

FLORIN CHELARU

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WORK EXPERIENCE

Google Inc., Cloud Monitoring	Feb 2017
<i>Senior Software Engineer; Tech Lead; Engineering Manager</i>	–
Lead a team of engineers at Google, in the Cloud Monitoring group and designed large parts of a cloud resource metadata ingestion system consisting of several micro-services and an API. Some of the technologies used include Cloud Spanner, Cloud Pub/Sub and Kubernetes Platform.	Jan 2022
Twinfog Inc. angel.co/twinfog	July 2016
<i>Co-Founder & CTO</i>	–
Founded Twinfog, a geolocation-based Reddit for helping communities of expats. I designed and implemented the Twinfog cross platform mobile app using Xamarin (C# .NET) for the UI, and ASP.NET MVC, SQL Server and Azure Cloud Services for the backend. Demo video: youtu.be/oINy4qSrMiM	December 2017
MIT Computer Science and Artificial Intelligence Laboratory	August 2015
<i>Postdoctoral Associate</i>	–
Expanded on the work done for my Ph.D. by designing a series of open-source visualization libraries for <i>genetic variants</i> analysis. Base library code available at: github.com/florin-chelaru/vis.js .	June 2016
University of Maryland Center for Bioinformatics and Computational Biology	Jan 2011
<i>Graduate Research Assistant</i>	–
Research Assistant in the Center for Bioinformatics and Computational Biology, at the University of Maryland, College Park. I created and developed Epiviz (epiviz.org), a big data visualization and analysis tool for epigenetics and genomics. Featured in Nature Methods and adopted by Genentech. GitHub project: github.com/epiviz .	Jun 2015
Rocket Fuel Inc., Artificial Intelligence Team	2014,
<i>Software Engineer</i>	Jun–Sep
Worked on the Artificial Intelligence (Ad Prediction) team. Designed and implemented ML, probabilistic models and a visualization tool to improve conversion rates for online ads.	2013,
	Jun–Sep
Facebook Inc., Spam Detection Team (Site Integrity)	2012,
<i>Software Engineer</i>	May–Aug
Worked on the Spam Detection team, using Machine Learning models to aid in the detection of malicious users and content.	
University of Maryland Department of Computer Science	Jan 2011
<i>Graduate Teaching Assistant</i> for the following classes:	–
CMSC702 – Computational Systems Biology (Instructor: Dr. Hector Corrada)	Dec 2012
CMSC433 – Parallelism and Multithreading in Java (Instructors: Dr. Adam Porter, Dr. Tom Yeh)	
CMSC420 – Data Structures (Instructor: Professor Hanan Samet)	
Microsoft Inc., Office Team (Lync Server)	Jun 2010
<i>Software Engineer</i>	–
Designed database optimization software for improving the performance of the Lync Communication Server.	Jan 2011
Microsoft Inc., Bing Team (Search Domain Relevance)	Sep 2008
<i>Software Engineer in Test</i>	–
Designed and developed software for measuring the quality of web search results in the Bing search engine. Created a visualization tool for monitoring the quality and relevance of search result captions.	Jun 2010

	<p>Code40 Inc. Romania <i>Undergraduate Internship</i> Designed and implemented components of a web server application for micro-loans: caching, back-end data validation, error handling.</p>	<p>2007, Jul–Aug</p>
EDUCATION	<p>University of Maryland, College Park <i>Doctor of Philosophy, Computer Science</i> Advisor: Dr. Héctor Corrada Bravo</p> <p>Dissertation: Epiviz: interactive visual analytics software for genomics Relevant coursework: Machine Learning (H. C. Bravo); Information Visualization (B. Shneiderman); Neural Modeling (J. Reggia); Computational Linguistics (K. H. Seitz); Computational Genomics (C. Kingsford); Functional Genomics (H. C. Bravo); Computer Vision (Y. Aloimonos).</p>	<p>Jan 2011 – May 2015</p>
	<p>University Al. I. Cuza, Iași, Romania <i>Bachelor of Science, Computer Science</i> Class Rank: 8 of 176 Advisor: Dr. Liviu Ciortuz</p> <p>Bachelor's dissertation: Artificial Intelligence in Computer Go Relevant coursework: Machine Learning; Bioinformatics; Neural Modeling; Evolutionary Algorithms; Artificial Intelligence; Graph Theory; Algorithm Design; Probabilities and Statistics; Calculability, Decidability and Complexity; Cryptography; Antivirus Technologies; Software Engineering and Design Patterns; C/C++; C# and .NET Framework; Java; Relational Databases and SQL.</p>	<p>Sep 2004 – Jun 2008</p>
PUBLICATIONS	<p>J. Wagner*, F. Chelaru*, J. Kancherla*, J. N. Paulson*, A. Zhang, V. Felix, A. Mahurkar, N. Elmqvist, H. C. Bravo, “Metaviz: interactive statistical and visual analysis of metagenomic data”. <i>Nucleic Acids Research</i>, gky136, Feb. 2018 https://doi.org/10.1093/nar/gky136</p> <p>F. Chelaru* and H. C. Bravo, “Epiviz: a view inside the design of an integrated visual analysis software for genomics”. <i>BMC Bioinformatics</i>, 16 Suppl 11, S4. http://doi.org/10.1186/1471-2105-16-S11-S4</p> <p>F. Chelaru*, L. Smith, N. Goldstein, and H. C. Bravo, “Epiviz: interactive visual analytics for functional genomics data,” <i>Nature Methods</i>, vol. 11, no. 9, pp. 938–940, Aug. 2014. http://dx.doi.org/10.1038/nmeth.3038</p> <p>H. C. Bravo*, F. Chelaru, L. Smith and N. Goldstein, “epivizr: R Interface to epiviz web app,” Bioconductor package: 1.4.2.</p> <p>F. Chelaru* and L. Ciortuz, “Combining old-fashioned computer go with monte carlo go,” in <i>Proceedings of the 2008 10th International Symposium on Symbolic and Numeric Algorithms for Scientific Computing, SYNASC 2008</i>, 2008, pp. 216–222. http://dx.doi.org/10.1109/SYNASC.2008.77</p> <p>S. Iftene* and F. Chelaru, “The general Chinese remainder theorem,” in <i>International Scientific Journal of Computing</i>, vol. 6, issue 1, pp. 44-50, 2007. http://www.computingonline.net/archieve/IJC_2007_06_1_05.pdf</p>	
SOFTWARE PROJECTS	<p>Music with Ms. Johnson (vioara-cu-susanna.ro) — Jan 2023 github.com/florin-chelaru/music-with-susanna The professional presentation page of Susanna Johnson, violin, viola and general music teacher. <i>TypeScript, ReactJS, Material UI</i></p> <p>Epiviz (epiviz.org) — Aug 2014 github.com/epiviz A web visualization tool used to aid in the analysis and exploration of large functional genomics data. <i>JavaScript (jQuery, d3.js, WebSockets), PHP, MySQL, R/Bioconductor, Python</i></p>	

Epivizr Bioconductor R Package — Aug 2014

github.com/epivizr/epivizr, epivizr.github.io

An R package that provides WebSocket communication to the Epivizr web app for interactive visualization of genomic data. Objects in R/bioc interactive sessions can be displayed in genome browser tracks or plots to be explored by navigation through genomic regions.

Authors: H. C. Bravo, F. Chelaru, L. Smith, N. Goldstein*

IsoCreator (iso-creator-cs.sourceforge.net) — Feb 2007

A .NET app used to create ISO 9660 Joliet CD/DVD images from folders on the local machine.

C#, .NET Framework 2.0

TECHNOLOGIES USED Cloud & Distributed Systems: Kubernetes, Google Cloud Platform (worked on the GCP team).
Frontend: React.js, TypeScript, Xamarin/MAUI.

Backend: Java (J2EE), ASP.NET, Python, Node.js.

Databases: Google Cloud Spanner, Mongo DB, Microsoft SQL Server, MySQL.

LANGUAGES Romanian – native, English – fluent, German – B1 level.

REFERENCES

Dr. Héctor Corrada Bravo

Senior Principal Scientist at Genentech

hcorrada@gmail.com

hcbravo.org

Dr. Mihai Pop

Professor, Department of Computer Science, University of Maryland, College Park

Director, University of Maryland Institute for Advanced Computer Studies (UMIACS)

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pop-lab.org

Dr. Jack van Ryswyck

Senior Research Scientist at Snowflake

[linkedin.com/in/jack-van-ryswyck-532a205](https://www.linkedin.com/in/jack-van-ryswyck-532a205)

Dr. Michael Benisch

VP of Engineering at Woven Planet

[linkedin.com/in/michael-benisch-19055530](https://www.linkedin.com/in/michael-benisch-19055530)