CAMILLO IACOMETTI

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Leiden, Netherlands



EDUCATION

MSc in Cellular & Molecular Biology (international)

University of Turin, Italy Final Grade: 110 /110

Dissertation: "Roo retroelement insertion clusters in D.melanogaster: experimental validation and roles in

alternative transcription"

BSc in Biological Sciences 2015

University of Genoa, Italy Final Grade: 103 /110

Dissertation: "Study of the effect of emerging contaminants on Artemia salina's development: a mosaic

embryo model"

RESEARCH EXPERIENCE

PhD Student 09.2021 – Present

Institute of Biology Leiden (IBL), Leiden University

Supervisor: Lennart Schada von Borzyskowski – Natural and Synthetic Microbial Metabolism

Currently working on genetic tool validation and development for non-model marine Alphaproteobacteria. My project is aimed at harnessing the biotechnological potential of Roseobacter group bacteria for bio-geoengineering applications such as bioremediation of pollutants in marine environments.

Research Assistant 02.2020 – 08.2021

Max-Planck Institute of Molecular Plant Physiology (MPIMP)

Supervisors: Arren Bar-Even & Steffen Lindner - Systems & Synthetic Metabolism group

Rational engineering of *E.coli* central metabolism aimed at uncovering latent metabolic pathways that can replace canonical glycolysis for novel bioproduction routes.

Master Thesis Student 07.2017 – 02.2018

Institute of Evolutionary Biology (IBE: CSIC-UPF), Barcelona Biomedical Research Park (PRBB) Supervisors: Miriam Merenciàno & Josefa González - Functional and Evolutionary Genomics lab

Characterization of *de novo roo* retrotransposon insertions in promoter regions across different strains of *D.melanogaster*. Study of the effects of specific *roo* insertions on fly fecundity and on the alternative transcription of a candidate cold-stress gene.

Bachelor Thesis Student 11.2012 – 01.2014

Hearth, Environment & Life Sciences Department (DISTAV), University of Genoa

Supervisors: Chiara Gambardella & Carla Falugi - Developemental Biology lab

Bachelor thesis project on the evaluation of the effects that emerging contaminants, as organophosphate pesticides, exert on *A.salina* viability and cholinergic system during development, according to the EU regulation on *Registration, Evaluation and Authorization of Chemicals* (REACH).

RELEVANT TECHNICAL SKILLS

Genetic Engineering

<u>E.coli:</u> gene knock-out/in using P1 phage transduction, λ-red recombineering, Direct mutagenesis using Multiplex Automated Genomic Engineering (MAGE), gene overexpression from plasmid and from genome.

Marine Alphaproteobacteria: gene knock-out with classic homologous recombination-based approach, isolation of gDNA, characterization of genomic constitutive promoters' activity by fluorescent reporter measurement, conjugation with *E.coli* ST18.

Molecular Cloning

Construction of plasmids through Gibson or Ligation-mediated assembly, preparation and transformation of chemically-competent (DH5α) and electrically-competent *E.coli* cells (SIJ488), Preparation of competent cells (chem. & electrochem.).

2018

Molecular biology & Microbiology:

DNA extraction from tissues, microbial cultures, PCR mixtures & Gel, quantification by spectrophotometry.

RNA extraction from tissues (D.melanogaster) and microbial cultures (E.coli) & purification, cDNA synthesis, quantification;

Relative quantitation of gene expression by Real-Time qRT-PCR (AB 7900HT Fast Real-Time PCR System), comparative Ct analysis.

Oligonucleotides design for DNA amplification (PCR), for MAGE and for PCR-mediated site-directed mutagenesis, PCR protocols, Gel electrophoresis. Sequence Analysis of DNA products and whole genomic DNA (pairwise alignment, consensus sequence, multiple alignment, identification of mutations & indels).

Solid & Liquid culture media preparation (LB, MX, M9), Growth Experiments in tube and 96-well plates (Biotek & Tecan plate readers)

Biochemistry

Protein expression using *E.coli* BL21, protein extraction and purification using AKTA Start FPLC system, SDS-gel electrophoresis; *In vitro* enzyme kinetic assays with Agilent Cary UV 60 spectrophotometer; enzyme quantification (Bradford protein assay); ¹³C-based metabolic flux analysis, MS data analysis of interested AAs; Immunohystochemical localization of cholinestherases in *A.salina*.

Bioinformatics & IT

Basic Python code, Basic Unix; MatLab: growth plots generation

NGS reads (.fastq) mapping and analysis using <u>breseq</u> pipeline or *Geneious*, Sanger seq. reads mapping and analysis using *Geneious*, GraphPad Prism (enzyme kinetics analysis, growth curve and fluorescence plotting)

Databases (e.g. ENCODE, EcoCyc, Biocyc, Metacycic, NCBI), Office 365 (MS Word, Excel, Power Point), figures editing with Adobe Illustrator

LANGUAGES

ITALIAN - Native speaker

ENGLISH – Fluent (FIRST certificate, Level B2 – University of Cambridge, 2009 + MSc taught in English & international experience)

SPANISH - Fluent

FRENCH - Intermediate

AKNOWLEDGEMENTS

EDISU Scholarship
Region of Piedmont
2015 - 2016
2016 - 2017

Erasmus Plus-Traineeship 2016 - 2017

The first months of my master thesis project were funded through the Erasmus plus traineeship program.

EXTRACURRICULAR ACTIVITIES

Info operator at Science Festival of Genoa

October 2014

Teaching assistant in the *Image Analysis Laboratory* course held by Prof. Giancarlo Panzica (University of Turin). The course aimed at developing basic skills with image manipulation and quantification through *Fiji-ImageJ* software; understanding the characteristics of the digital image, the principles of morphometry and statistical interpretation.

March - June 2017

Private Tutor for students from primary school to bachelor

2016 - 2020

Volunteer for *m'lMporta*, no-profit association promoting environmental awareness and projects with local authorities, mainly in order to cope with the sanitation service deficit in my city area.

2018 – 2020

PUBLICATIONS

lacometti C., Marx K., Hönick M., Biletskaia V., Schulz-Mirbach H., Dronsella B., Satanowski A., Delmas V.A., Berger A., Dubois I., Bouzon M., Döring V., Noor E., Bar-Even A. and **Lindner S.N.** <u>Activating Silent Glycolysis Bypasses in *Escherichia coli*</u>. BioDesign Research, volume 2022, doi: https://doi.org/10.34133/2022/9859643

Merenciano M., **lacometti C**., González J. <u>A unique cluster of roo insertions in the promoter region of a stress response gene in *Drosophila* <u>melanogaster</u>. Mobile DNA, 2019 10:10 doi:10.1186/s13100-019-0152-9</u>

Gambardella C., Nichino D., **lacometti C.**, Ferrando S., Falugi C., Faimali M. <u>Long term exposure to low dose neurotoxic pesticides affects hatching, viability and cholinesterase activity of *Artemia sp.* Aquat Toxicol. 2018 Mar;196:79-89.</u>

doi: 10.1016/j.aguatox.2018.01.006

REFERENTS

Lennart Schada von Borzyskowski (Main Supervisor)

Assistant Professor @ Institute of Biology Leiden

Steffen Lindner (Project Supervisor)

Postdoc @ Charité - Biochemistry and Systems biology of Metabolism

Josefa González Pérez (MSc Thesis Supervisor)

CSIC Tenured Scientist @ Institute of Evolutionay Biology of Barcelona

Chiara Gambardella (BSc Thesis Supervisor)

NRC Researcher, Institute of Marine Sciences (ISMAR)

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- Charité Universitätsmedizin Berlin, Charitéplatz 110117 Berlin
- Pg. Marítim de la Barceloneta 37-49, 08003 Barcelona, Spain
- Via De Marini 6, 16149, Genova, Italy