

Andreas Venizelos

Postdoctoral Researcher in Cancer Genomics

K.G.Jebsen Center for Genome Directed Cancer Therapy

Dept. of Clinical Science

University of Bergen, Bergen, Norway

- 13 August 1988
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- s dauidven
- andreas.venizelos@yahoo.com

Social Network -

https://linkedin.com/in/dr-andreasvenizelos-297802ab

Languages

Greek mother tongue English German

IT skills –

Nowegian Bokmål

- Programming language: Perl, Python, R, R markdown
- Operating Systems: MS Windows, Linux, MacOS
- Bioinformatics Applications: Bioconductor, Biopython, BLAST, GALAXY, Genome Browsers (e.g. UCSC, HGMD), Alamut, Picard, Samtools, Bedtools, Beftools, gatk, IGV, FastQC
- General IT applications: MS Office, Adobe Suite, MS Visual Studio, Docker

Areas of Expertise

Bioinformatics and Systems biology Clonal evolution analysis Next generation sequencing data analysis Structural Bioinformatics Machine Learning Applications Gene Regulatory pattern analysis Gene Ontology

Working Experience

2021 - Present Postdoctoral Researcher Mohn Research Laboratory, Bergen, Norway

- Analysis and interpretation of WGS, WES and panel sequencing
- Investigating genomic alterations and subclonal evolution dynamics in breast cancer tumours during neoadjuvant chemotherapy.
- Identifying the molecular mechanisms underlying the development of high-grade gastroenteropancreatic neuroendocrine carcinomas (NEC).
- Assessing the potential role of early epigenetic alterations as an underlying cause of NEC.
- Consulting services for Roche Norway AS (lecture).
- Representative of K.G. Jebsen Academy for Genome-Directed Therapy in Cancer.

2016 - 2021

Researcher

Mohn Research Laboratory, Bergen, Norway

- Performing genome assembly and annotation.
- Developing reproducible bioinformatic pipelines for next generation cancer sequencing efforts, including targeted capture assays, WES, RNA-seq.
- Validating bioinformatic tools and clustering algorithms for clonal evolution analysis using public and in-house samples from our clinical
- Analyzing in-house genomics/transcriptomics data from patientderived tumor cells, and developing interactive data visualizations.
- Performing assays, PCR, QC, DNA extraction, and other molecular biology techniques.
- Preparing DNA libraries for NGS and executing sequencing runs on Illumina Sequencers.
- Consulting services for Cepton Strategies, Paris, France (1h via phone).
- Committee member for the Research School of Clinical Medicine Tasks assigned: Meetings arrangement, reviewing all relevant material before committee meetings, generating abstract books-timetables for poster-oral presentations, preparing the monthly agenda, contacting individuals, approving reports of committee meetings before their distribution.

2013 - 2016

Senior Medical

Centogene A.G., Rostock, Berlin, Germany

Technologist

- Expertise in next generation sequencing (Roche 454, Ion Torrent, Illumina), Sanger sequencing, PCR, Real-Time PCR, MLPA, MS-MLPA, nucleic acid extraction, cell cultures, excellent knowledge of sequencing analysis softwares (Seq Pilot, Mutation Surveyor), different databases and genetic tools (Alamut, HGMD, OMIM).
- Technical validation of raw data obtained from different techniques
- Supervision of a team of 7 technicians, mentored and trained team members via technical presentations, demonstrations, troubleshooting technical issues.
- Active participation in the preparation of the laboratory for CAP and ISO-13485 inspections.

Tasks assigned: writing of SOP and validation files, engagement and participation in different committees (metrology, telemetry).

2012 - 2013

Medical Technologist

401 Military Hospital of Athens, Greece

- Expertise in Sanger sequencing, high-throughput SNP genotyping, nucleic acid extraction; developed protocols for the screening of new genes.
- Responsible for the accurate orientation of samples to external laboratories, contact person of the laboratory for all genetic tests.

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Lab skills

- Hands on library preparation for NGS platforms (WGS, WES, Gene/Custom Panels)
- Sequencing on Illumina platforms (MiSeq, NextSeq500, HiSeq 2000/2500, NovaSeq)
- Sequencing on Ion Torrent platforms (Ion-PGM, Ion-Proton, Ion-Chef)
- DNA/RNA extraction; Blood, Snap-Frozen Tissue, Formalin Fixed Paraffin Embedded (FFPE)
- Covaris Sonication (M220)
- DNA quantification (QuantoFluor, Qubit)
- Quantitative PCR, Long-range PCR
- Sizing, quantitation, purity assessment of DNA using Bio-analyzer 2100 Agilent and TapeStation
- Sanger Sequencing

Education

Postgraduate Training

2016 – 2021 Ph.D. in Cancer Genomics University of Bergen, Norway

Title dissertation: Mutational characterization of cancers; Studies of Breast Cancers under Chemotherapy and Neuroendocrine Carcinomas **Opponents**: Dr. Kristine Misund, Dr. Thomas Berg, Dr. Agnete

 ${\bf Engelsen}$

2011 – 2012 M.Sc. in Molecular Virology University of Ioannina, Greece

- Molecular characterization of viruses via genotyping, training in nucleic acid extraction, PCR, RT-PCR, gel electrophoresis, Sanger sequencing, cloning.
- Training in sequencing analysis softwares (SeqPilot, Mutation Surveyor), different databases and genetic tools (OMIM, Blast, UCSC Genome Browser).

• Mentored and trained undergraduate students.

GPA: 10/10

Title dissertation: G20 gene molecular characterization of

cyanophages

Opponents: Prof. Theoni Trangas, Prof. Evangelos Briasoulis

2007 – 2011 B.Sc. in Biological Applications University of Ioannina, Greece

and Technology GPA: 7.32/10

Publications

2022 Andreas Venizelos, Christina Engebrethsen, Wei Deng, Jürgen

Geisler, Stephanie Geisler, Gjertrud T. Iversen, Turid Aas, Hildegunn S. Aase, Manouchehr Seyedzadeh, Eli Sihn Steinskog, Ola Myklebost, Sigve Nakken, Daniel Vodak, Eivind Hovig, Leonardo A. Meza-Zepeda, Per E. Lønning, Stian Knappskog, Hans P. Eikesdal: Clonal evolution in primary breast cancers under sequential epirubicin and docetaxel

monotherapy Genome Medicine (under review)

Venizelos, A., Elvebakken, H., Perren, A., Nikolaienko, O., Deng,

W., Lothe, I.M.B., Hjortland, G.O., Sundlöv, A., Svensson, J., Garresori, H., Kersten, C., Hofsli, E., Detlefsen, S., Krogh, M., Sorbye, H., Knappskog, S.: *The molecular characteristics of high-grade gastroenteropancreatic neuroendocrine neoplasms*. Endocrine-Related Cancer,

2021 Nov;29(1):1-14

2021 H P Eikesdal, S Yndestad, A Elzawahry, A Llop-Guevara, B Gilje, E

S Blix, H Espelid, S Lundgren, J Geisler, G Vagstad, A Venizelos, L Minsaas et al: Olaparib monotherapy as primary treatment in unselected triple negative breast cancer. Ann Oncol. 2021 Feb;32(2):240-

249.

Oral presentations at international congress

2021 Sorbye, H., Venizelos, A., Elvebakken, H., Perren, A., Nikolaienko,

O., Deng, W., Lothe, I.M.B., Hjortland, G.O., Sundlöv, A., Svensson, J., Garresori, H., Kersten, C., Hofsli, E., Detlefsen, S., Krogh, M., Knappskog, S. title: Molecular characteristics of high-grade gastroen-

teropancreatic neuroendocrine neoplasmsESMO Congress 2021,

Paris, France

Venizelos, A., Elvebakken, H., Perren, A., Nikolaienko, O., Deng, W., Lothe, I.M.B., Hjortland, G.O., Sundlöv, A., Svensson, J., Gar-

resori, H., Kersten, C., Hofsli, E., Detlefsen, S., Krogh, M., Sorbye, H., Knappskog, S. title: Mutational landscape of 109 high-grade gastroenteropancreatic neuroendocrine neoplasms G317th Annual ENETS

Conference, Barcelona, Spain

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Posters at international congress

Venizelos, A., Clausen, C., Deng, W., Geisler, J., Geisler, S., Aas, T.,

Aase, H., Seyedzadeh, M., Steinskog, E.S., Myklebost, O, Nakken, S., Vodak, D., Hovig, E., Meza-Zepeda, L.A., Lønning, P.E., Knappskog, S.*, Eikesdal, H.P.* *title: Whole exome sequencing (WES) of locally*

 $advanced\ breast\ cancers\ treated\ with\ monotherapy$

AACR Annual Meeting 2019, Atlanta, USA
Venizelos, A., Clausen, C., Knappskog, S., Eikesdal, HP. and Lønning

PE. title: Genetic alterations affecting treatment response in locally

advanced breast cancers

7th CCBIO Annual Symposium, Solstrand, Bergen, Norway

Awards

2020 ENETS; Top 3 oral presentation; highlights selection; title: Mutational

landscape of 109 high-grade gastroenteropancreatic neuroendocrine

 $neoplasms\ G3$

17th Annual ENETS Conference, Barcelona, Spain

Personal

Inerests Crossfit, Hiking, Skiing Others Full, Clean driving licence References Available upon request