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[Personal website](#)

Sex: Male | Date of birth: 17/02/2002 | Nationality: Chinese

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## RESEARCH EXPERIENCE

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**Project Leader (Finished) --- immunology 2022/07--2023/04**

**China University Life Science Competition (ULSC)**

**Project Name: Research on Macrophage Death Induced by Cytoplasm Membrane Vesicles(CMV)s Separated from Streptococcus Agalactiae Cells**

This research focus mechanism of host cell inflammation and cell death caused by the interaction of cytoplasm membrane vesicles(CMV)s released by streptococcus agalactiae(GBS) with mouse macrophages. Bacteria can transfer bacterial molecules to host cells through CMVs, and inducing cell death with these factors in the CMVs. In this research we studied on CMVs produced by GBS during vivo infection and their direct association with host cell pathology, conducted research on the extraction and identification of CMVs from GBS, as well as the effects of CMVs on macrophage proliferation in mice, the expression of macrophage inflammatory cytokines induced by CMVs, the death of macrophage induced by CMVs.

**Project Leader (Finished) --- Gene Engineering 2021/07--2023/05**

**National Innovation and Entrepreneurship Competition Fund**

**Project Name: Recombinant Tumor Vaccine for Breast Cancer HER2 Antigen Based on OMV**

This project aims to develop a novel recombinant tumor vaccine for breast cancer HER2 antigen based on bacterial outer membrane vesicle (OMV). The vaccine uses ClyA protein as a guiding sequence to locate HER2 antigen to the outer membrane of E. coli and elicit immune response in mouse macrophages. The vaccine is constructed by cloning and fusing ClyA and HER2 genes, expressed in E.coli, separated by overspeed centrifugation, and identified by molecular biology techniques.

**Project Leader (Finished) ---Bioinformatics 2024/12--2024/04**

**Graduation Projects**

**Project Name: Benchmarking Algorithms for Batch Correction in Single-cell ATAC-seq data Analysis**

The complexity of genome dataset is increasing, and current dataset often include different sample, generated across multiple conditions. The methods of intergration can minimize this complexity of dataset. In this project, we present a benchmarking study of data intergration methods in complex intergration tasks,Through systematic benchmarking, we aim to assess the efficacy, accuracy, and scalability of different integration methodologies. By leveraging a diverse set of datasets reflecting real-world scenarios, providing insights into the performance of these software tools under varying conditions. The project's outcomes will not only inform the selection of optimal integration methods but also contribute valuable knowledge to the evolving landscape of single-cell genomics research.

## EDUCATION AND TRAINING

### Biological Chemical Engineering (Double major: English) 2020/09--2024/06

Inner Mongolia University(IMU) (Monitor of the Class)

Address: No. 24, Zhaojun Road; Yuquan District, Hohhot City, Inner Mongolia 010000

Major Courses(Bold font represents a score exceeding 85): **Bioinformatics**, **Probability Theory and Mathematical Statistics**, **Professional Foreign Language (English)**, General Biology, **Instrumental Analysis**, **Cell Biology**, Physiology, **Microbiology**, **Biological Statistics**, Biochemistry, College Chemistry, **Cell Engineering**, Advanced Mathematics, **Genetic engineering**, Molecular biology, **Principle of chemical engineering**

Comprehensive Evaluation: Rank 1 in 30

### Bioinformatics 2023/12--2024/04

Westlake University (Visiting student)

No.18 Shilongshan Road, Xihu District, Hangzhou, Zhejiang, 310024

Zhang Kai Lab ([Lab website](#))

Major job: Finishing the content of the **graduation project** and help the RA to do **some data analysis job**. Also supplement laboratory teaching documents.

Learned Skills: Basic concepts and methods of **Computational biology**, become proficient in **data science**. Ability of **making English presentation**.

### Chemical Biology 2024/09--Now

University of Geneva and École polytechnique fédérale de Lausanne(EPFL)

Geneva and Lausanne, Switzerland

Major Courses: Elements in bioinformatics, Machine Learning(Auditor), Microscopy and imaging course

## PERSONAL SKILLS

Self assessment	Data processing	Data Mining	English Writing	Literature reading	Problem solving
	Competent	Basic	Familiar	Proficient	competent
❖	Familiar with Systems such as Windows, MacOS, <b>Ubuntu or CentOS</b> , as well as some tools like “DepMap”, “UCSC genome browser” or GEO database, can manage my code with Github				
❖	Understanding the basic syntax of <b>Python</b> and familiar with some packages of python such as NumPy, Pandas, Matplotlib				
❖	Familiar with <b>Jupyter notebook</b> and VSCode				

## HONORS & PUBLICATIONS

Honors	09.2024, Merit-based Fellowship from SNE Chemical biology
	09/2023, Selected as the candidate of “JingYing Scholar” Training Program of Inner Mongolia University
	04/2023, Third-Class Model Scholarship of Inner Mongolia University
	04/2023, Undergraduate First-class Academic Scholarship of Inner Mongolia University
	05/2022, 2021 Outstanding Youth League Cadre of Inner Mongolia University
	05/2022 University-level Merit Student of Inner Mongolia University
	04/2022, Undergraduate Second-class Academic Scholarship of Inner Mongolia University
	05/2021, 2020 Outstanding League Member of Inner Mongolia University
	04/2021, Undergraduate Third-class Academic Scholarship of Inner Mongolia University
Publications	[1]Sun Cai, Wu Yanhao, <b>Guo Jiayi</b> . Research Progress in the Application of OMV in Tumor Vaccines[J]. <i>Chinese Bulletin of Life Sciences</i> , 2022,34(11):1431-1441.DOI:10.13376/j.cbls/2022157,
	[2]Wang S, <b>Guo J</b> , Bai Y, et al. Bacterial outer membrane vesicles as a candidate tumor vaccine platform. <i>Front Immunol</i> . 2022;13:987419. doi:10.3389/fimmu.2022.987419