Mirko LEDDA

Steven J. Knapp Lab Plant Sciences & Strawberry Breeding Program University of California, Davis, USA

Email: maledda@ucdavis.edu Webpage: https://mirkoledda.github.io

RELEVANT SKILLS

Computing	Programming	 Python, R, Matlab (+ some C, Perl, HTML, CSS and JavaScript). Algorithms, software and web app development and distribution. Code optimization, coverage and profiling analysis.
	Data science	 Statistics, algebra, calculus and probability theory. Machine learning (incl. Tensoflow, Keras and scikit-learn). Big data analysis on HPC (incl. GPUs) and data visualization.
Biology	Bioinformatics	 Next-gen 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.Biology wet-lab management.Safety and quality control (incl. MP, SOP and GLP).
Language		English: Fluent French: Native Italian: Native German: Basic

EDUCATION

Ph.D. in Integrative Genetics and Genomics with Emphasis in Biotechnology (DEB)University of California at Davis, CA, USA

B.Sc. in Life Sciences with Emphasis in Biotechnology

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

RESEARCH EXPERIENCE

Postdoctoral scholar - UC Davis, CA

Supervisor: Prof. Steven J. Knapp

Topic: Computational and statistical methods for genomic prediction, the *in silico* design of genotype markers in complex genomes, and genetic linkage and association studies with complex phenotypes.

Ph.D. researcher - UC Davis, CA

Sep 2014-Apr 2019

Apr 2019-current

Supervisor: Prof. Sharon Aviran

Topic: Computational and statistical methods for the analysis of high-throughput RNA structure probing experiments and RNA secondary structure predictions.

2008

RESEARCH EXPERIENCE (CONTINUED)

Research intern - 23andMe, Moutain View, CA

Jul 2018-Sep 2018

Supervisor: Dr. Babak Alipanahi

Topic: Finemapping genetic association studies using deep learning.

Research Assistant - Nestlé Research Center, Lausanne, Switzerland

Apr 2009-Apr 2014

Supervisor: Prof. Johannes le Coutre

Topic: Genetic bases of taste perception. Taste physiology and receptor pharmacology.

Soldier specialist in biological weapons - Swiss Army, Labor Spiez, Switzerland Sep 2008-Sep 2014 Supervisor: Dr. Christian Beuret (5 months, then part-time 3 weeks per year)

Topic: Lab methods for the rapid identification of pathogenic bacteria, viruses and toxins.

Undergraduate researcher - University of Palermo, Italy

Oct 2007-Apr 2008

Supervisor: Prof. Anna Maria Puglia

Topic: Strategies for the study of genes with unknown functions in *Streptomyces*.

TEACHING EXPERIENCE

Reader - Chemical Engineering Thermodynamics Laboratory, UC Davis 2019

IOR: Prof. Bruce Gates and Prof. Jiandi Wan, Level: Graduate

Duties: Graded laboratory reports.

Teaching assistant - Advanced Genetic Analysis (GGG201A), UC Davis 2018

IOR: Prof. Danika Bannash and Prof. David Segal, Level: Graduate

Duties: Support to student and led a discussion session.

Guest Lecturer - Quantitative Genetics and Selection Theory (PLS298), UC Davis 2018

IOR: Prof. Steve Knapp, Level: Graduate Duties: 1h30 lecture on Machine Learning.

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

IOR: Mirko Ledda, Level: Undergraduate, Graduate and Professor

Duties: 4h workshop on Machine Learning.

Guest Lecturer - Topics in BME: Computational Genomics (BIM189C), UC Davis 2017

IOR: Prof. Sharon Aviran, Level: Upper level undergraduate

Duties: Two 2h lectures on Machine Learning.

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 and breeding theory in lab sessions.

Course development - Quantitative Genetics and Selection Theory (PLS298), UC Davis 2015

IOR: Prof. Steve Knapp, Level: Graduate

Duties: Preparation of the teaching material as it was a new class.

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) patteRNA: 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]

Manuscripts submitted / in-preparation

Ledda M., Cobo N., Knapp S.J. (2019) PolyOligo: A Webapp and Software to Design KASP, CAPS and PCR Genotyping Assays in Polyploid and Complex Genomes, *in preparation for G3: Genes, Genomes, Genetics* [doi]

Picot D.A., **Ledda M.**, Feldmann M.J., Hardigan M.A., Poorten T.J., Heffelfinger C., Cole G.S., Acharya C.B., Dellaporta S., Knapp S.J. (2019) Genealogy Spanning the 300 Year History of Garden Strawberry, *in preparation* [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)

PRESENTATIONS AND POSTERS

2019 ASHS Annual Conference - Tropicana, Las Vegas, NV

2019

Ledda M., Cobo N., Lorant A., Hardigan M.A. and Knapp S.J., PolyOligo: A Bioinformatic Platform for Identifying Target DNA Sequences for the Development of Sub-Genome Specific DNA Markers in Polyploid/Complex Genomes. *Poster*

[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

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

MEMBERSHIPS / HONORS

Graduate Student member - American Society for Horticultural Science (ASHS)	2019
Student member - The RNA Society	2018
Nominated for membership - Phi Kappa Phi ($\Phi K\Phi$) Honor Society	2018
Nominated for membership - Golden Key International Honour Society	2015-2018

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.

COMMUNITY SERVICES

IGG representative for the Graduate Student Association (GSA) - UC Davis 20		
Volunteer for "Skype a Scientist" - AECI Charter High School, Bronx, NY, USA	2019	
eMentor for the Biotechnology Academy Program - Sheldon High School, Sacramento, CA, U	JSA 2019	
IGG Annual Colloquium organizer - UC Davis	2017	
DEB volunteer judge for the Teen Biotech Challenge 2017 - DEB, UC Davis	2017	
Student mentor for Topics in BME: Computational Genomics (BIM189C) - UC Davis	2017	
DEB volunteer judge for the Teen Biotech Challenge 2016 - DEB, UC Davis	2016	
Volunteer for "Science in the Siskiyous" - Dunsmuir High School, Dunsmuir, CA, USA	2015	
Volunteer for "Science vs Fiction" - Senior Center, Davis, CA, USA	2015	
Mentor for incoming international IGG students - UC Davis	2015	

HOBBIES/INTERESTS

Sports (Soccer, Alpine Ski, GoKart), Travels, Hiking, DIY enthusiast.

References upon request