

DONGYUAN WU

GRADUATE STUDENT

Department of Biostatistics
University of Florida
2004 Mowry Rd, CTRB 5th Floor
Gainesville, FL 32610
☎ +1 (352) 870 4802
✉ dongyuanwu@ufl.edu
📁 [dongyuanwu.github.io](https://github.com/dongyuanwu)

Last updated on May 23, 2022

EDUCATION

University of Florida

Doctor of Philosophy in Biostatistics

Gainesville, FL

Aug. 2020 - present

University of Florida

Master of Science in Biostatistics

Gainesville, FL

Aug. 2018 - May 2020

Minzu University of China

Bachelor of Science in Applied Statistics

Beijing, China

Sept. 2013 - July 2017

WORK EXPERIENCE

Graduate Research Assistant

Department of Biostatistics, University of Florida

Gainesville, FL

Aug. 2020 - present

The Role and Mechanisms of Lipid and Lipoprotein Dysregulation in Sepsis

- Analyzed demographic and clinical data with descriptive statistics, examined the existence of significant differences among different groups of sepsis patients.
- Aligned RNA sequencing data into expression matrix, did further downstream analyses such as differential expression analysis, gene co-expression network analysis, functional annotation, and so on.
- Dealt with missing values and unbalanced issues of lipidomics data, and then conducted suitable comparisons based on different clinical outcomes, phenotypes, etc.

Graduate Teaching Assistant

Department of Biostatistics, University of Florida

Gainesville, FL

Aug. 2020 - present

Grading and office hour responsibilities

- PHC 6052: Introduction to Biostatistical Methods (Fall 2021)
- PHC 6051: Biostatistical Methods II (Spring 2021)
- PHC 6020: Clinical Trial Methods (Spring 2021)
- PHC 6790: Biostatistical Methods Using SAS (Fall 2020)
- PHC 6050: Statistical Methods for the Health Sciences I (Fall 2020)
- PHC 6089: Public Health Computing (Fall 2020)

OPS Student Research Assistant

*Department of Community Health and Family Medicine,
University of Florida*

Gainesville, FL

May 2019 - Apr. 2020

- Collaborated with investigators to identify problems and provided biostatistical consultation including analysis of data, interpretation of results, and preparation of reports.

Research Assistant

*Institute of Basic Research in Clinical Medicine,
China Academy of Chinese Medical Sciences*

Beijing, China

July 2017 - June 2018

- Provided a variety of statistical analysis for projects using appropriate statistical and computing methodologies, and assisted in the interpretation and presentation of results.
- Imported data from SQL, generated reproducible templates for adverse drug reaction weekly report, monthly report, and annual report by using R Markdown, and assisted to develop a platform that can automatically display these reports for different drugs.

**RESEARCH
EXPERIENCE**
Cell-Cell Communication on Spatial RNA-seq Data

Supervisor: Dr. Susmita Datta

Gainesville, FL

Jan. 2022 - present

Immune Cell Responses to SARS-CoV-2 Infection

Supervisor: Dr. Susmita Datta

Gainesville, FL

Apr. 2021 - Feb. 2022

- Preprocessed datasets, such as filtering, normalization, and resampling.
- Applied MAST for multiple single-cell RNA sequencing data related to immune cells, to figure out the differential expression genes among healthy donors, mild COVID-19 patients, and severe COVID-19 patients.
- Made gene co-expression networks using PIDC based on healthy, mild COVID-19, and severe COVID-19, separately. Then found the hub genes in modules and detected the differentially connected genes.
- Built random forests to calculate the Gini importance scores for genes.
- Did functional annotation for those detected important genes to improve the insight of gene expression and network patterns associated with immune cells for SARS-CoV-2 infection.

Analysis of Alzheimer's Disease scRNAseq Data

Supervisor: Dr. Susmita Datta

Gainesville, FL

Sept. 2019 - Apr. 2020

- Used 2-dimensional t-SNE plots to visualize the single-cell RNA sequencing data.
- Preprocessed dataset, such as splitting, filtering, normalization, and clustering.
- Applied hurdle models specifically designed for sequencing-based single-cell gene expression data, including CRE and MAST, to detect differentially expressed genes between AD-pathology and no-pathology.
- Did network analysis and functional annotation for those differentially expressed genes.

Risk Factors of ADR for a Traditional Chinese Medicine

Supervisor: Dr. Wei Yang

Beijing, China

Oct. 2016 - May 2017

- Processed 30,888 data in advance, such as data cleaning, data standardization.
- Analyzed all data with descriptive statistics, examined the existence of significant differences between normal group and adverse drug reaction (ADR) group by applying various hypothesis testing methods, and reported the results by using R Markdown.
- Associated resampling methods, including RUS, ROS, and SMOTE, with classification algorithms, such as decision trees, AdaBoost, random forests, and LASSO, to improve the accuracy of classifiers for the minority class (i.e., ADR group).

Evaluation of TCM Clinical Practice Guidelines

Supervisor: Dr. Wei Yang

Beijing, China

Aug. 2015 - Sept. 2016

- Processed more than 20,000 items of Traditional Chinese Medicine (TCM) data in advance, including data cleaning, data standardization.
- Analyzed data in 76 TCM syndrome and 11 subjects with descriptive statistics, and reported the results by using R Markdown based on the idea of reproducible research.

REFEREED PUBLICATIONS

DelRocco, N., Wang, Y., **Wu, D.**, Yang, Y., Shan, G. (2022). New Confidence Intervals for Relative Risk of Two Correlated Proportions. *Statistics in Biosciences*. doi: 10.1007/s12561-022-09345-7

Wu, D., Zhang, R., and Datta, S. (2022). Unraveling T Cell Responses for Long Term Protection of SARS-CoV-2 Infection. *Frontiers in Genetics*, 13:871164. doi: 10.3389/fgene.2022.871164

Malaty, J., **Wu, D.**, and Datta, S. (2021). Flexible Nasal Endoscopic Procedures in Family Medicine: Indications and Effectiveness. *Family Medicine*, 53(10), 886-889. doi: 10.22454/FamMed.2021.332061

Ellis, D., **Wu, D.**, and Datta, S. (2021). SAREV: A review on statistical analytics of single-cell RNA sequencing data. *Wiley Interdisciplinary Reviews: Computational Statistics*, e1558. doi: 10.1002/wics.1558

ORAL PRESENTATIONS

“*UTRCOV2: Unraveling T cell responses for long term protection of SARS-COV-2 infection.*” Annual International Conference on Critical Assessment of Massive Data Analysis. July 2021. Virtual.

POSTER PRESENTATIONS

“*UTRCOV2: Unraveling T cell responses for long term protection of SARS-COV-2 infection.*” UF College of Medicine Research Day. April 2022. Gainesville, FL.

“*COVID-19: Reduced Lung Function and Increased Psycho-emotional Stress.*” UF College of Public Health & Health Professions Research Day. February 2021. Virtual.

SOFTWARES

- **PairedRR**: An Shiny app to calculate the confidence intervals for relative risk of two correlated proportions.
- **RSBID**: An R package to provide resampling strategies for binary imbalanced datasets.

AWARDS AND HONORS

Student Travel Award	Department of Biostatistics, University of Florida	2021
Grinter Fellowship	University of Florida	2020
Outstanding Master Graduate Certificate of Excellence	Department of Biostatistics, University of Florida	2020
First-class Scholarship (TOP 3%)	International Center, University of Florida	2019
Honorable Mention	Minzu University of China	2017
First Prize in Beijing Region	Interdisciplinary Contest in Modeling	2016
Wu Xianhong Scholarship	National Mathematical Modeling Contest	2016
	College of Science, Minzu University of China	2015, 2016

SKILLS

R, SAS, Python, L^AT_EX, C/C++, WinBUGS, MySQL

CERTIFICATES

- [SAS Certified Professional: Advanced Programming Using SAS 9.4](#) (2020)
- [SAS Certified Specialist: Base Programming Using SAS 9.4](#) (2019)
- [Applied Data Science with Python Specialization](#) by UMich on Coursera (2020)
- [Genomic Data Science Specialization](#) by JHU on Coursera (2020)
- [Data Science Specialization](#) by JHU on Coursera (2019)