

# Shengliang NI

2025/05/02 update

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[nslbotnslbot.github.io](https://github.com/nslbotnslbot)/[nslbotcola.github.io](https://github.com/nslbotcola)

## EDUCATION

### The University of Tokyo (current)

PhD of Sciences (Computational Biology and Medical Sciences)

Supervisor: Martin Frith

Tokyo, Japan

Expected 2025

### The University of Tokyo

Master of Sciences (Computational Biology and Medical Sciences)

Supervisor: UI-TEI Kumiko

Tokyo, Japan

Received 2021

### Zhejiang University

Bachelor of Agronomy (Agriculture and Biotechnology)

Hangzhou, China

Received 2018

## RESEARCH INTERESTS

I study genomic elements such as pseudogenes, repetitive sequences, and non-coding RNA loci across various genomes—including those of eukaryotes, prokaryotes, and specialized cell lines like breast cancer—using bioinformatics and statistical approaches.

## RESEARCH

If the conference is associated with journal, names of the journal will be listed.

\* indicates co-first authors

(Selected research)

1. Ni S\*, Shu S\*, Kato M, Saitoh N and Frith M C. Repeat elements enriched in cis-regulatory regions act in cancer cell transition to estrogen-independence. **GIW ISCB-Asia 2023** (recommend to *NAR Genomics and Bioinformatics*).
2. Frith M C, Ni S. DNA conserved in diverse animals since the Precambrian controls genes for embryonic development *Molecular Biology and Evolution*. (2023).
3. Zhang H\*, Ni S\* and Frith M C. An immune-suppressing protein in human endogenous retroviruses *Bioinformatics Advances* (2023, GIW ISCB-Asia 2022 associated).

(Other posters and presentations)

1. Ni S\*, Shu S\*, Kato M, Saitoh N, Frith M C, Palihati M, Carninci P. Repeat elements enriched in cis-regulatory regions act in cancer cell transition to estrogen-independence. **Poster. Cold Spring Harbor Asia meeting on Chromatin, Epigenetics & Transcription (2024).**
2. Ni S\*, Shu S\*, Kato M, Saitoh N and Frith M C. Investigating the role of repeat elements in promoting the transition of MCF-7 cells to LTED cells. **Poster. The 46<sup>th</sup> annual meeting of MBSJ (2023).**
3. Ni S\*, Shu S\*, Kato M, Saitoh N and Frith M C. Investigating the role of repeat elements in promoting the transition of MCF-7 cells to LTED cells. **Poster. The 68<sup>th</sup> HGA (2023).**

4. Ni S\*, Shu S\*, Kato M, Saitoh N and Frith M C. Investigating the role of repeat elements in promoting the transition of MCF7 cells to LTED cells. **Oral Presentation. The 11<sup>th</sup> IIBMP (2022).**
5. Ni S\* and Frith M C. K The protein fossil record in prokaryote genomes: a hidden treasure. **Oral Presentation. The 11<sup>th</sup> IIBMP (2022).**
6. Ni S\* and Ui-Tei K. Design and validation of antiviral siRNAs for targeting the conserved regions in coronavirus. **Poster. The 43<sup>rd</sup> annual meeting of MBSJ (2020).**
7. Ni S\* and Ui-Tei K. Design and validation of antiviral siRNAs for targeting the conserved regions in coronavirus. **Poster. The 25<sup>th</sup> annual meeting of RNA Biology (2020).**

## INTERNSHIPS AND EXPERIENCE

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- Data Scientist Training/Education Program (DSTEP) 2022-2024
- Procter & Gamble 2<sup>nd</sup> IT Tech Challenge (Hackathon), Tokyo, Japan Sep 2020
- Research Student, Supervisor: UI-TEI Kumiko 2018-2019

## AWARDS

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- Ushio Scholarship 2022-2024
- Zhejiang Rural Credit Cooperative International Exchange Scholarship 2017
- Scholarship for Special Major, Zhejiang University 2016, 2017
- Scholarship for Excellence in Arts and Sports, Zhejiang University 2015

## SKILLS

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**Languages:** English (Fluent); Chinese (Native); Japanese

### Technical Skills:

- **Programing:** UNIX shell (first priority), Python, R, MySQL, HTML5, PHP, C/C++.
- **Computational skill:**
  - Experiences and knowledge of statistics and multi-omics data analysis in bioinformatics (LAST, DESeq2, Seurat, edgeR, clusterProfiler, etc.);
  - Fundamental applications of machine learning and deep learning (scikit-learn, XGBoost);
  - Basic knowledge of quantum computation (Qiskit).