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## RELEVANT SKILLS

<b>Computing</b>	Programming	<ul style="list-style-type: none"><li>• Python, R, Matlab (+ some C and Perl).</li><li>• Software development and release.</li><li>• Shell scripting and analysis on high-performance clusters.</li></ul>
	Data science	<ul style="list-style-type: none"><li>• Machine Learning, statistics, algebra, calculus and probability theory.</li><li>• Algorithms and statistical methods development.</li><li>• Big data management and analysis.</li></ul>
<b>Biology</b>	Bioinformatics	<ul style="list-style-type: none"><li>• High-throughput sequencing (incl. library prep).</li><li>• Genomics, transcriptomics, metabolomics and GWAS.</li><li>• Common bioinformatics tools/pipelines.</li></ul>
	Engineering	<ul style="list-style-type: none"><li>• Receptors biochemistry.</li><li>• Molecular, structural and cell biology.</li><li>• Bioprocesses and bioreactors.</li></ul>
<b>Business</b>	Management	<ul style="list-style-type: none"><li>• Project management and team building.</li><li>• Effective oral and written communication.</li><li>• Teaching, consulting and mentoring.</li></ul>
	Processes	<ul style="list-style-type: none"><li>• Intellectual properties.</li><li>• Biological lab management.</li><li>• Safety and quality control (incl. MP, SOP and GLP).</li></ul>

## EDUCATION

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

University of California at Davis, CA, USA

2014- 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

### **B.Sc. in Life Sciences with Emphasis in Biotechnology**

2004-2008

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

**Research intern** - 23andMe, Mountain View, CA 2018-Current  
Supervisor: Dr. Babak Alipanahi  
Development of machine learning algorithms for finemapping genetic association studies.

**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  $\text{Ca}^{2+}$ -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 Part-time: 3 weeks per year

- 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.

**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.

<b>Guest Lecturer</b> - 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.	

## TEACHING EXPERIENCE (CONTINUED)

<b>Teaching assistant</b> - 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.	
<b>Course development</b> - 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

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

## MEMBERSHIPS / HONORS

<b>Student member</b> - The RNA Society	2018
<b>Nominated for membership</b> - Phi Kappa Phi ( $\Phi\Kappa\Phi$ ) Honor Society	2018
<b>Nominated for membership</b> - Golden Key International Honour Society	2015-2018

## COMMUNITY SERVICE

<b>Graduate Student Association (GSA) representative</b> - UC Davis	2015-Current
Representative for the IGG graduate program.	
<b>IGG Annual Colloquium organizer</b> - UC Davis	2017
Member of the organizing committee.	
<b>DEB volunteer for the Teen Biotech Challenge 2017</b> - DEB, UC Davis	2017
Judge for System and Computational Biology websites.	
<b>Mentor for Topics in BME: Computational Genomics (BIM189C)</b> - UC Davis	2017
Mentored three students for their final projects.	
<b>DEB volunteer for the Teen Biotech Challenge 2016</b> - DEB, UC Davis	2016
Judge for System and Computational Biology websites.	
<b>Volunteer for "Science in the Siskiyous"</b> - Dunsmuir High School, Dunsmuir, CA, USA	2015
Presented biology research and taught basic genetic concepts to three 9 <sup>th</sup> to 12 <sup>th</sup> grade high-school classes.	
<b>Volunteer for "Science vs Fiction"</b> - Senior Center, Davis, CA, USA	2015
Presented common scientific misconceptions followed by an open discussion with seniors.	
<b>Mentor for the IGG program</b> - UC Davis	2015
Mentor for all incoming international IGG students and mentor for a 1 <sup>st</sup> year IGG student.	

## PRESENTATIONS AND POSTERS

- [BC]2 Basel Computational Biology Conference** - Congress Center, Basel, Switzerland 2017  
**Ledda M.** and Aviran S., patterRNA: 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)

- Radecki P.\*, **Ledda M.\*** and Aviran S. (2018) Automated Recognition of RNA Structure Motifs by Their SHAPE Data Signatures, *Genes* 9(6) [doi]
- Ledda M.** and Aviran S. (2018) patterRNA: transcriptome-wide search for functional RNA elements via structural data signatures, *Genome Biology* 19(28) [doi]
- 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.K.\*, **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)

## **LANGUAGES**

English: Fluent

French: Native

Italian: Native

German: Basic

## **HOBBIES/INTERESTS**

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

**References upon request**