

Al-powered biochemistry: targeting NAD cofactor essentiality to combat antimicrobial-resistant pathogens

Supervisor: Prof. Leonardo Sorci

Department S.I.M.A.U.

Division of Bioinformatics and Biochemistry,
Faculty of Engineering

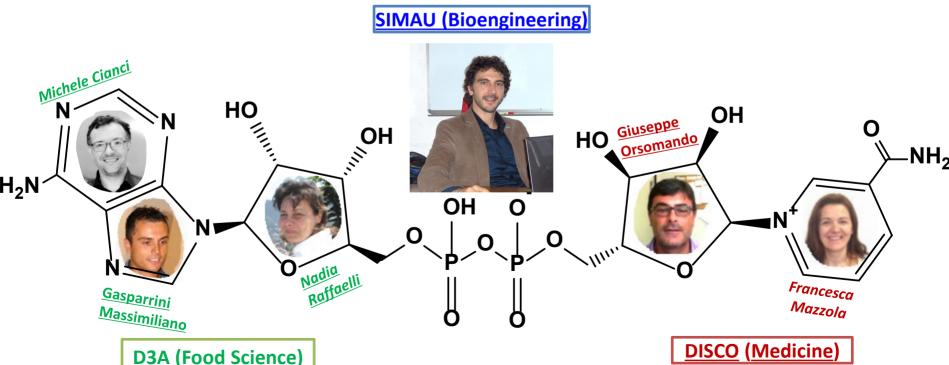
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Research Group Description

Leonardo Sorci, Ph.D. in Biomedical Biotechnologies, Associate Professor of Bioinformatics and Biochemistry (*ORCID*, *CV link*, *publications*, H-index 24), former Staff Scientist at Sanford-Burnham-Prebys medical discovery institute, San Diego, California, joins forces with other colleagues at UNIVPM to bring their complementary skills in the area of Biochemistry (**BIOS-07/A Biochemistry**), sharing their long-term expertise in **NAD cofactor** genomics, metabolism and enzymology.

Interdepartmental Bioinformatics and Biochemistry lab (IBBL)





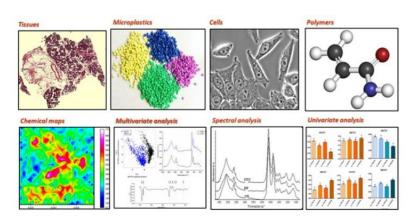
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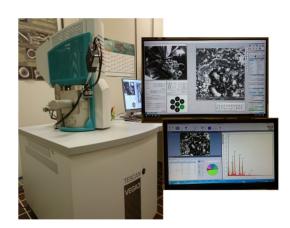
The **Department of Science and Engineering of Materials, Environment and Urban Planning (SIMAU)** is a structure in which the **confluence of different expertises** yield high-level teaching and high-profile international research in the field of **Science of Matter** and **Earth Sciences** with a special focus toward the **Environment**.

It operates within the **Engineering Faculty** offering teachers specialised in the so-called «hard sciences» (**Chemistry** and **Physics**) as well as teachers involved in more «applicative» fields, such as **Materials Engineering**, **Geotechnics**, **Geology**, **Environmental Engineering** and **Urban Planning**.

- TECHNICAL ARCHITECTURE
- BIOINFORMATICS AND BIOCHEMISTRY
- ECOLOGY
- ECONOMICS AND MARKETING
- EXPERIMENTAL PHYSICS
- CHEMICAL FOUNDATION OF TECHNOLOGIES







- APPLIED GEOLOGY AND HYDROGEOLOGY
- ENVIRONMENTAL AND SANITARY CHEMICAL ENGINEERING
- GEOTECHNICAL ENGINEERING
- MATERIALS SCIENCE AND TECHNOLOGY
- URBAN PLANNING TECHNIQUES



Lab skills/expertise: wet-lab approaches relevant to antimicrobial research



GENE CLONING

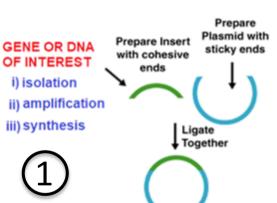
Transform

Bacteria

Plate on

Incubate Plate Overnight

Selective Media





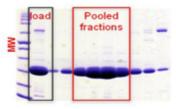
PROTEIN EXPRESSION



PROTEIN PURIFICATION

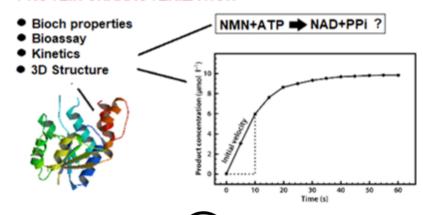


Column Chromatography



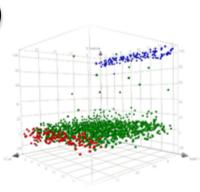


PROTEIN CHARACTERIZATION





- HTS screening
- Cheminformatics artifact filter
- Hit confirmation and selection
- Dose response (IC50, Ki)
- Inhibition mechanism
- Counterscreen





Lab skills/expertise: in silico and Al-powered approaches relevant to antimicrobial research

Gene/protein identification and prediction by sequence (homology) and comparative genomics (gene-fusion, clustering, co-regulation) approaches

Gene ? LPEKRYNHSLRVAETAIKLAEIYDGDTSKVELAGVLHDFCKYDDLGKMS
L EKRY HSL V +TA++LA IY+ DT K +AG++HD K K+

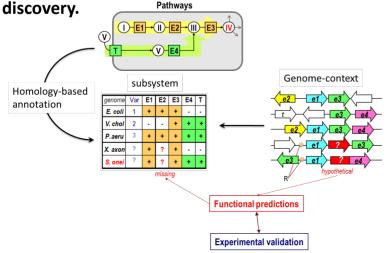
Gene X LGEKRYKHSLGVXDTAVRLAGIYNEDTEKARIAGLVHDCAKKLPGEKII

Organism A ?

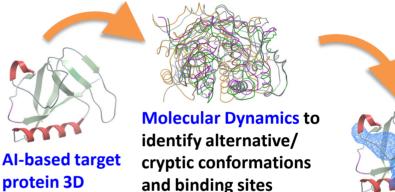
Organism B

Organism C X

Genomics and network-based, including emerging graph-neural networks and trasformer models to identify unexplored target genes for antimicrobial discovery

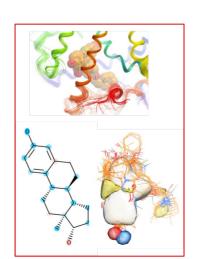


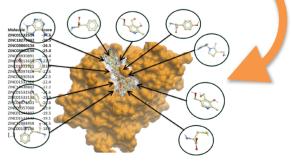
As an integration of, or when experimental 3D structure of the target is unavailable...



protein 3D prediction (includes oligomerization)

Identification of *orthosteric* and *allosteric* binding sites (pocket finder AI-tools)





High quality 2D fingerprint and 3D pharmacophore models used to lead discovery



Project Idea: Al-powered biochemistry: targeting NAD cofactor essentiality to combat antimicrobial-resistant pathogens

Background: NAD cofactor metabolism, which is essential for all pathogens, contains several promising drug targets—specifically, between two and five key enzymes that could be exploited to combat infectious diseases. This approach is especially valuable for addressing bacterial infections, given the growing crisis of multidrug resistance. Since bacteria and other pathogens have not yet evolved defenses against them, new NAD-based antibiotics are expected to remain effective for decades to come.

Objectives: By combining advanced Al-driven biochemistry and drug discovery with our deep expertise in NAD enzymology and structural biology, we aim to develop novel drug candidates targeting pathogens most prone to antibiotic resistance, thus addressing a critical public health challenge. Our target pathogens include, but are not limited to:

🜠 (nadB) De novo (nadA) Downstream (pncB) Salvage Uptake (nadV) (nadM) **NMN** NMNAT NMPRT

Acinetobacter baumannii and other gram-negative enterobacteria, which represent an active and growing threat in Europe due to their high propensity for developing antibiotic resistance. *Mycobacterium tuberculosis*, a re-emerging deadly pathogen driven by the rise of multi-drug-resistant strains. *Plasmodium species (malaria parasites)*, whose potential re-emergence is linked to climate change, increased global travel, and migration patterns. *Schistosoma mansoni*, a parasitic flatworm causing significant disease burden in endemic regions. parasite (flatworm)

Supervisor's curated publications pertinent to the project.

PNAS, 2008

Chemistry & Biology, 2009

Journal of Biological Chemistry 2010

mBIO, 2014

ACS Chemical Biology, 2019

PLOS Pathogens, 2020

Frontiers in microbiology, 2021

Pharmaceuticals, 2024