei Kurnosov, PhD Russian-Armenian University, Armenia



Lilit Nersisyan, PhD Armenian Bioinformatics Institute, Armenia

Dates 13.01.2022-31.07.2023

Led by



Lilit Grigoryan, PhD Stanford University, USA

Dates 20.07.2023-10.08.2023

Advanced Molecular Biology

This year-long course covered the core concepts of molecular and cellular biology, genomes and genetic mechanisms, and much more. Topics included DNA replication, membrane structure, cell communication, cancer, reproduction, development, tissues, and the immune system. Dr. Aleksei Kurnosov initially led the course and it was continued with Dr. Lilit Nersisyan. Dr. Kurnosov is a Molecular Biologist with a PhD from

the Institute of Bioorganic Chemistry in Moscow, and has a keen interest in education. Dr. Kurnosov has also taught at the Russian-Armenian University in Yerevan.

Basics of Immunology

This two week minicourse covered the rapidly advancing field of the immune system in humans and across other species. Students discovered how every cell in a multicellular organism, from the tiniest microorganism to the largest vertebrates, fends off harmful invaders like bacteria, viruses, fungi, and larger parasites. There was a focus on the two layers of defense mechanisms in vertebrates

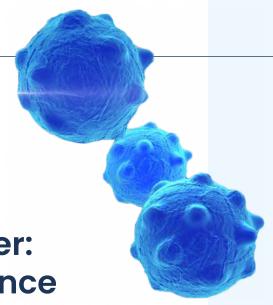
such as humans – the rapid, general "innate" responses, and the precise, specialized "adaptive" responses.
The course was led by Lilit Grigoryan, who holds a PhD in Immunology from Stanford University in the United States. Dr. Grigoryan had also previously served as a guest lecturer in immunology at ABI's OMICSS-21 and -22 summer schools.

Capstones

In 2023, several ABI researchers graduated from their university programs defending theses covering the projects they have been working on at ABI.

			0+
Emma Hovhannisyan			
Bachelor's degree	"Genetic Variability and Disease Resistance Analysis of Grapevine from Armenia"	American University of Armenia (AUA)	Supervisor: Hans Binder, PhD Co-Supervisor: Maria Nikoghosyan
Nelli Vardazaryan			
Master's degree	"The Effects of Geographical and Nutrient Variation on Bee Microbiome Composition: a Meta-Analysis"	Institute of Molecular Biology (IMB), National Academy of Sciences (NAS) of Armenia	Supervisor: Lilit Nersisyan, PhD
Yeva Bareghamyan			
Master's degree	"Mapping Gene Markers of Therapy Response in Liver Metastases of Colorectal Cancer"	Yerevan State University (YSU)	Supervisor: Arsen Arakelyan, PhD
Anush Baloyan			
Bachelor's degree	"Validating Population Structure Analysis Using Self Organizing Maps, a Case Study of Armenian Grapevines"	American University of Armenia (AUA)	Supervisor: Hans Binder, PhD

RESEARCH



Metastatic Cancer: Resisting Resistance

Liver metastases significantly exacerbate the prognosis for various cancer diseases, including colorectal cancer. Our exploration of the molecular mechanisms underlying this phenomenon utilized bioinformatics and gene expression data from hundreds of tumor samples. Through our analysis, we uncovered that metastasis and prior treatments can give rise to fibrotic, non-immunogenic tumor states that elude

standard therapies, including immunotherapies, by acquiring epigenetic plasticity in cancer cells.

Our studies advocate for potential strategies to overcome treatment resistance through refined precision medicine options. This project will continue through collaboration with and support from Agenus, aiming to pave the way for innovative solutions in the battle against cancer.

Principal Investigator Hans Binder, PhD

agenus

In collaboration with



Ashekyan O, Shahbazyan N, Bareghamyan Y, Kudryavzeva A, Mandel D, Schmidt M, Loeffler-Wirth H, Uduman M, Chand D, Underwood D, Armen G, Arakelyan A, Nersisyan L, Binder H. **Transcriptomic Maps of Colorectal Liver Metastasis: Machine Learning of Gene Activation Patterns and Epigenetic Trajectories in Support of Precision Medicine.** *Cancers.* 2023;15:15:3835. doi:10.3390/cancers15153835





Principal Investigator Hans Binder, PhD

FAST DISCOVERING THE FUTURE

In collaboration with



The grapevine stands as one of the earliest domesticated fruit crops, revered for both its fruit and its contribution to winemaking. Armenia, recognized as one of the ancient origins of grapevine domestication and wine production, holds a rich heritage in this regard. Our research endeavors focus on unraveling the diversity of Armenian vine accessions through a comprehensive analysis of their whole genomes.

To achieve this, we employed DNA sequencing and harnessed the power of machine learning knowledge mining. Through these methods, we gained insights

into the domestication history of grapevines and their correlations with phenotypic traits, including cultivar utilization and a potential resistance against powdery mildew.

The implications of our findings extend to addressing challenges in viticulture, particularly those posed by climate change. This project, one of the nine Advanced Research Programs supported by FAST, is intricately woven into international collaborations with partners in Georgia, central Europe, and China. Through our collective efforts, we aim to contribute to viticulture practices and foster innovation in the face of changing climate.

Margaryan K, Nikoghosyan M, Baloyan A, Gasoyan H, Hovhannisyan E, Galstyan L, Konecny T, Arakelyan A, Binder H. **Machine learned -based visualization of the diversity of vine genomes worldwide and in Armenia using SOMmelier.** *BIO Web Conf.* 2023;68, 01009. doi:10.1051/bioconf/20236801009

Konecny T, Nikoghosyan M, Binder H. Machine learning extracts marks of thiamine's role in cold acclimation in the transcriptome of Vitis vinifera. Front. Plant Sci. 2023;14. doi:10.3389/fpls.2023.1303542

Single-Cells in Space and Time

Cells serve as the fundamental building blocks within tissues and organs, orchestrating life in both space and time, thereby impacting processes related to disease and aging. Genomic sequencing technologies have entered the era of spatial single-cell resolution.

A dedicated team of junior researchers at ABI analyzes single-cell transcriptomics data, in cooperation with wet lab scientists from Leipzig University. Our studies focus on elucidating the mechanisms

underlying wound healing in the skin, unraveling the spatial microarchitecture of skin tumors (melanoma), and uncovering molecular communication links within the mouse brain.

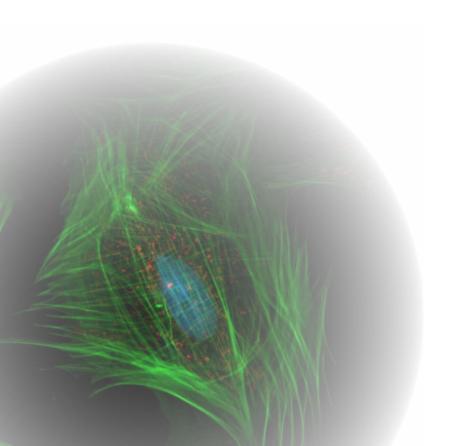
We have discovered that fibroblasts undergo identity changes during skin development and exert a profound influence on the invasiveness of cancer cells. Through this interdisciplinary collaboration, we aim to advance our understanding of health, disease, and aging processes on a cellular level.

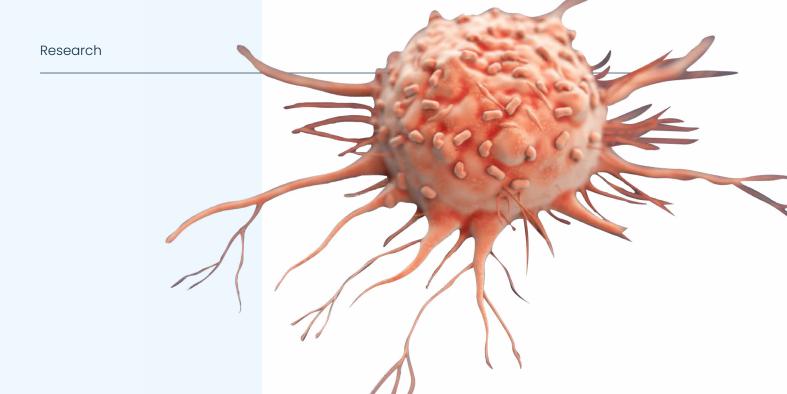
Principal Investigator

Hans Binder, PhD

In collaboration with







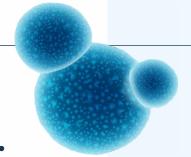
Fighting Cancer: Harnessing Flies for Precision Medicine

Team leads Hans Binder, PhD Nicolas Borisov, PhD

Funding
VIVAN

Vivan collaborates with ABI to bridge the gap in cancer drug research. Fruit flies, serving as experimental models, offer a unique avenue for testing drug combinations tailored for precision medicine. Since 2022, ABI has led a team of bioinformaticians and data scientists in direct involvement with Vivan's R&D. Our research centers on genetic biomarker discovery for precision cancer diagnostics and treatment decisions, integrating

machine learning and drug screening with genetic fly models. This collaboration aims to advance personalized cancer therapeutics, leveraging innovative approaches to enhance treatment options.



Microbiome Dynamics: the Effects of Disease

Our bodies are inhabited by thousands of diverse species of microbes, such as bacteria and fungi, forming a complex ecosystem. Microbes cooperate or compete in different parts of the food chain, as they interact with their host. Our team is developing new computational tools to explore the chain of events that are involved in altering the composition of microbial communities, and to determine how such changes may lead to diseases in human patients.

In collaboration with
Karolinska Institutet (Sweden)
and the IMB NAS RA, ABI
investigates the response of
microbial communities to
antibiotics, focusing on drugresistant Staphylococcus
aureus strains found locally
in Armenia. Utilizing novel
sequencing technologies
and a broad range of

data analysis approaches, we investigate changes happening at the level of DNA and RNA upon antibiotic perturbations. Similar to humans, honeybees harbor thousands of gut bacteria, collectively known as the gut microbiome. The gut microbiome composition is an important factor regulating the health and social behavior of honeybees and may affect the recently observed decline in global honeybee populations. In collaboration with Yerevan State University and Utrecht University (The Netherlands), we are exploring honeybee gut microbiome variations across diverse geographies, spanning the USA, Europe, Armenia, and New Zealand, and study their response to herbicides and pesticides using the most current sequencing and data analysis approaches.

Principal Investigator Lilit Nersisyan, PhD

In collaboration with



INSTITUTE OF MOLECULAR BIOLOGY

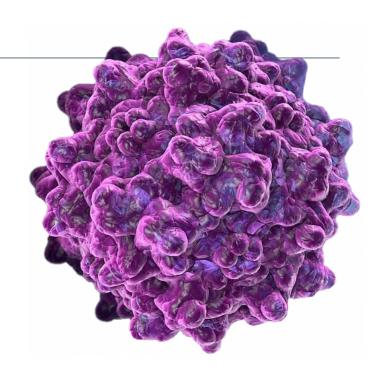






Huch S, Nersisyan L., Ropat M, Barrett D, Wu M, Wang J, Valeriano VD, Vardazaryan N, Huerta-Cepas J, Wei W, Du J, Steinmetz LM, Engstrand L, Pelechano V. **Atlas of mRNA translation and decay for bacteria.** *Nat Microbiol.* 2023;8:1123–1136

doi:10.1038/s41564-023-01393-z



Gene Therapies: Designing Vehicles for Disease Cure

Principal Investigator Lilit Nersisyan, PhD

In collaboration with





Some diseases are caused by the lack of a functional gene in the tissue preventing it from performing its functions. Gene therapies offer promise for treating such diseases. Gene therapies deliver the functional copy of a gene to the target tissue using viral capsules as delivery vectors. However, viral vectors are expensive to make and those add to the price of such therapies. Indeed, a recently released gene therapy to treat a rare genetic bleeding disorder called Haemophilia B, was coined as the most expensive drug on the market, costing \$3.5 million per dose. There is a need to optimize the production of these

vectors, to increase efficiency of delivery and reduce costs.

The team at ABI collaborates with the Wyss Institute at Harvard University and a USbased company Rejuvenate Bio to aid their efforts in designing more efficient viral vectors targeting age-related diseases. They utilize directed mutagenesis to prepare a diverse set of chimeric adeno-associated viral (AAV) vectors and select those that appear to be more efficient in the delivery of the gene to the target tissue. We developed a software tool to aid in the analysis and selection of the most efficient chimeric vectors.

Cell Fate Prediction: Single-Cell Developmental Trajectories

The cells in our bodies undergo functional and structural changes. During normal development, these changes give rise to the diversity of cell states forming the tissues and organs in our body. During disease, such changes may give rise to dysfunctional or malignant cells. Singlecell transcriptomics helps understand similarities between different cell types and build possible trajectories leading from one cell state to the other.

These trajectories are governed by changes in gene expression patterns

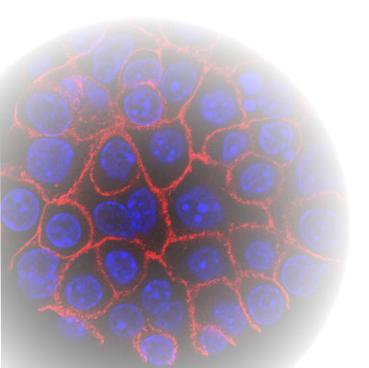
or RNA velocities of the genes. The single-cell RNA sequencing datasets not only contain information about the present state of gene expression patterns in the cells but also about the past and future states. With the support of a Science Committee development grant co-financed by Flagship Pioneering, our team utilizes the recent development of single-cell RNA sequencing technologies to use the hidden information about the past and future states of the cells and construct trajectories of cell state transitions that lead to healthy development or disease progression.

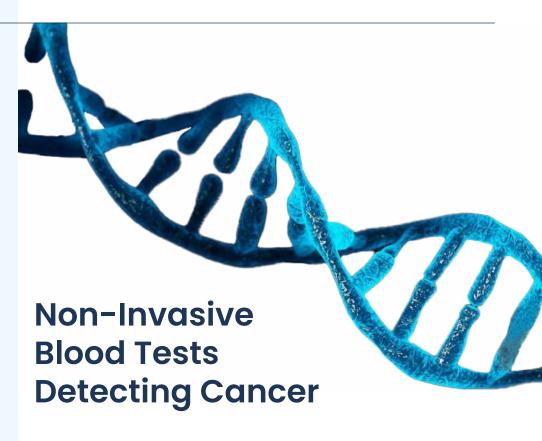
Principal Investigator Lilit Nersisyan, PhD

Funding









Principal Investigator Lilit Nersisyan, PhD

In collaboration with



University of Nevada, Reno

Blood is circulated by red blood cells that deliver oxygen to our tissues, the immune cells that fight infections, and the plasma that transports big and small molecules. Within the plasma, there is a small amount of circulating cell-free DNA, which originates from different tissues in our bodies. These DNA fragments have recently gained attention as possible biomarkers for noninvasive tests that could help us detect cancer and other diseases at earlier stages and/or frequently monitor the progress of cancer without the need for invasive biopsies. Such biomarkers are also referred to as liquid biopsies.

Our team concentrates on telomeres, specific chromosomal sites that are usually found at the tips of our chromosomes. Telomere length and sequence variation are known to contribute to the development of age-related diseases and cancers. Small amounts of telomeric sequences can also be found within circulating cellfree DNA pieces. Leveraging publicly available sequencing datasets and the tools we developed, we compare telomeric sequences in healthy individuals and those suffering from cancer and other disorders to discover cancer-specific biomarkers.

Publications

Ashekyan O, Shahbazyan N, Bareghamyan Y, Kudryavzeva A, Mandel D, Schmidt M, Loeffler-Wirth H, Uduman M, Chand D, Underwood D, Armen G, Arakelyan A, Nersisyan L, Binder H. **Transcriptomic Maps of Colorectal Liver Metastasis: Machine Learning of Gene Activation Patterns and Epigenetic Trajectories in Support of Precision Medicine.** *Cancers.* 2023;15:15:3835. doi:10.3390/cancers15153835

Avagyan S, Binder H. **Subtyping or not subtyping—Quo vadis for precision medicine of colorectal cancer**. *Transl Cancer Res. 2023;12:5:1067-1072*. doi: 10.21037/tcr-23-133

Everest-Dass A, Nersisyan S, Maar H, Novosad V, Schröder-Schwarz J, Freytag V, Stuke JL, Beine MC, Schiecke A, Haider MT, Kriegs M, Elakad O, Bohnenberger H, Conradi LC, Raygorodskaya M, Krause L, von Itzstein M, Tonevitsky A, Schumacher U, Maltseva D, Wicklein D, Lange T. **Spontaneous metastasis xenograft models link CD44 isoform 4 to angiogenesis, hypoxia, EMT and mitochondria-related pathways in colorectal cancer.** *Mol Oncol.* 2023. doi:10.1002/1878-0261.13535

Hakobyan S, Stepanyan A, Nersisyan L, Binder H, Arakelyan A. **PSF toolkit: an R package for pathway curation and topology-aware analysis.** *Front Genet.* 2023; 14:1264656. doi:10.3389/fgene.2023.1264656

Huch S, Nersisyan L., Ropat M, Barrett D, Wu M, Wang J, Valeriano VD, Vardazaryan N, Huerta-Cepas J, Wei W, Du J, Steinmetz LM, Engstrand L, Pelechano V. **Atlas of mRNA translation and decay for bacteria.** *Nat Microbiol.* 2023;8:1123–1136. doi:10.1038/s41564-023-01393-z

Konecny T, Nikoghosyan M, Binder H. **Machine learning extracts marks of thiamine's role in cold acclimation in the transcriptome of Vitis vinifera.** *Front. Plant Sci. 2023;14.* **doi:10.3389/fpls.2023.1303542**

Margaryan K, Nikoghosyan M, Baloyan A, Gasoyan H, Hovhannisyan E, Galstyan L, Konecny T, Arakelyan A, Binder H. **Machine learned -based visualization of the diversity of vine genomes worldwide and in Armenia using SOMmelier.** *BIO Web Conf.* 2023;68, 01009. doi:10.1051/bioconf/20236801009

Shkurnikov M, Nersisyan S, Averinskaya D, Chekova M, Polyakov F, Titov A, Doroshenko D, Vechorko V, Tonevitsky A. **HLA-A*01:01 allel e diminishing in COVID-19 patients population associated with non-structural epitope abundance in CD8+ T-cell repertoire.** *PeerJ.* 2023;11:e14707. **doi:10.7717/PEERJ.14707**

Zhiyanov A, Engibaryan N, Nersisyan S, Shkurnikov, M, Tonevitsky A. **Differential** co-expression network analysis with DCoNA reveals isomiR targeting aberrations in prostate cancer. *Bioinform*. 2023:39:2:btad051. doi:10.1093/bioinformatics/btad051w

CONFERENCES & EVENTS



Viticulture 2.0: A Conference in Armenia

Renowned institutions and viticulture experts from Armenia, Sweden, France, and Germany convened for the "Viticulture 2.0: From Science to Vineyards" workshop, hosted by ABI and the IMB NAS RA on May 3 and 4. Supported by Alliance of National and International Science Organizations (ANSO) (Armenia-China-France) and ProGriCC (Production of Grapes in a Changing Climate) projects, the event featured insightful presentations and a captivating journey to the heart of Armenian viticulture in Vayots Dzor.



Grapevine Genomics Summer School in Italy

From June 25 to 28, Anush Baloyan and Tomas Konecny, sponsored by GRAPEDIA (Grapevine Genomics Encyclopedia), a platform aimed at establishing open access for the grapevine research community, attended the <u>summer school</u> at the Edmund Mach Foundation in Trento, Italy. The program provided insights into Grapedia's technology architecture, prototype progress, and upcoming challenges in grapevine data analysis. This training, targeting the scientific community, aimed to shape Grapedia's future through lectures and hackathon sessions.



Machine Learning Summer School in the UK

Hripsime Gasoyan attended the Oxford Machine Learning Summer School (OxML) from July 8 to 16, with funding from FAST (Foundation of Armenian Science and Technology). The program comprised MLx Fundamentals, covering topics such as linear algebra, optimization, and statistical machine learning, and MLx Cases, focusing on real-world ML issues, including data collection, transfer learning, and model interpretation.



Grapevine Genomics Workshop in Armenia

A collaborative mini-workshop at FAST (Foundation of Armenian Science and Technology), took place on September 5, organized by the ABI Vine group, connected experts from Armenia and Georgia through the Swedish Institute funded project (ProGriCC grant), as well as potential stakeholders from the vine industry, Armenian Vine and Wine Association and Armenian National Agrarian University (ANAU). Discussions centered on vine bioinformatics, the ADVANCE project, genetic diversity in Armenian grapes, Georgia's soils, vine genomes, functional genomics, and most importantly, future cooperation in all directions.



GRAPEDIA Annual Meeting in Spain

The GRAPEDIA (Grapevine Genomics Encyclopedia) Annual Meeting in Valencia, Spain, held from September 11 to 13, gathered computational biology experts for grapevine research. Participants shared insights on databases and FAIR genomic data-sharing principles. Notably, Maria Nikoghosyan delivered an oral presentation titled "SOMmelier: analyzing genomic data and annotating grapes, particularly Armenian associations."



Wine Industry and Global Climate Change Challenges: A Workshop in Georgia

ABI actively participated in the five-day workshop titled "Wine Industry and Global Climate Change Challenges" held in Telavi, Georgia, from October 2 to 6. Dr. Hans Binder, Armine Asatryan, Emma Hovhannisyan, and Tomas Konecny, representing ABI, shared valuable insights into the ongoing advancements in vine genomics and bioinformatics.





Lilit Nersisyan participated in the EMBL-EBI course on "Managing a Bioinformatics Core Facility," gaining valuable insights into optimal practices for overseeing local and national bioinformatics infrastructures from November 7 to 9 in Hinxton, UK.



Exploring Vine-Related Bioinformatics: ProGriCC Winter Workshop in Sweden

The ProGriCC winter workshop in Uppsala, Sweden took place from December 11 to 14. This workshop served as a gathering for ProGriCC partners and researchers from GRAPEDIA.

The primary objective was to discuss the vine-related bioinformatics program and explore its continuation in the upcoming years. Both Dr. Hans Binder and Tomas Konecny presented at the Public Symposium on "Research in the Field of Winemaking under Climate Change."

PEOPLE

Director

Lilit Nersisyan, PhD

Board of Trustees

Hans Binder, PhD Chairman University of Leipzig, Germany Arsen Arakelyan, PhD **Board** member Institute of Molecular Biology National Academy of Sciences (IMB NAS), Armenia Aram Adourian, PhD **Board Member** Flagship Pioneering, USA

Advisory Board

Jonathan Schug, PhD University of Pennsylvania, **USA**

Hovhannes Harutyunyan, MBA Pilant, Germany

Rafi Baghdjian Center for Scientific

Innovation and Education Foundation, Armenia

Peter Stadler, PhD University of Leipzig, Germany

Hripsime Hovakimyan Kuzevska K.A.K. Doo, Macedonia

Ogsen Gabrielyan, PhD Boehringer Ingelheim, Germany

Eduard Avetisyan European Molecular Biology Laboratory (EMBL), Germany

Research Team

*Some of the researchers have contributed part time, remotely, and/or in-kind

Hans Binder, PhD, Pl

Siras Hakobyan,

PhD (c)

Nelli Vardazaryan,

PhD (c)

Lilit Galstyan, MSc (c)

Lusine Adunts.

BSc (c)

Meline Mkrtchyan,

MSc (c)

Kristine Margaryan, PhD

Sveta Mnatsakanyan,

MD Resident

Levon Galstyan,

MSc (c) (alumnus)

Ohanes Ashekyan,

PhD (c) (alumnus)

Lilit Nersisyan, PhD, Pl

Maria Nikoghosyan,

PhD (c)

Hripsime Gasoyan,

MSc (c)

Emma Hovhannisyan, BSc

Arpine Grigoryan,

BSc (c)

Shengchang Duan,

MSc (c)

Luiza Stepanyan, BSc (c)

Tatevik Jalatyan,

MSc (c) (alumna)

Nerses Shahbazyan,

MD (c) (alumnus)

Anna Kudryavtseva,

MSc (c) (alumna)

Nikolay Borisov, PhD

Tomas Konecny,

PhD (c)

Anush Baloyan,

BSc (c)

Melina Tamazyan, BSc

Gayane Aydinyan,

BSc (c)

Davit Tarverdyan,

BSc (c)

Armine Asatryan, PhD

Susanna Avagyan,

MSc (c) (alumna)

Daniil Igumnov,

MS (alumnus)

Daria Laricheva,

MSc (c) (alumna)

Operations Team

Karine Shahgaldyan

HR Administrative Assistant

Sona Melikjanyan

Graphic Designer

Volodya Poghosyan

Full-Stack Developer

Araz Chiloyan

Operations and

Development Manager

Antranik Wartanian

Systems Administrator

Mariam Hovsepyan

Legal Expert

Khachatur Ashotyan

Systems Administrator

Mentors and Volunteers

Jakub Wudarski, PhD University of Nevada

Erik Aznauryan, PhD Harvard University

Marta Tortegrossa, PhD Leipzig University

Susanna Avagyan, MSc (c) Stanford University

Sevag MakdissianBirthright Armenia, operations

Stepan Nersisyan, PhD Thomas Jefferson University

Arsen Arakelyan, PhD IMB NAS RA

Juliane Liou, PhD Leipzig University

Phillip Guevorgian YerevaNN

Tigran PetrosyanList.am, operations

Valeri Vardanyan, PhD University of Tokyo

Maria Schmidt, PhD Leipzig University

Chrats Melkonian, PhD Utrecht University

Ketrin Boyacioglu Birthright Armenia

Alumni

Tatevik Jalatyan

MSc (c) at University of Oxford, England

Daniil Igumnov

Researcher, GERO.AI, Serbia

Anna Kudryavtseva

Bioinformatician, First Moscow State Medical University named after I.M. Sechenov, Russia

Nerses Shahbazyan

Bioinformatics Scientist at BostonGene, Armenia

Susanna Avagyan

MS (c) at Stanford University, USA

Daria Laricheva

MSc (c) at Brandeis University, USA

Levon Galstyan

MS (c) at Johns Hopkins University, USA

Ohanes Ashekyan

PhD (c) at University of Ottawa, Canada

CELEBRATING INDIVIDUAL ACHIEVEMENTS



Avagyan

Editorial publication in
Translational Cancer
Research;

MSc Candidate at
Stanford University

Susanna



Jalatyan Chevening Scholarship for Oxford University; ABI Curiosity Award

Tatevik



Konecny

Publication in
Frontiers in Plant
Science

Tomas



Hakobyan

Publication in Frontiers in Genetics

Siras



Nikoghosyan

Oral presentation at GRAPEDIA
Annual Meeting

Maria



Emma
Hovhannisyan

ABI Curiosity

Award



Vardazaryan

Young Scientist
Support Program
Grant Recipient



Galstyan

MSc Candidate
at Johns Hopkins
University

Levon



Arpine
Grigoryan

Erasmus+ Program
at Lille University



Sean Mintz MIT Internship, MIT's MISTI program



Nerses
Shahbazyan
Publication

in Cancer



Ohanes
Ashekyan
Publication
in Cancer

FUNDING

Budget

In 2023, both our sources of income and expenses experienced a remarkable surge, more than tripling compared to the previous year. In our inaugural year, we relied exclusively on donations to drive the development of our human resources and lay the groundwork for diversifying our funding streams. However, over the past two years, our financial landscape has evolved significantly, with industry collaborations and research grants emerging as substantial contributors

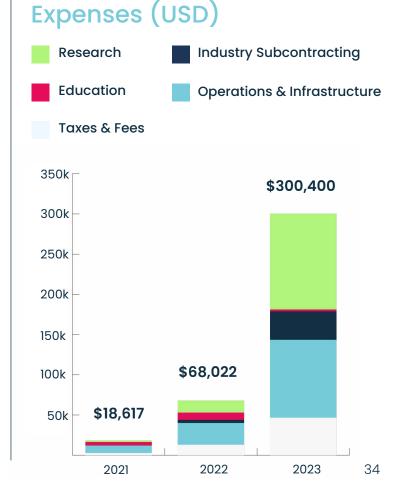
to our organization's income.

Notably, alongside donations, these collaborations and grants have played a pivotal role in fortifying our financial foundation. Equally crucial are the in-kind contributions made by numerous researchers who have generously dedicated their time to our projects. It's important to note that these invaluable contributions, though not explicitly reflected in our expense charts, have significantly

enriched the scope and impact of our initiatives.

Fueled by these combined efforts, we have successfully secured the necessary funds to embark on the continuous expansion of our team in 2024, ensuring the continuity and growth of our ongoing activities. Looking ahead, our commitment to furthering our mission remains steadfast, and we plan to sustain this momentum through continued fundraising activities, industry collaborations, and research grants.





Donors

As we strive for further growth and the realization of additional projects, we extend our gratitude to all those who have been instrumental in our journey so far. With the unwavering support of our private donors and organizations, we are poised for continued success in the pursuit of our organization's goals.

Donation Matching Organizations

Adobe Armenia Inc. Synopsys Inc. VMware AMD Inc. Netflix Spotify

Donation Platforms

Benevity Global Giving ReArmenia ARPA Institute

Industry Collaborations

agenus



Research & Infrastructure Support Grants











DONATE



donate.abi.am

FOLLOW US





abi.am