# **Diego Cotella**

Curriculum vitae

# PERSONAL DATA

Born in Ovada (Italy) in 1974 Living in Novara

# **BIO AND EDUCATION**

| 1987 – 1993 | Secondary School Diploma in Mechanical Engineering, ITIS "C.Barletti", Ovada         |
|-------------|--|
| 1993 – 1999 | Laurea (MSc) in Biology, Università del Piemonte Orientale (UPO) Alessandria         |
| 2000 – 2003 | Doctorate (PhD) in Environmental Sciences, UPO Alessandria                           |
| 1999 – 2002 | R&D staff scientist, Innosense srl, Colleretto Giacosa                               |
| 2004 – 2006 | Postdoctoral researcher, Dresden University of Technology (Germany)                  |
| 2006 - 2010 | Research Associate (Assegnista di Ricerca), UPO Novara                               |
| 2010 – 2011 | Postdoctoral researcher, University of Medicine and Dentistry of New Jersey (UMDNJ), |
|             | Piscataway (USA)   |

## **UNIVERSITY CAREER**

| 2021 to date | Associate Professor in Cell and Experimental Biology (BIOS-10/A), UPO |
|--------------|---|
| 2010 – 2021  | Assistant Professor in Experimental Biology (BIO/13), UPO             |
| 2006 - 2010  | Contract professor, Recombinant DNA Technologies, UPO                 |

## **UNIVERSITY POSITIONS**

| 2023 to date | Thesis Advisory Committee member, PhD Program "Food, health & longevity studies"     |
|--------------|--|
| 2023 to date | Delegate of the International Relations for the School of medicine                   |
| 2015 to date | Member, State exam and Graduation Commission of the Degrees in Physiotherapy,        |
|              | Biotechnology, Medical Biotechnology, Nursing  |
| 2022-2024    | Departmental delegate for the University Orientation Programs                        |
| 2011 - 2018  | Thesis Advisory Committee member, PhD Program "Biotechnologies for the Human Health" |
|              | and Science and Medical Biotechnologies"   |

# TEACHING

| 2025 to date | Biology (pre-Med course)  |
|--------------|---|
| 2021 to date | Synthetic Biology (MSc Medical Biotechnology)                   |
| 2019 to date | Biology (BSc in Nursery)  |
| 2015 to date | Cell Biology (BSc in Biotechnology)                             |
| 2011 to date | Applied Biology (BSc in Healthcare Professions)                 |
| 2006 – 2010  | Laboratory of Recombinant DNA technology (BSc in Biotechnology) |

## THIRD MISSION (ORIENTATION, PUBLIC ENGAGEMENT, ETC.)

| 2022 to date      | University Representative for UniStem   |
|-------------------|---|
| 2020 – 2022, 2025 | Departmental Representative for "European Researchers' Night"                                     |
| 2019 to date      | Teacher, "Scientific Degrees" project. Seminars for high school teachers                          |
| 2016 to date      | Teacher, educational workshops for high schools and "Soft Skills and Orientation Pathways" (PTCO) |
| 2012 – 2024       | Departmental Representative for Orientation   |
| 2011 – 2024       | Teacher, preparation courses for entrance tests in Medicine and Health Professions                |

#### **SCIENTIFIC MEMBERSHIP**

| 2012 to date | Member, the Italian association of Biotechnologists                     |
|--------------|---|
| 2013 to date | Member, the Italian society of Biophysics and Molecular Biology (SIBBM) |
| 2012 to date | Member, the Italian Association of Biology and Genetics (AIBG)          |

#### Awards

| 2019      | National Scientific Habilitation to Associate Professor in Molecular Biology (BIO/11) |
|-----------|---|
| 2016      | National Scientific Habilitation to Associate Professor in Applied Biology (BIO/13)   |
| 2006      | EU Marie European Reintegration Grant (ERG)   |
| 2004-2006 | EU Marie Curie Host Development Program Postdoctoral Fellowship                       |
| 2000-2002 | PhD Fellowship from the Fondazione per le Biotecnologie (Turin)                       |

#### **MAIN FIELDS OF INTEREST**

- 1. Long non-coding RNAs (IncRNAs)
- 2. Biomolecular interactions
- 3. Cell biotechnology
- 4. Phage display
- 5. Microbial engineering and living therapeutics

## **CURRENT RESEARCH FOCUS**

#### 1. The non-coding genome and the Long noncoding RNAs (IncRNAs)

The discovery of long noncoding RNA (IncRNA) represents a significant advance in cell biology. Our goal is to develop new tools to study the interaction between IncRNA and proteins, to understand their function. To this end, we focus on SINEUP, a family of natural antisense (NAT) IncRNAs, whose effect is to promote the translation of specific target mRNAs. Understanding the interaction between proteins and SINEUP could greatly enhance our understanding of the molecular activity of SINEUP.

#### 2. Engineering cell factories for the sustainable production of biopharmaceuticals.

Recombinant proteins are fundamental resources for basic and applied research, as well as for biotechnological applications. They can be produced in a variety of expression systems although mammalian cells are the first choice when post-translational processing is required for function. This project aims at engineering CHO (Chinese Hamster Ovary) and other mammalian cells, to develop novel cell lines capable to produce proteins at elevated level. To this regard, I use my expertise to design, validate and finally apply the SINEUP<sup>™</sup> and other genetic tools to engineer mammalian cell factories to improve the production processes of recombinant proteins.

#### 3. Engineering a new generation of live microbial therapeutics for human disease intervention.

A growing body of research highlights the critical role of microbes in human health and disease, opening new avenues for therapeutic intervention. This project explores the development of engineered live biotherapeutics, non-pathogenic bacteria modified through synthetic biology to detect environmental cues, degrade harmful compounds, and deliver targeted therapies. By addressing key metabolic and immunological pathways implicated in disease, these living medicines offer a promising approach to conditions with significant unmet clinical needs. The research also considers essential design, regulatory, and patient-centred factors for advancing these therapies toward clinical application.

#### **CURRENT FUNDED PROJECTS**

#### **FUNDING PROGRAM**

**2023** – Italian Ministry of University and Research (PRIN 2022 PNRR call)

#### **TITLE OF THE PROJECT**

Decoding dysfunctional RNA transcription and processing by dominant haploinsufficiency in autism spectrum disorders (ASD): rescue by SINEUP translational regulators

**2023** – Italian Autism Foundation (FIA) (2022 call) SINEUP RNAs: a novel RNA molecule platform for the treatment of haploinsufficiency in autism spectrum disorders (ASD).

## **TOP FIVE PAPERS**

1] Morra M, Marradi D, Gandini L, Ivagnes V, Ottolini G, Bovio A, Jabali G, Maraschi L, Ayomide Dada I, Chawanda TV, Gorla M, Tarasiuk O, Mocchetti C, Soluri MF, Boccafoschi F, Sblattero D, **Cotella D** (2025). A non-hypothesisdriven practical laboratory activity on functional metagenomics: "fishing" protein-coding DNA sequences from microbiomes. *Front Bieng Biotech* 13, https://doi.org/10.3389/fbioe.2025.1602982

**2]** Espinoza S, Bon C, Valentini P, Pierattini B, Tettey Matey A, Damiani D, Pulcrano S, Sanges R, Persichetti F, Takahashi H, Carninci P, Santoro C, **Cotella D**, Gustincich S (2021). SINEUPs: a novel toolbox for RNA therapeutics. *Essays Biochem* EBC20200114; <u>https://doi/doi:10.1042/EBC20200114</u>

**3]** Fasolo F, Patrucco L, Volpe M, Bon C, Peano C, Mignone F, Carninci P, Persichetti F, Santoro C, Zucchelli S, Sblattero D, Sanges R, **Cotella D**\*, Gustincich S\*. The RNA-binding protein ILF3 binds to transposable elements sequences in SINEUP lncRNAs. *FASEB J*, 33(12):13572-13589. DOI: 10.1096/fj.201901618RR

**4]** Patrucco L, Peano C, Chiesa A, Guida F, Luisi I, Boria I, Mignone F, De Bellis G, Zucchelli S, Gustincich S, Santoro C, Sblattero D, **Cotella D**. (2015) Identification of novel proteins binding the AU-rich element of  $\alpha$ -prothymosin mRNA through the selection of open reading frames (RIDome). RNA Biol. 12(12):1289-300. doi: 10.1080/15476286.2015.1107702.

**5]** Patrucco L, Peano C, Chiesa A, Guida F, Luisi I, Boria I, Mignone F, De Bellis G, Zucchelli S, Gustincich S, Santoro C, Sblattero D, **Cotella D** (2015). Identification of novel proteins binding the AU-rich element of  $\alpha$ -prothymosin mRNA through the selection of open reading frames (RIDome). *RNA Biol.* 12(12):1289-300. doi: 10.1080/15476286.2015.1107702.