

Curriculum vitae

Assistant Professor

Department of Computing Sciences, Bocconi University

Tangherloni Andrea

07/02/1990

Via Guglielmo Röntgen 1, 20136, Milan, Italy

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Work experience

Assistant Professor

SEPTEMBER 2023 - CURRENT

Department of Computing Sciences, Bocconi University, Italy

Bocconi Institute for Data Science and Analytics, Bocconi University, Italy

Postdoctoral Researcher (paid on ERC 772970 - MicroC)

DECEMBER 2022 - AUGUST 2023

Bocconi Institute for Data Science and Analytics, Bocconi University, Italy

Department of Computing Sciences, Bocconi University, Italy

Postdoctoral Researcher

DECEMBER 2020 - NOVEMBER 2022

Department of Human and Social Sciences, University of Bergamo, Italy

Research Associate (paid on ERC 677501 - ZF_Blood)

NOVEMBER 2018 - NOVEMBER 2020

Department of Haematology, University of Cambridge, Cambridge, UK

Wellcome - MRC Stem Cell Institute, Cambridge, UK

Visiting worker at Wellcome Trust Sanger Institute, Hinxton, UK

Education and training

PhD in Computer Science

NOVEMBER 2015 - FEBRUARY 2019

Department of Informatics, Systems and Communication, University of Milano-Bicocca, Italy

Dissertation title: *"High-Performance Computing to tackle complex problems in life sciences"*. Supervisors: Prof. Daniela Besozzi and Prof. Paolo Cazzaniga

Master of Science in Computer Science

NOVEMBER 2013 - OCTOBER 2015 (110/110 summa cum laude)

Department of Informatics, Systems and Communication, University of Milano-Bicocca, Italy

Dissertation title: *"Design and development of a large-scale biochemical simulator accelerated on GPUs"*. Supervisor: Prof. Giancarlo Mauri

Bachelor of Science in Computer Science

OCTOBER 2010 - OCTOBER 2013 (107/110)

Department of Informatics, Systems and Communication, University of Milano-Bicocca, Italy

Dissertation title: *"Development of a food database for feature-based classification and retrieval"*. Supervisor: Prof. Raimondo Schettini

Education/training abroad

Erasmus+ Traineeship Program (A.Y. 2017/18, 05/04/2018 – 05/07/2018) with the highest evaluation (90/90): visiting PhD student at the Department of Computer Science and Technology (The Computer Laboratory), University of Cambridge, Cambridge, UK

ExtraUE Exchange Program (A.A. 2016/17, 06/06/2017 – 06/08/2017) with the highest evaluation (100/100): visiting PhD student at the Department of Cancer Biology, School of Medicine, Vanderbilt University, Nashville, TN, USA

Research interests

Artificial Intelligence and Machine Learning
Computational Intelligence
High-Performance Computing
Computational Biology and Systems Biology
Bioinformatics
Mathematical modelling and simulation of complex biological systems
Biomedical image analysis

Fellowships, awards, and prizes

Bocconi Junior Grant (2025): € 10.000

Bocconi Junior Grant (2024): € 10.000

Second place at the Start Cup Bergamo 2021 (Start Cup School and Competition) with the project “ReHubAI: il rimedio a portata di click!”, Department of Management Information and Production Engineering, and Center for Young and Family Enterprise (CYFE), University of Bergamo, Italy

PhD Scholarship (2015 - 2018) founded by the Italian Ministry of Education, University of Milano-Bicocca, Italy

Fellowship for the Erasmus+ Traineeship Program (2018): visiting PhD student at the Department of Computer Science and Technology, University of Cambridge, Cambridge, UK (3 months)

Fellowship for the ExtraUE Exchange Program (2017): visiting PhD student at the Department of Cancer Biology, School of Medicine, Vanderbilt University, Nashville, TN, USA (2 months)

Teaching activity

First and second level master's degrees

Lectures (4 hours, A.Y. 2022/23): Neural networks and deep learning, Second Level Master's Degree in Methods and Data Analysis in Biomedical Research (MEDAL), University of Milano-Bicocca, Italy

Master's degrees

Lecturer (20 hours, A.Y. 2024/25): Bioinformatics, Master's Degree in Artificial Intelligence, Bocconi University, Italy

Lecturer (32 hours, A.Y. 2024/25): Deep Learning and Reinforcement Learning, Master's Degree in Artificial Intelligence, Bocconi University, Italy

Lecturer (32 hours, A.Y. 2023/24): Deep Learning and Reinforcement Learning, Master's Degree in Artificial Intelligence, Bocconi University, Italy

Laboratory lectures (16 hours, A.Y. 2021/22): ICT Systems Management,

Master's Degree in Management Engineering, University of Bergamo, Italy

Laboratory tutor (30 hours, A.Y. 2017/18): Data Technology and Machine Learning, Master's Degree in Computer Science, University of Milano-Bicocca, Italy

Bachelor's degrees

Lecturer (8 hours, A.Y. 2024/25): Machine Learning, Bachelor's Degree in Economics, Management and Computer Science, Bocconi University, Italy

Lecturer (30 hours, A.Y. 2024/25): Machine Learning and Artificial Intelligence, Bachelor's Degree in Mathematical and Computing Sciences for Artificial Intelligence, Bocconi University, Italy

Lecturer (30 hours, A.Y. 2023/24): Machine Learning and Artificial Intelligence, Bachelor's Degree in Mathematical and Computing Sciences for Artificial Intelligence, Bocconi University, Italy

Lecturer (40 hours, A.Y. 2023/24): Computer Science, Bachelor's Degrees, Bocconi University, Italy

Teaching assistant (30 hours, A.Y. 2022/23): Topics in Computational Modelling: from Information Theory to Evolutionary Models, Bachelor's Degree in Mathematical and Computing Sciences for Artificial Intelligence, Bocconi University, Italy

Laboratory lectures (6 hours, A.Y. 2022/23): Fundamentals of informatics and statistics, Bachelor's Degree in Psychological Studies, University of Bergamo

Laboratory tutor (30 hours, A.Y. 2021/22): Computer Science, Bachelor's Degree in Biotechnology, University of Milano-Bicocca, Italy

Laboratory lectures (6 hours, A.Y. 2020/21): Fundamentals of informatics and statistics, Bachelor's Degree in Psychological Studies, University of Bergamo

Laboratory lectures (24 hours, A.Y. 2016/17): Computer Architecture, Bachelor's Degree in Computer Science, University of Milano-Bicocca, Italy

E-learning tutor (30 hours, A.Y. 2014/15): Operating Systems and Computer Networks, Bachelor's Degree in Computer Science, University of Milano-Bicocca, Italy

Other

Masterclass at "VisionnAlres", Bocconi University, Milan, 2025. Title of the talk: *"Machine Learning e biologia computazionale: come l'AI sta cambiando la ricerca biomedica"*

AI Instructor at the Summer School "Recent Trends In Artificial Intelligence" (2025), Bocconi University

AI Instructor at the Summer School for High School Students (2024 and 2025), Bocconi University

Lessons and laboratory (June 21 - 30, 2023): Transcriptomics Advanced Course, EMBL-EBI, Wellcome Genome Campus, Hinxton, UK

Lessons and laboratory (24 hours May 30 - 31 and June 1, 2017): Python for Beginners, Micron Semiconductor Italia S.R.L., Milan, Italy

Supervision of Students and Post-Doctoral Fellows

Research Assistants

2023-current: Supervision of four Research Assistants in Computer Science, Department of Computing Sciences, Bocconi University, Italy

First level master's degrees

2021: co-supervision of a first-level Master's thesis (with Dr Marabini Leonardo)

Bachelor's and Master's students

2023-current:

- Supervision of six Master's theses in Computer Science, Department of Computing Sciences, Bocconi University, Italy
- Supervision of ten Bachelor's theses in Computer Science, Department of Computing Sciences, Bocconi University, Italy

2019: Co-supervision of two Master's theses in Computer Science, University of Milano-Bicocca; University of Cambridge, UK

2017-2018: Co-supervision of three Bachelor's theses in Computer Science, University of Milano-Bicocca, Italy

Conferences and talks

Conferences

Speaker at the IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology, Tainan, Taiwan 2025. Title of the talk: *"GENESIS: Generating scRNA-Seq data from Multiome Gene Expression"*

Speaker at the IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology, Tainan, Taiwan 2025. Title of the talk: *"Knowledge-Enriched Cell-Type Annotation in Single-Cell Transcriptomics via LLM Embeddings"*

Speaker at the IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology, Natal, Brazil 2024. Title of the talk: *"A Fast Feature Selection for Interpretable Modeling Based on Fuzzy Inference Systems"*

Speaker at the IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology, Natal, Brazil 2024. Title of the talk: *"Forest-based Evolutionary Algorithm for Reconstructing Boolean Gene Regulatory Networks"*

Speaker at the IEEE Congress on Evolutionary Computation, Yokohama, 2024. Title of the talk: *"A modified EACOP implementation for Real-Parameter Single Objective Optimization Problems"*

Speaker at the IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology, The Netherlands, 2023. Title of the talk: *"Consensus clustering strategy for cell type assignments of scRNA-seq data"*

Speaker at the IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology, Canada, 2022. Title of the talk: *"Multi-objective Optimization for Marker Panel Identification in Single-cell Data"*

Speaker at the IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology, Canada, 2022. Title of the talk: *"A Deep Learning Pipeline for the Automatic cell type Assignment of scRNA-seq Data"*

Speaker at the IEEE International Conference on Bioinformatics and Biomedicine, Virtual, 2021. Title of the talk: *"Integration of Multiple scRNA-Seq Datasets on the Autoencoder Latent Space"*

Speaker at the 17th International Conference on Computational Intelligence Methods For Bioinformatics And Biostatistics, Virtual, 2021. Title of the talk: *"Genetic Algorithms for the identification of marker panels in single-cell RNA Data"*

Speaker at the IEEE Congress on Evolutionary Computation, Virtual, 2021. Title of the talk: *"The Impact of Representation on the Optimization of Marker Panels for Single-cell RNA Data"*

Speaker at the 24th International European Conference on Parallel and Distributed Computing, Turin, Italy, 2018. Title of the talk: *"High Performance Computing for Haplotyping: Models and Platforms"*

Speaker at the 17th International Workshop on Network Tools and Applications in Biology, Palermo, Italy, 2017. Title of the talk: *“GenHap: A Novel Computational Method Based on Genetic Algorithms for Haplotype Assembly”*

Speaker at the 12th International Meeting on Computational Intelligence Methods For Bioinformatics And Biostatistics, Naples, Italy, 2015. Title of the talk: *“Deterministic Simulations of Large-Scale Models of Cellular Processes Accelerated on Graphics Processing Units”*

Seminars

Invited speaker at “Multiomics in Personalized Medicine Workshop”, TÜSEB Aziz Sancar Research Centre, Ankara, 2025. Title of the talk: *“AI Meets Biology: Can AI Help Us Analyse Multi-Omics Data?”*

Invited speaker at “Get a Taste of Bocconi”, Bocconi University, Milan, 2024. Title of the talk: *“AI Evolution: From Leonardo to ChatGPT”*

Invited speaker at “Orientamatica (PRISTEM, “Modelli matematici nella realtà”), Bocconi University, Milan, 2024. Title of the talk: *“Introduction to Artificial Intelligence”*

Invited speaker at the Milan meeting on “Next Generation Sequencing”, University of Milano-Bicocca, Italy, 2022. Title of the talk: *“Automatic cell-type annotation of scRNA-seq data: an overview”*

Invited speaker at the Internal Seminar Series, Wellcome - MRC Stem Cell Institute, Cambridge, UK, 2019. Title of the talk: *“Advanced Machine Learning approaches for the downstream analysis of scRNA-Seq data”*

Invited speaker at the Department of Computer Science and Technology, University of Cambridge, Cambridge, UK, 2018. Title of the talk: *“High-Performance Computing Applied to Computational Intelligence in Systems Biology and Genome Analysis”*

Invited speaker at the 1st Computational Intelligence and Biomathematics, Rio de Janeiro, Brasile, 2017. Title of the webinar: *“Biomedical Image Segmentation and Analysis Using Machine Learning and Computational Intelligence Techniques”* (co-author Dr. Leonardo Rundo)

Invited speaker at the Department of Cancer Biology, Vanderbilt University, Nashville, TN, USA, 2017. Title of the talk: *“High Performance Computing to Tackle Complex Problems in Life Sciences”*

Program Committee and Organization Committee

Conferences

Member of the Organizing Committee of the IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology, 2027

Member of the Organizing Committee of the IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology, 2023

Member of the Organizing Committee of the IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology, 2023

Member of the Program Committee of the 14th International Conference on Bioinformatics, Biocomputational Systems and Biotechnologies, 2022

Member of the Program Committee of the 8th International Conference on Machine Learning, Optimization, and Data Science, 2022

Member of the Program Committee of the 7th International Conference on Machine Learning, Optimization, and Data Science, 2021

Member of the Program Committee of the 6th International Conference on Machine Learning, Optimization, and Data Science, 2020

Special Sessions and workshops

Co-chair of the Special Session “Deep Learning in Computational Biology and Biomedicine: from Biomedical Data to Drug Discovery”, IEEE International Joint Conference on Neural Networks, 2025

Co-chair of the Special Session “*Candidate Solution Representation and Fitness Landscape Manipulation*”, IEEE Congress on Evolutionary Computation, 2023

Co-chair of the Special Session “*Candidate solution representation and fitness landscape manipulation*”, IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology, 2022

Member of the Program Committee of the Special Session “*Candidate solutions representation and fitness landscape manipulation*”, IEEE Congress on Evolutionary Computation, 2021

Co-chair of the Workshop “*Machine Learning and Computational Intelligence in multi-omics and medical image analysis*”, 16th International Conference on Artificial Intelligence Applications and Innovations, 2020,

Local Organizer of the International Workshop on Bio-Design Automation, 2019

Co-chair of the Special Session “*Machine Learning and Computational Intelligence in multi-omics and medical image analysis*”, 16th International Conference on Computational Intelligence methods for Bioinformatics and Biostatistics, 2019

Member of the Program Committee of the post-proceedings of the 16th International Conference on Computational Intelligence methods for Bioinformatics and Biostatistics, 2019, LNCS

Member of the Program Committee of the Special Session “*Fast and Efficient Solutions for Computational Intelligence Methods in Bioinformatics, Systems and Computational Biology*”, 15th International Conference on Computational Intelligence methods for Bioinformatics and Biostatistics, 2018

Member of the Program Committee of the Special Session “*Modeling and Simulation Methods for System Biology and System Medicine*”, 15th International Conference on Computational Intelligence methods for Bioinformatics and Biostatistics, 2018

Member of the Program Committee of the Special Session “*Advances in High-Performance Bioinformatics and Biomedicine*”, 26th Euromicro International Conference on Parallel, Distributed, and Network-Based Processing, 2018

Member of the Program Committee of the Special Session “*Advances in High-Performance Bioinformatics, Systems and Synthetic Biology*”, 25th Euromicro International Conference on Parallel, Distributed, and Network-Based Processing, 2017

Special Session “*Parallel and distributed high performance computing solutions for computational intelligence methods*”, IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology, 2016

Referee activity

Peer-Reviewed journals

Mathematical Biosciences and Engineering (AIMS Press); Applied Soft Computing (Elsevier); Artificial Intelligence in Medicine (Elsevier); Biosafety and Health (Elsevier); Computational and Structural Biotechnology Journal (Elsevier); Computer Methods and Programs in Biomedicine (Elsevier); Computers in Biology and Medicine (Elsevier); Expert Systems With Applications (Elsevier); Journal of Industrial Information Integration (Elsevier); Knowledge-based Systems (Elsevier); Neurocomputing (Elsevier); Parallel Computing Systems &

Applications (Elsevier); Frontiers in Bioengineering and Biotechnology (Frontiers); Frontiers in Cell and Developmental Biology (Frontiers); Frontiers in Genetics (Frontiers); Frontiers in Immunology (Frontiers); Frontiers in Neurology (Frontiers); Frontiers in Oncology (Frontiers); Frontiers in Plant Science (Frontiers); Computational and Mathematical Methods in Medicine (Hindawi); IEEE Access (IEEE); IEEE Computational Intelligence Magazine (IEEE); IEEE Transactions on Artificial Intelligence (IEEE); Fundamenta Informaticae (IOS Press); Concurrency and Computation: Practice and Experience (John Wiley & Sons); Algorithms (MDPI); Applied Sciences (MDPI); BioChem (MDPI); Biomolecules (MDPI); Cancers (MDPI); Diagnostics (MDPI); Electronics (MDPI); Genes (MDPI); Journal of Imaging (MDPI); Mathematics (MDPI); Sensors (MDPI); Signals (MDPI); Sustainability (MDPI); Symmetry (MDPI); Bioinformatics (Oxford Academic); Nucleic Acids Research (Oxford Academic); PLoS Computational Biology (PLOS); BMC Bioinformatics (Springer Nature); BMC Medical Informatics and Decision Making (Springer Nature); Communications Biology (Springer Nature); International Journal of Machine Learning and Cybernetics (Springer Nature); Nature Computational Science (Springer Nature); Nature Methods (Springer Nature); The Journal of Supercomputing (Springer Nature); Engineering Optimization (Taylor & Francis); Smart Science (Taylor & Francis)

International conferences

Euromicro International Conference on Parallel, Distributed, and Network-Based Processing; IEEE Congress on Evolutionary Computation; IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology; International Conference on Bioinformatics, Biocomputational Systems and Biotechnologies; International Conference on Computational Intelligence methods for Bioinformatics and Biostatistics; International Conference on Machine Learning, Optimization, and Data Science

PhD students

Reviewer for a PhD thesis at the University of Eastern Finland and Sapienza University of Rome

Transfer of Status Assessor for a PhD student at the University of Oxford

Editorial activities

Editor in Chief

River Series “Computational Approaches in Biology and Medicine” (River Publishers)

Associate Editor

2022 - Current: PLoS ONE, Public Library of Science

2022 - Current: BMC Bioinformatics, Springer-Nature

2022 - 2025: Heliyon (Computer Science Section), Cell Press

Review Editor

“*Computational Genomics*”, Frontiers in Genetics, Frontiers in Bioengineering and Biotechnology, and Frontiers in Plant Science

Topic Editor

“*Biomedical Engineering*”, Applied Sciences (MDPI)

Guest Editor

“*Artificial Intelligence Techniques for Medical Imaging and Computational Biology*”, Special Issue, Applied Sciences (MDPI)

"Artificial Intelligence Approaches for the Analysis and Integration of Omics Data",
Research Topic in Integrative Bioinformatics, Frontiers in Bioinformatics
"Machine Learning Applied to Medical Imaging and Computational Biology", Special
Issue, Applied Sciences (MDPI)

Ancillary activities

2024-2025: Moderator of the department website Department of Computing Sciences, Bocconi University, Italy
2024-Current: Co-chair of the IEEE CIS Task Force on Advanced Representation in Biological and Medical Search and Optimization
2023-Current: Member of the IEEE Bioinformatics and Bioengineering Technical Committee (BBTC) Members
2022-Current: Member of the IEEE CIS Task Force on Advanced Representation in Biological and Medical Search and Optimization
2020: Member of the Stem Cell Institute Public Engagement network, Wellcome – MRC Stem Cell Institute, Cambridge, UK
2015-2018: PhD Representative, Department of Informatics, Systems and Communication, University of Milano-Bicocca

Other information

Author/co-author of more than 40 publications in international peer-reviewed journals, international conference proceedings, and book chapters.
Google Scholar: 1817 citations; h-index = 18
Scopus: 1293 citations; h-index = 16

Languages

Italian: mother tongue
English: excellent

Date

03/10/2025

Signature

A handwritten signature in black ink, written over a horizontal line. The signature is stylized and appears to be 'F. A. M.' followed by a flourish.

Publications

Journals

1. **Tangherloni A†**, Riva SG, Myers B, Buffa FM, Cazzaniga P (2023). *MAGNETO: Marker pAnels GeNEraTor with multi-Objective optimization*. **Journal of Biomedical Informatics**, 147:104510. DOI: 10.1016/j.jbi.2023.104510
2. Papetti DM, **Tangherloni A**, Farinati D, Cazzaniga P, Vanneschi L (2023). *Simplifying Fitness Landscapes using Dilation Functions evolved with Genetic Programming*. **IEEE Computational Intelligence Magazine**, 18(1): 22-31. DOI: 10.1109/MCI.2022.3222096
3. Riva SG, Cazzaniga P, Nobile MS, Spolaor S, Rundo L, Besozzi D, **Tangherloni A†** (2022). *SMGen: A generator of synthetic models of biochemical reaction networks*. **Symmetry**, 14(1): 119. DOI: 10.3390/sym14010119
4. Castelli M, Manzoni L, Mariot L, Nobile MS, **Tangherloni A** (2022). *Salp Swarm Optimization: a Critical Review*. **Expert Systems with Applications**, 189: 116029. DOI: 10.1016/j.eswa.2021.116029
5. **Tangherloni A†**, Nobile MS, Cazzaniga P, Capitoli G, Spolaor S, Rundo L, Mauri G, Besozzi D (2021). *FiCoS: a fine- and coarse-grained GPU-powered deterministic simulator for biochemical networks*. **PLoS Computational Biology**, 17(9): e1009410. DOI: 10.1371/journal.pcbi.1009410
6. **Tangherloni A†**, Ricciuti F, Besozzi D, Liò P, Cvejic A (2021). *Analysis of single-cell RNA sequencing data based on autoencoders*. **BMC Bioinformatics**, 22(1): 309. DOI: 10.1186/s12859-021-04150-3
7. Ranzoni AM*, **Tangherloni A***, Berest I*, Riva SG, Myers B, Strzelecka PM, Xu J, Panada E, Mohorianu I, Zaugg JB, Cvejic A (2021). *Integrative Single-Cell RNA-Seq and ATAC-Seq Analysis of Human Developmental Hematopoiesis*. **Cell Stem Cell**, 28(3): 472-487. DOI: 10.1016/j.stem.2020.11.015
8. Rundo L*, **Tangherloni A***, Cazzaniga P, Mistri M, Galimberti S, Woitek R, Sala E, Mauri G, Nobile MS (2021). *A CUDA-powered method for the feature extraction and unsupervised analysis of medical images*. **The Journal of Supercomputing**. DOI: 10.1007/s11227-020-03565-8
9. Rundo L*, **Tangherloni A***, Tyson DR*, Betta R, Militello C, Spolaor S, Nobile MS, Besozzi D, Lubbock ALR, Quaranta V, Mauri G, Lopez CF, Cazzaniga P (2020). *ACDC: Automated cell detection and counting for time-lapse fluorescence microscopy*. **Applied Sciences**, 10(18), 6187. DOI: 10.3390/app10186187
10. Besozzi D, Manzoni L, Nobile MS, Spolaor S, Castelli M, Vanneschi L, Cazzaniga P, Ruberto S, Rundo L, **Tangherloni A** (2020). *Computational Intelligence for Life Sciences*. **Fundamenta Informaticae**, 171(1-4), 57-80. DOI: 10.3233/FI-2020-1872
11. **Tangherloni A**, Spolaor S, Cazzaniga P, Besozzi D, Rundo L, Mauri G, Nobile MS (2019). *Biochemical parameter estimation vs. benchmark functions: A comparative study of optimization performance and representation design*. **Applied Soft Computing**, 81: 105494. DOI: 10.1016/j.asoc.2019.105494
12. Rundo L*, **Tangherloni A***, Cazzaniga P, Nobile MS, Russo G, Gilardi MC, Vitabile S, Mauri G, Besozzi D, Militello C (2019). *A novel framework for MR image segmentation and quantification by using MedGA*. **Computer Methods and Programs in Biomedicine**, 176(1): 59-172. DOI: 10.1016/j.cmpb.2019.04.016
13. Rundo L*, **Tangherloni A***, Nobile MS, Militello C, Besozzi D, Mauri G, Cazzaniga P (2019). *MedGA: A Novel Evolutionary Method for Medical Image Enhancement in Medical Imaging Systems*. **Expert Systems with Applications**, 119: 387-399. DOI: 10.1016/j.eswa.2018.11.013
14. **Tangherloni A†**, Spolaor S, Rundo L, Nobile MS, Cazzaniga P, Mauri G, Liò P, Merelli I, Besozzi D (2019). *GenHap: A Novel Computational Method Based on Genetic Algorithms for Haplotype Assembly*. **BMC Bioinformatics**, 20(4): 172. DOI: 10.1186/s12859-019-2691-y
15. Rundo L, Han C, Nagano Y, Zhang J, Hataya R, Militello C, **Tangherloni A**, Nobile MS, Ferretti C, Besozzi D, Gilardi MC, Vitabile S, Mauri G, Nakayama H, Cazzaniga P (2019). *USE-Net: Incorporating Squeeze-and-Excitation blocks into U-Net for prostate zonal segmentation of multi-institutional MRI datasets*. **Neurocomputing**, 365: 31-43. DOI: 10.1016/j.neucom.2019.07.006

16. Rundo L, Militello C, **Tangherloni A**, Russo G, Vitabile S, Gilardi MC, Mauri G (2018). *NeXt for neuro-radiosurgery: A fully automatic approach for necrosis extraction in brain tumor MRI using an unsupervised machine learning technique*. **International Journal of Imaging Systems and Technology**, 28(1): 21-37. DOI: 10.1002/ima.22253
17. **Tangherloni A**, Nobile MS, Besozzi D, Mauri G, Cazzaniga P (2017). *LASSIE: simulating large-scale models of biochemical systems on GPUs*. **BMC Bioinformatics**, 18(1): 246. DOI: 10.1186/s12859-017-1666-0
18. **Tangherloni A**†, Nobile MS, Cazzaniga P, Besozzi D, Mauri G (2017). *Gillespie's Stochastic Simulation Algorithm on MIC Coprocessor*. **The Journal of Supercomputing**, 73(2): 676-686. DOI: 10.1007/s11227-016-1778-8
19. Nobile MS, Cazzaniga P, **Tangherloni A**, Besozzi D (2017). *Graphics Processing Units in Bioinformatics, Computational Biology and Systems Biology*. **Briefings in Bioinformatics**, 18(5): 870-885. DOI: 10.1093/bib/bbw058

Book, book chapters, and contributed volumes

1. Cascino DL, Gatti G, Unwith S, Matarazzo LS, Riva SG, Damiani G, **Tangherloni A**† (2025). *Biomedical Data Science: A Step-by-Step Guide to Analysis and Interpretation*, River Publishers
2. Rundo L, Han C, Zhang J, Hataya R, Nagano Y, Militello C, Ferretti C, Nobile MS, **Tangherloni A**, Gilardi MC, Vitabile S, Nakayama H, Mauri G (2020). *CNN-Based Prostate Zonal Segmentation on T2-Weighted MR Images: A Cross-Dataset Study*. In **Neural Approaches to Dynamics of Signal Exchanges**, 151: 269-280, Springer. DOI: 10.1007/978-981-13-8950-4_25
3. Cazzaniga P, Nobile MS, **Tangherloni A**, Besozzi D (2018). *Accelerating stochastic simulations of mechanistic models of biological systems: Advantages and issues in the parallelization on Graphics Processing Units*. In **Quantitative Biology: Theory, Computational Methods, and Models**, 423-440, MIT Press.
4. Rundo L, Militello C, **Tangherloni A**, Russo G, Lagalla R, Mauri G, Gilardi MC, Vitabile S (2018). *Computer-assisted Approaches for Uterine Fibroid Segmentation in MRgFUS Treatments: Quantitative Evaluation and Clinical Feasibility Analysis*. In **Quantifying and Processing Biomedical and Behavioral Signals, Smart Innovation, Systems and Technologies**, 103: 229-241, Springer. DOI: 10.1007/978-3-319-95095-2_22

Conference Proceedings

1. Egin A, **Tangherloni A**, Dantcheva A (2025). *Now You See Me, Now You Don't: A Unified Framework for Expression Consistent Anonymization in Talking Head Videos*. Accepted at ICCV 2025 (Workshop CV4BIOM)
2. Fabbriatore A, Buffa FM, **Tangherloni A**† (2025). *Knowledge-Enriched Cell-Type Annotation in Single-Cell Transcriptomics via LLM Embeddings*. Proceeding of **International Conference on Computational Intelligence in Bioinformatics and Computational Biology**, pp. 1-8
3. Riva SG, Myers B, Buffa FM, **Tangherloni A**† (2025). *GENESIS: Generating scRNA-Seq data from Multiome Gene Expression*. Proceeding of **International Conference on Computational Intelligence in Bioinformatics and Computational Biology**, pp. 1-8
4. Papetti DM, **Tangherloni A**, Coelho V, Besozzi D, Cazzaniga P, Nobile MS (2025). *We Are Sending You Back... to the Optimum! Fuzzy Time Travel Particle Swarm Optimization*. In **Applications of Evolutionary Computation, Lecture Notes in Computer Science**, 15613: 160-175, Springer. DOI: 10.1007/978-3-031-90065-5_10
5. **Tangherloni A**†, Cazzaniga P, Stranieri N, Buffa FM, Nobile MS (2024). *A Fast Feature Selection for Interpretable Modeling Based on Fuzzy Inference Systems*. Proceeding of **International Conference on Computational Intelligence in Bioinformatics and Computational Biology**, pp. 1-8, IEEE. DOI: 10.1109/CIBCB58642.2024.10702107
6. Stranieri N, Buffa FM, **Tangherloni A**† (2024). *Forest-based Evolutionary Algorithm for Reconstructing Boolean Gene Regulatory Networks*. Proceeding of **International Conference on Computational Intelligence in Bioinformatics and Computational Biology**, pp. 1-8, IEEE. DOI: 10.1109/CIBCB58642.2024.10702181
7. **Tangherloni A**†, Coelho V, Buffa FM, Cazzaniga P (2024). *A modified EACOP implementation for*

8. Riva SG, Myers B, Cazzaniga P, Buffa FM, **Tangherloni A†** (2023). *Consensus clustering strategy for cell type assignments of scRNA-seq data*. Proceeding of **International Conference on Computational Intelligence in Bioinformatics and Computational Biology**, pp. 1-8, IEEE. DOI: 10.1109/CIBCB56990.2023.10264908
9. Coelho V, Papetti D, **Tangherloni A**, Cazzaniga P, Besozzi D, Nobile MS (2023). *The Domination Game: Dilating Bubbles to Fill Up Pareto Fronts*. Proceeding of **Congress on Evolutionary Computation**, pp. 1-8, IEEE. DOI: 10.1109/CEC53210.2023.10253982
10. **Tangherloni A†**, Riva SG, Myers B, Cazzaniga P (2022). *Multi-objective Optimization for Marker Panel Identification in Single-cell Data*. Proceeding of **International Conference on Computational Intelligence in Bioinformatics and Computational Biology**, pp. 1-8, IEEE. DOI: 10.1109/CIBCB55180.2022.9863042
11. Riva SG, Myers B, Cazzaniga P, **Tangherloni A†** (2022). *A Deep Learning Pipeline for the Automatic cell type Assignment of scRNA-seq Data*. Proceeding of **International Conference on Computational Intelligence in Bioinformatics and Computational Biology**, pp. 1-8, IEEE. DOI: 10.1109/CIBCB55180.2022.9863034
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