

Mirko LEDDA

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OBJECTIVES To obtain a 3-month industry internship in computational biology as part of the requirements of the Designated Emphasis in Biotechnology graduate program. Have 5 years of industry experience. Result oriented and thrives in large multinational teams and interdisciplinary research environments.

SKILLS

- **Computing:**
 - Programming
 - * Python, R, Matlab (+ some C and Perl).
 - * Software development and release.
 - * Shell scripting and analysis on high-performance clusters.
 - * Common bioinformatics tools/pipelines.
 - Data science
 - * Machine Learning.
 - * Statistics, algebra, calculus and probability theory.
 - * Algorithms and statistical methods development.
 - * Big data management and analysis.
- **Biology:**
 - High-throughput sequencing (incl. library prep).
 - Genomics, transcriptomics, metabolomics and GWAS
 - Receptors biochemistry.
 - Molecular, structural and cell biology.
 - Bioprocesses and bioreactors.
- **Management/Business:**
 - Project management and team building.
 - Effective oral and written communication.
 - IP process.
 - Biological lab management and safety (incl. MP, SOP and GLP).
 - Teaching, consulting and mentoring.

EDUCATION

Ph.D. in Integrative Genetics and Genomics with Emphasis in Biotechnology (DEB*)

University of California at Davis, CA, USA

Expected graduation in Early 2019

Thesis title : Hairpin in a haystack: structure-guided search for functional RNA elements
Thesis advisor : Prof. Sharon Aviran, Biomedical Engineering & Genome Center, UC Davis
DEB advisor : Dr. Judith A. Kjelstrom, director, UC Davis Biotechnology Program

*The [DEB graduate program](#) is an inter-graduate group program involving faculty and students from 29 STEM disciplines. It is focused on providing cross-disciplinary training in critical areas of biomolecular research, promoting interdisciplinary team science, bioethics, entrepreneurship and professionalism as well as coordinating training in a biotechnology or life

B.Sc. in Life Sciences with Emphasis in Biotechnology

2008

University of Applied Sciences (HES-SO), Sion, Switzerland

Topic : Strategies for the study of genes with unknown functions in *Streptomyces*
Advisors : Prof. Sergio Schmid, Institute of Life Technologies, University of Applied Sciences (HES-SO), Sion, Switzerland
Prof. Anna Maria Puglia, Dept. of Biological, Chemical and Pharmaceutical Sciences and Technologies, University of Palermo, Italy

RESEARCH/WORK EXPERIENCE

Ph.D. researcher - UC Davis, CA

2014-Current

Supervisor: Prof. Sharon Aviran

Computational and statistical methods development for nucleic acid structures. My contributions include:

- Characterized the statistical properties of RNA structure profiling data.
- Contributed to, and developed all the statistical bases and strategies for, *RNAprob*, a tool for data-directed RNA secondary structure prediction using probabilistic modeling.
- Conceived and developed *patteRNA*, a Machine Learning algorithm for the rapid mining of RNA secondary structure motifs from transcriptome-wide structure profiling data .
- Mentored new lab members and undergraduate summer students.
- Acting as lab manager and lab safety officer.
- Actively consulted in multiple research projects to tackle complex problems.

Research Assistant - Nestlé Research Center, Lausanne, Switzerland

2009-2014

Supervisor: Prof. Johannes le Coutre

Project and Lab manager for research on taste perception physiology. My contributions include:

- Performed genome-wide association studies (GWAS) of human metabolism and taste perception;
 - Discovered and patented new biomarkers of the health-status of the gastrointestinal tract.
 - Developed statistical methods to analyze human taste phenotypic data.
 - Discovered new genetic drivers of bitter taste perception, most notably for caffeine.
- Established the single cell Ca^{2+} -imaging technique in the lab and developed computational tools for automated image data analysis.
- Functionalized and *in vitro* validated of several human, feline and rat GPCRs/TRP channels in mammalian cells (HEK, HeLa, CHO, Chem-1 and primary rat DRG neurons).
- Performed receptor-interaction studies using siRNAs and co-expression approaches.
- Developed a method for the expression and purification of water-insoluble proteins in *E. coli*.
- Managed research projects and activities. Produced deliverable on time.
- Mentored students and interns.
- Acted as Lab security officer, supervised supplies and ensured proper functioning of lab equipment.

Soldier specialist in biological weapons - Swiss Army, Labor Spiez, Switzerland 2008-2014

Supervisor: Dr. Christian Beuret

- Development and validation of laboratory techniques for rapid identification of pathogenic bacteria, viruses and toxins.
- Team leader for research tasks.

TEACHING EXPERIENCE

- Guest Lecturer** - Quantitative Genetics and Selection Theory (PLS298), UC Davis 2018
IOR: Prof. Steve Knapp
Level: Graduate
Duties: Gave a 1h30 lecture on fundamental concepts in Machine Learning.
Evaluation: Very positive oral feedback from the students.
- Lecturer** - Machine Learning Workshop for the Plant Sciences Dept., UC Davis 2017
IOR: Mirko Ledda
Level: Undergraduate, Graduate and Professors
Duties: 4h workshop on Machine Learning for the Ross-Ibarra, Knapp and Runcie Labs.
Evaluation: Very positive oral feedback. Prof. Ross-Ibarra twitted about it [here](#).
- Guest Lecturer** - Topics in BME: Computational Genomics (BIM189C), UC Davis 2017
IOR: Prof. Sharon Aviran
Level: Upper level undergraduate
Duties: Two 2h lectures on fundamental concepts in Machine Learning.
Evaluation: 4.7/5 - 67% excellent and 33% very good ratings.
- Teaching assistant** - Quantitative Genetics and Selection Theory (PLS298), UC Davis 2016
IOR: Prof. Steve Knapp
Level: Graduate
Duties: Teaching R programming and the mathematical bases of selection theory in lab sessions.
Evaluation: Very positive oral feedback from the students.
- Course development** - Quantitative Genetics and Selection Theory (PLS298), UC Davis 2015
IOR: Prof. Steve Knapp
Level: Graduate
Duties: Preparation of the teaching material for this newly proposed class.

AWARDS

- UC Davis Graduate Student Travel Award** - UC Davis 2017
Competitive award to cover the cost to attend, as a speaker, the *2017 [BC]2 Basel Computational Biology Conference* in Basel, Switzerland.
- Registration Bursary** - Wellcome Genome Campus Scientific Conferences 2016
Competitive award to cover the cost to attend, as a speaker, the *2016 Computational RNA Biology Conference* in Cambridge, UK.
- Summer Graduate Student Researcher Award** - UC Davis 2016
3-months support for graduate research in engineering, computer science, and disciplines with engineering-related applications and methods.

MEMBERSHIP

- Student member** - The RNA Society 2018

COMMUNITY SERVICE

Representative for the IGG graduate program.

Graduate Student Association (GSA) - UC Davis

2015-Current

Member of the organizing committee.

IGG Annual Colloquium - UC Davis

2017

Judge for System and Computational Biology websites.

Teen Biotech Challenge 2017 - DEB, UC Davis

2017

Mentored three students for their final projects.

Topics in Biomedical Engineering: Computational Genomics - UC Davis 2017

Judge for System and Computational Biology websites.

Teen Biotech Challenge 2016 - DEB, UC Davis

2016

Presented biology research and taught basic genetic concepts to three 9th to 12th grade high-school classes.

Science in the Siskiyou - Dunsmuir High School, Dunsmuir, CA, USA

2015

Presented common scientific misconceptions followed by an open discussion with seniors.

Science vs Fiction - Senior Center, Davis, CA, USA

2015

Mentor for all incoming international IGG students and mentor for a 1st year IGG student.

IGG program - UC Davis

2015

PRESENTATIONS AND POSTERS

[BCJ2 Basel Computational Biology Conference - Congress Center, Basel, Switzerland 2017
Ledda M. and Aviran S., patteRNA: Transcriptome-wide search for functional RNA elements via structural data signatures. *Speaker - 20min talk*

Genome Research Day - 23andMe, Mountain View, CA 2017
Ledda M. and Aviran S., Transcriptome-wide search for functional RNA elements via structural data signatures. *Poster*

Computational RNA Biology Conference - Wellcome Trust, Cambridge, UK 2016
Ledda M., Deng F., Vaziri S., and Aviran S., Data-directed RNA secondary structure prediction using probabilistic modeling. *Speaker - 15min talk*

PUBLICATIONS (* INDICATES CO-AUTHORSHIP)

Ledda M. and Aviran S. (2018)
patteRNA: transcriptome-wide search for functional RNA elements via structural data signatures, *Genome Biology* in press

Choudhary K., Shih N.P., Deng F., **Ledda M.**, Li B. and Aviran S. (2016)
Metrics for rapid quality control in RNA structure probing experiments, *Bioinformatics* 32(23): 2575-3583 [\[doi\]](#)

Deng F.*, **Ledda M.***, Vaziri S. and Aviran S. (2016)
Data-directed RNA secondary structure prediction using probabilistic modeling, *RNA* 22(8): 1109-19 [\[doi\]](#)

Michlig González S., Meylan Merlini J., Beaumont M., **Ledda M.**, Tavenard A., Mukherjee R., Camacho S and le Coutre J. (2016)
Acute Effects of single ingestion of TRPV1, TRPA1 and TRPM8 agonists on the energetic metabolism and the autonomic activity in healthy subjects, *Scientific Reports* 6: 20795 [\[doi\]](#)

Rueedi R.*, **Ledda M.***, Nicholls A.W., Salek R.M., Marques-Vidal P., Morya E., Sameshima K., Montoliu I., Da Silva L., Collino S., Martin F-P., Rezzi S., Steinbeck C., Waterworth D.M., Waeber G., Vollenweider P., Beckmann J.S., le Coutre J., Mooser V., Bergmann S., Genick U.K., Kutalik Z. (2014)
Genome-wide association study of metabolic traits reveals novel gene-metabolite-disease links, *PLoS Genetics* 10(2) [\[doi\]](#)

Ledda M.*, Kutalik Z.*, Destito M.C.S., Souza M.M., Cirillo C. a., Zamboni A., Martin N., Morya E., Sameshima K., Beckmann J.S., le Coutre J., Bergmann S., Genick U.K. (2013)
GWAS of human bitter taste perception identifies new loci and reveals additional complexity of bitter taste genetics, *Human Molecular Genetics* 23: 259-267 [\[doi\]](#)

Godinot N., Yasumatsu K., Barcos M.E., Pineau N., **Ledda M.**, Viton F., Ninomiya Y., le Coutre J. and Damak S. (2013)
Activation of tongue-expressed GPR40 and GPR120 by non caloric agonists is not sufficient to drive preference in mice, *Neuroscience* 250: 20-30 [\[doi\]](#)

Montoliu I.*, Genick U.*, **Ledda M.**, Collino S., Martin F.P., Le Coutre J. and Rezzi S. (2013)
Current status on genome-metabolome-wide associations: An opportunity in nutrition research, *Genes and Nutrition* 8: 19-27 [\[doi\]](#)

Genick U.K., Kutalik Z., **Ledda M.**, Souza Destito M.C., Souza M.M., Cirillo C. a., Godinot N., Martin N., Morya E., Sameshima K., Bergmann S., le Coutre J. (2011)
Sensitivity of genome-wide-association signals to phenotyping strategy: The PROP-TAS2R38 taste association as a benchmark, *PLoS One* 6(11) [\[doi\]](#)

PATENTS

Genick U.K., **Ledda M.**, Montoliu I., Le Coutre J., Rezzi S., Collino S., Martin F.P., Da Silva L.
Genetic and urine-derived markers of human metabolic and gut microbial states
European Patent Office *EP2687845 A1* (issued in 2014)
US Patent Office *US Patent 20,150,160,191* (Issued in 2015)

HOBBIES

Soccer, Alpine Ski and GoKart
Hiking, traveling and taking (too) many pictures
Building servers at home

References upon request