



# LORENZO MAEDDU

## MACHINE LEARNING SCIENTIST

### PERSONAL PROFILE

I am a machine learning scientist with experience in teaching, research, and multidisciplinary projects.

### WORK EXPERIENCE



#### Senior Data Scientist (Knowledge Graph Insights)

AstraZeneca - Göteborg, Sweden

Sep 2022 - Current Position

- Developed graph-mining frameworks for large biomedical knowledge graphs to investigate novel pharmaceutical insights



#### Machine Learning Scientist (PhD experience - Harvard)

Harvard Medical School, Brigham and Women's Hospital, Loscalzo Lab

Jun 2021 - Sept 2021

- Tested 6000+ drugs with network pharmacology and machine learning strategies to uncover effective treatments for the COVID-19 infection



#### Protein-Protein Interactions Prediction Challenge

Institute for Network Medicine Foundation

Nov 2020 - Sept 2021

- Worked with an international team of academics from Harvard University and Sapienza University of Rome, to develop a competition for predicting new protein-protein interactions



#### Machine Learning Scientist in the AI task-force for COVID-19

Confederation of Laboratories for Artificial Intelligence Research in Europe May 2020 - Dec 2020

- Released a public bioinformatic dataset on COVID-19 for drug repurposing tasks ([link](#))
- Developed a drug repurposing deep learning model for COVID-19 (submitted paper)



#### Data Scientist

Sapienza University of Rome

Sep 2017 - Dec 2017

- Developed a Java-based framework for extraction and analysis of social temporal web data ([link](#))

### TEACHING



#### Machine Learning Teacher

UnitelmaSapienza University of Rome

Jun 2022 - Aug 2022

- Recorded 17 Machine Learning lessons



#### Data Science Teacher

LUISS Guido Carli university

Feb 2022 - May 2022

- Teaching assistant for the "Data Visualization" course at the LUISS Guido Carli university.



#### Machine Learning Scientist

Sapienza University of Rome

Dec 2020 - Feb 2021

- Wrote the teaching material for a course of Machine Learning ([link](#))



#### Computer Science Teacher

Sapienza University of Rome

Feb 2019 - May 2019

- Teaching assistant for Python at the "Fundamentals of Programming" course ([link](#))

### EDUCATION



#### PhD in Innovative Biomedical Technologies in Clinical Medicine

Sapienza University of Rome

Nov 2018 - Jun 2022

- Research Fields: Machine Learning, Graph-Mining, Data-Mining
- Areas of Application: Network Biology, Network Pharmacology



#### MSc, Computer Science (in English)

Sapienza University of Rome

Oct 2016 - Oct 2018

- Grade: 110/110, with honors (GPA 4.00)
- Machine Learning, Deep Learning and Big Data analytics



#### BSc, Computer Science (in Italian)

Sapienza University of Rome

Sep 2013 - Oct 2016

- Grade: 108/110 (GPA 4.00)
- Thesis in network analysis using Big Data technologies (Hadoop, MapReduce, Java)

### CONTACT ME AT

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### SOFT SKILLS

Empathy

Enthusiasm

Problem Solving

Networking

Teamwork

### HARD SKILLS

Italian, native proficiency

English, working proficiency

Multidisciplinary capabilities

Research

Machine learning

### PROGRAMMING SKILLS

Python, Java, SQL, Numpy,

Pandas, Sklearn, Keras, Pytorch,

Data structures, NLP, Networkx

### CERTIFICATIONS

Neural Networks and Deep Learning ([link](#))

### AWARDS

Winner of the "Enabling Sustainable Cities Through Blockchain Challenge" (2019, WomENCourage)

## JOURNAL ARTICLES

### Assessment of community efforts to advance network-based computational prediction of protein-protein interactions

Xu-Wen Wang, Lorenzo Madeddu et al.

Submitted - 2022|

- <https://doi.org/10.1101/2021.09.22.461292>

### Deep Graph Networks for Drug Repurposing with Multi-Protein Targets

Davide Bacciu, Federico Errica, Alessio Gravina, Lorenzo Madeddu, Marco Podda, Giovanni Stilo

Submitted - 2022|

### A Network-Based Analysis of Disease Modules from a Taxonomic Perspective

Lorenzo Madeddu, Giorgio Grani, Paola Velardi

JBHI - 2021|

- <https://doi.org/10.1109/JBHI.2021.3106787>

### A Feature-Learning-Based Method for the Disease-Gene Prediction Problem

Lorenzo Madeddu, Giovanni Stilo, Paola Velardi

IJDMB -

- <https://doi.org/10.1504/IJDMB.2020.109502>

## CONFERENCE ARTICLES

### Integrating Categorical and Structural Proximity in Disease Ontologies

Lorenzo Madeddu, Giorgio Grani, Paola Velardi

EMBC - 2021|

- <https://doi.org/10.1109/EMBC46164.2021.9630114>

### Challenges and Solutions to the Student Dropout Prediction Problem in Online Courses

Bardh Prenkaj, Giovanni Stilo, Lorenzo Madeddu

CIKM - 2020|

- <https://doi.org/10.1145/3340531.3412172>

### Predicting disease genes for complex diseases using random watcher-walker

Lorenzo Madeddu, Giovanni Stilo, Paola Velardi

SAC - 2020|

- <https://doi.org/10.1145/3341105.3373979>

### Predicting Disease Genes Using Connectivity and Functional Features

Lorenzo Madeddu, Giovanni Stilo, Paola Velardi

BIBM - 2019|

- <https://doi.org/10.1109/BIBM47256.2019.8982929>

## BOOK CHAPTERS

### Deep Learning Methods in Network Biology

Lorenzo Madeddu, Giovanni Stilo

World Scientific - 2021|

- <https://doi.org/10.1142/q0322>

### Ontological and Connectivity Structure of Disease-Gene Modules in the Human Interactome

Lorenzo Madeddu, Giovanni Stilo

Springer Nature - 2021|

- [https://doi.org/10.1007/978-3-030-58080-3\\_76-1](https://doi.org/10.1007/978-3-030-58080-3_76-1)

## FEATURED TALKS

### AI for Health and Medicine Intelligent Information Mining - Research Group

OpenDI

Mar 2022|

- A presentation of the current research projects of the lab at the Computer Science department.

### AI for Health and Medicine Intelligent Information Mining - Research Group

Ital-IA

Feb 2022|

- A presentation of the current research projects of the lab.

### Deep Learning in Network Biology

Dido Workshop

Nov 2021|

- A comprehensive overview of fundamental concepts, deep learning methods and critical challenges of Network Biology in Network Biology, Network Medicine, and Network Pharmacology.