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.

RELEVANT SKILLS

Computing Programming • Python, R, Matlab (+ some C and Perl).

· Software development and release.

• Shell scripting and analysis on high-performance clusters.

Data science • Machine Learning.

Statistics, algebra, calculus and probability theory.
Algorithms and statistical methods development.

· Big data management and analysis.

Biology Bioinformatics • High-throughput sequencing (incl. library prep).

• Genomics, transcriptomics, metabolomics and GWAS.

• Common bioinformatics tools/pipelines.

Engineering • Receptors biochemistry.

Molecular, structural and cell biology.

• Bioprocesses and bioreactors.

Business Management • Project management and team building.

• Effective oral and written communication.

· Teaching, consulting and mentoring.

Processes • Intellectual properties.

• Biological lab management.

Safety and quality control (incl. MP, SOP and GLP).

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 science company.

EDUCATION (CONTINUED)

B.Sc. in Life Sciences with Emphasis in Biotechnology

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

2008

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 datadirected 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 Ca²⁺-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.

TEACHING EXPERIENCE (CONTINUED)

Lecturer - Machine Learning Workshop for the Plant Sciences Dept., UC Davis 2017

IOR: Mirko Ledda

Level: Undergraduate, Graduate and Professor

Duties: Organized a 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: Gave 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: Taught 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: Prepared 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-month 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

Graduate Student Association (GS	A) representative - UC Davis	2015-Current
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Representative for the IGG graduate program.

IGG Annual Colloquium organizer - UC Davis 2017

Member of the organizing committee.

DEB volunteer for the Teen Biotech Challenge 2017 - DEB, UC Davis 2017

Judge for System and Computational Biology websites.

Mentor for Topics in BME: Computational Genomics (BIM189C) - UC Davis 2017

Mentored three students for their final projects.

DEB volunteer for the Teen Biotech Challenge 2016 - DEB, UC Davis 2016

Judge for System and Computational Biology websites.

COMMUNITY SERVICE (CONTINUED)

Volunteer for "Science in the Siskiyous" - Dunsmuir High School, Dunsmuir, CA, USA 2015 Presented biology research and taught basic genetic concepts to three 9^{th} to 12^{th} grade high-school classes.

Volunteer for "Science vs Fiction" - Senior Center, Davis, CA, USA

2015

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

Mentor for the IGG program - UC Davis

2015

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

PRESENTATIONS AND POSTERS

[BC]2 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]

PUBLICATIONS (CONTINUED)

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/INTERESTS

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

References upon request