# Weijia Xiong

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## **EDUCATION**

Columbia University, Mailman School of Public Health, New York, NY

expected May 2021

Master of Science, Biostatistics GPA: 3.92

East China Normal University, Shanghai, China

June, 2019

Bachelor of Science, Statistics GPA: 3.6

University of California at Los Angeles, Los Angeles, CA

Summer, 2017

Summer Program, Statistics

#### PROFESSIONAL EXPERIENCE

Roche Shanghai, China

# Statistical Programming Analysis (SPA) Intern

July, 2018 – Dec, 2018

- Participated in development of AutoReport RShiny app to streamline regulatory reporting process and integrate all related data files and documents in a one-stop shop; currently online and actively used.
- Completed design of GitHub web pages for team projects.
- Assisted team for with user acceptance testing (UAT), including questionnaire creation, data collection and final review.

## RESEARCH EXPERIENCE

# Columbia University Research

New York, NY

# **Exploration of Super Learner algorithm**

 $Aug,\,2020-Present$ 

Research assistant, advised by Dr. Caleb Miles

- Learned and reproduced the analysis of OTNY data, predicting three months of education enrollment or employment among unenrolled and unemployed at baseline via Super Learner algorithm.
- Applied Super Learner to Heart disease dataset and compared the predicted classification error rate with other machine learning methods (eg. GLM, Lasso, SVM, etc).

# An application of Hidden Markov model for mobile mental health study

June, 2020 – Present

Research assistant, advised by Dr. Linda Valeri

- Simulated outcome survey data and covariates (e.g. mobile-communication data like phones and calls, season, monthly clinical diagnostic scores) with mutual relations (e.g. AR (1), nonlinear pricewise function) between latent disease states.
- Created a mixed-effects Hidden Markov model with a multinomial logistic regression model for time-dependent covariates.
- Explored the causal effect of mobile-communication data and evaluated the prediction performance.
- Re-performed the model considering different missing data scenarios (MCAR, MAR, and MNAR) with different imputation methods (MICE, missForest, and median) or removing. Compared the results of prediction error rate.

# Longitudinal microbiome data analysis

Apr, 2020 - Present

Research assistant, advised by Dr. Jianhua Hu and Dr. Jian Wang

- Participated in a vaginal longitudinal microbiome research. Assisted in proposing a new cluster method for overall trajectory.
  - Imputed missing values via the KNN method.
  - Reproduced the clustering work for vaginal community state type and performed other methods (Hierarchical, K-means)
  - Conducted comparison of preterm birth classification for different methods (Lasso, Random Forest, SVM, etc).
- Participated in a longitudinal analysis of the microbiome in acute myeloid leukemia (AML) patients via Logistic Functional Additive Regression Method.
  - Performed data pre-processing and analysis for compositional data.
  - Implemented the algorithm to predict the infection of AML and assisted in building the package

## Analysis of vehicle collision in New York City

Sep, 2019 – Dec, 2019

Research assistant, advised by Dr. Jeff Goldsmith

- Analyzed NYPD motor vehicle collisions-crashes data with related covariates weather, traffic light, restaurant and holiday.
- Built the regression model and analyzed the correlation between potential factors and vehicle collisions.
- Created a website with vehicle collision map and shiny dashboard for data visualization and conclusions.

## **PUBLICATION**

Bao, Y\*, Xiong, W\*(2020). One-round communication efficient distributed M-estimation (submitted)

Bao,  $Y^*$ , Liu,  $W^*$ , Mao,  $X^*$  and **Xiong, W**\*. Byzantine-Tolerant distributed multiclass sparse discriminant Analysis (manuscript in preparation)

## TECHNICAL SKILLS

Computer skills: R, Matlab, LaTex, SAS, VS Code, Git