JIANFENG SUN

computational biology, artificial intelligence, data science

I am working broadly in computational biology at the University of Oxford. I use mathematical algorithms to model biological systems in structural biology, molecular biology and cancer biology. I am carving out a niche for myself as a leading researcher in UMI-assisted research and computational drug discovery.



EDUCATION

2020 2017

Ph.D., Dr. rer. nat., Artificial Intelligence Structural Biology

Technical University of Munich

Munich, Germany

- · biological protein science, evolutionary biology
- · computational deep learning, statistics
- specifics deep learning for predicting protein interaction sites

2014 2016 M.Eng., Software Engineering and Computational Biology

Beijing Forestry University

Paeijing, China

- · biological genetics
- · computational algorithm design, web application development
- · specifics mathematical modelling for quantitative trait loci (QTLs) detection

2014 2011 B.S.Mgt.Sci., Accounting (minor dual-degree)

Nanjing Tech University

Nanjing, China

· economics

2014 2010 B.Sci., Information and Computer Science (i.e., Computational Mathematics)

Nanjing Tech University

Nanjing, China

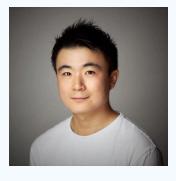
- · mathematics
- statistics
- · programming

RESEARCH EXPERIENCE

2025 2021 Postdoctoral Researcher in Single-cell Sequencing Analysis

University of Oxford

- Oxford, United Kingdom
- · biological transcriptomics, protein science
- · computational deep learning, statistics
- · duty computational analysis of single-cell data in sarcoma diseases and algorithm design for accurate long-read sequencing technology



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CONTACT

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- github.com/2003100127
- **y** Jianfeng_Sunny
- in jianfeng-sun-2ba9b1132
- @ 2003100127.github.io

LANGUAGE SKILLS

| Python |
|---------------|
| R |
| PHP (Laravel) |
| Bash |
| SQL |
| HTML-5 |
| Javascript |

Last updated on 2025-04-07.

2021 2020

Post-doctoral Studies in Transmembrane Proteome Analysis

Technical University of Munich

- Munich, Germany
- biological protein isoforms, mutations, variants, interaction sites
- computational machine learning, statistics
- duty Occurrences of mutation sites and interaction sites in the human transmembrane proteome



PROFESSIONAL SERVICE

Current 2024

Current 2024

Full member of Sigma XI, The Scientific Research Honor Society

Oxford, United Kingdom

Young Editorial Member of iMeta

Oxford

Oxford

Oxford, United Kingdom



THESIS

2021 2020 Prediction of residue contacts and interaction sites in transmembrane proteins using deep learning

Technical University of Munich

Munich, Germany

- · Ph.D. Defense
- · committee chairman Prof. Bernhard Küster
- examiner Prof. Burkhard Rost
- · link https://mediatum.ub.tum.de/doc/1577512



ACADEMIC PUBLICATION

Journal Articles, Conference Proceedings, Book Chapters, etc.

Please refer a full list of my publications at the sites below

- · Google Scholar
- ORCID
- ResearchGate

I updated my latest research at these websites on a regular basis. I primarily published articles in the topics of methods, computational tools, and their applications in molecular and disease biology.

PATENT

Current 2022

Silkworm silk gland recombinant expression vector for expressing human epidermal growth factor and preparation method and application thereof

China National Intellectual Property Administration (CNIPA)

China

- · inventors Dingpei Long, Jian Cheng, Jianfeng Sun, Zhonghuai Xiang, and Fangyin Dai (by order)
- · patent No. CN112852876A
- · link https://patents.google.com/patent/CN112852876A/en?og=cn112852876
- · my role algorithm design for deducing protein functions based on their experimental and predicted structures



FUNDING

2024 2023

Cancer Research UK (CRUK) Oxford Development Fund

University of Oxford

Oxford, United Kingdom

- Project Title Single-cell spliceosome map establishment of immune cells
- · Award Ref CRUKDF-MAY23-AC/JS



SOFTWARE COPYRIGHT

2016

The web system for differentiating dynamic complex traits based on growth curve

Beijing Forestry University

Peijing, China

- · Copyright No. 2016SRBJ047
- · Register Beijing Forestry University
- · Developer Jianfeng Sun

2016 2015

Climate Change Adaptation Information System for Beijing Landscape and Forestry Industry

Beijing Forestry University

Peijing, China

- Copyright No. 2016SR098798
- · Register Zhibo Chen & Jianfeng Sun
- · Developer Jianfeng Sun



SCIENTIFIC PROGRAM

Current 2022

Workshop in computational single-cell sequencing analysis

Koç University

Istanbul, Turkey

· my role - teaching assistant

I applied the golden rule to a protein material study: protein structures determine their functions. This is used to screen synthetic proteins with similar biological activities to commericialised proteins.

Invited pre-doctoral programs by the International Max Planck Current Research School for Computational Biology and Scientific 2017 Computing (IMPRS-CBSC), Parlin, Germany Max Planck Institute for Molecular Genetics CONFERENCE 2019 international conference on machine learning and cybernetics 2019 (ICMLC) Kobe Convention Center • The LSTM Network for residue-residue contacts prediction CPEN TALK Artificial Intelligence Applications on Biomedical Data 2025 Zoom online - RobotGym GmbH (German Company) Oxford, United Kingdom • Introduction of advanced artificial Intelligence methods, including language models and variational inference techniques, to the application in biomedical Machine Learning in Structural Biology 2020 Tecent VooV meeting online - Koushare Academic Platform Munich, Germany · Presentation and Q&A AWARD Third Prize of The 8th China (Shenzhen) Innovation & 2024 **Entrepreneurship International Competition (Milan Division)** Milan, Italy Milan 2nd rank among 20 screened teams in 2020 Munich Impact 2020 **Hackathon Programming Competition** Munich, Germany Munich **China National Scholarship** 2016 Peijing, China Beijing Forestry University 2015 China National Scholarship 2015 Peijing, China Beijing Forestry University 2014 University Third-prize Scholarship of Nanjing Tech University 2014 Nanjing, China Nanjing Tech University 2013

| 2013 | University First-prize Scholarship of Nanjing Nanjing Tech University | Tech University ♥ Nanjing, China | | | |
|-------------------------------|--|--|------------------------|--|--|
| 2012 2011 | University Third-prize Scholarship of Nanjing Nanjing Tech University | § Tech University ● Nanjing, China | | | |
| 2011 | University Third-prize Scholarship of Nanjing Nanjing Tech University | Tech University ♥ Nanjing, China | | | |
| 2011 | University Third-prize Scholarship of Nanjing Nanjing Tech University | § Tech University ● Nanjing, China | | | |
| 2011 | Top 8 Finalist in the Nanjing Regional Finals o | of 2011 China's Got | | | |
| 2010 | Wanda Plaza | • Nanjing, China | | | |
| | sponsor of this event - Dagexing Co., Ltdkind - dancing | | | | |
| > | COMPUTATIONAL TOOL | | | | |
| Current 2025 | UMIche | • | | | |
| | University of Oxford | Oxford, United Kingdom | | | |
| | Website - https://2003100127.github.io/umiche | | l am pass | | |
| Current 2025 | mclUMI | _ | techniqu | | |
| | University of Oxford | Oxford, United Kingdom | for biolog develope | | |
| | • Website - https://2003100127.github.io/mclumi | | methods available | | |
| Current | Tresor | | release r | | |
| 2025 | University of Oxford | Oxford, United Kingdom | | | |
| | Website - https://2003100127.github.io/tresor | | | | |
| Current 2025 | PyPropel | | | | |
| | University of Oxford | Oxford, United Kingdom | | | |
| | Website - https://2003100127.github.io/pypropel | | | | |
| Current | TMKit | | | | |
| 2023 | University of Oxford | Oxford, United Kingdom | | | |
| | • Website - https://2003100127.github.io/tmkit | | | | |
| Current | DeepdIncUD | | | | |
| 2024 | University of Oxford | Oxford, United Kingdom | | | |
| | Website - https://2003100127.github.io/deepdlncud | | | | |

I am passionate about fantastic techniques applied in data science for biological systems. I have developed dozens of tools and methods, with nearly 15 publicly available to check and use. I will release more!!

