

ZIQUIANG CHEN

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EDUCATION

University at Buffalo, Buffalo, NY

Expected 2020

Ph.D. in Biostatistics, GPA: 3.9/4.0

University at Buffalo, Buffalo, NY

Jul 2015

M.A. in Biostatistics, GPA: 3.9/4.0

The Second Institute of Oceanography, Hangzhou, China

Jul 2013

M.S. in Marine Biology, GPA: 3.7/4.0

Shandong University, Weihai, China

Jul 2009

B.Sc. in Biological Science, GPA: 3.0/4.0

DISSERTATION RESEARCH

Topic: Selected topics in statistical methods for drug and diagnostic device development.

Thesis Advisor: Dr. Gregory E. Wilding

- Unconditional exact tests based on multiple comparisons for Poisson rate.
- ROC analysis for multistage diagnostic testing procedures in the presence of intermediate results.

PROFESSIONAL EXPERIENCE

HTG Molecular Diagnostic, Inc.

Tucson, AZ

Biostatistician, Intern

Jan 2018 - Jan 2019

- Participated in several collaborative projects in Companion Diagnostics (CDx) branch.
- Work duties included statistical analysis plan, exploratory analysis and data visualization, literature review, method implementation and development, deliverable preparation and presentation.

School of Social Work, University at Buffalo

Buffalo, NY

Research Assistant

Jun 2019 - pres.

- Provided support to research projects, performed statistical analysis, wrote up reports and statistical section for grant proposals and papers.

Clinical and Translational Science Institute

Buffalo, NY

Research Assistant

Aug 2014 - pres.

- Planned and managed more than 50 research projects under supervision.
- Experienced in collaboration work for statistical consulting and quantitative analysis, including experimental design, clinical trials, longitudinal studies and meta-analysis, etc.
- Built an “in-house” R library of various statistical analyses for consulting purpose.

Department of Biostatistics, University at Buffalo

Buffalo, NY

Research Assistant

Jun 2014 - Jun 2016

- Supported professors in methodology development in genomics data and survey data.
- Parallel computing through Unix server on compute cluster.

PROFESSIONAL SERVICE

Reviewer, *Emerging Infectious Diseases*

2020 - pres.

SOFTWARE AND SKILLS

Proficient in R, SAS, SPSS, Arc-GIS, Graphpad Prism and MS Office.

SELECTED PROJECTS

Intern works:

Diffuse Large B-Cell Lymphoma (DLBCL) Cell of Origin Assay LoD feasibility study.

- The goal of this project was to evaluate the LoD of CDx assay for identifying DLBCL subtypes.
- Participated in assay data read-in and manipulation, exploratory visualization, principal component analysis and probit model.

Lung Subtyping Assay (LSA) classifier build.

- Built classifier to subtype non-small-cell lung cancer and used in samples from Veliparib clinical trial.
- Participated in statistical analysis plan, QC metric investigation, classifier build, repeatability, analytical sensitivity and robustness studies.

Tumor Mutational Burden (TMB) exploratory analysis.

- Member of a two-person team that conducted TMB association study and enrichment study.
- Responsible for analysis of RNAseq and whole exome sequencing data, including visualization, feature selection, clustering, and p-value adjustment via permutation.

Collaborative works:

Salivary metals, age, and gender correlate with cultivable oral *Candida* carriage levels.

- Addressed the LoD issue in metal measurement using a conditional expected value algorithm.
- Evaluated and utilized zero inflated negative binomial model (ZINB) for excess zeros in outcome.

Disease modifying effect in Alzheimer's disease with cholinesterase inhibitors.

- Longitudinal analysis with autoregressive linear mixed-effects model, multiple comparison, profile plots.

Methodology development:

Confidence interval of Cronbach's alpha with application to complex survey data.

- Developed a novel bootstrap method for the inference of the alpha coefficient.
- Incorporated complex survey design and applied in NCS-R data.

Logistic regression with misclassification in both outcome and predictors

- Developed a two-step correction method to estimate the parameters in logistic regression with misclassification.
- Evaluated and compared the new method with some existing methods and applied in NHANES data.

A novel exact method for Top-K family-wise error control

- Proposed an exact test for the inference of top-K selected features in a two stage study.
- Developed an R package and corresponding Shiny apps.

HONORS AND AWARDS

Best Young Researchers' Award, The Upstate Chapter of American Statistical Association	2017
Poster of Distinction at UB Research Day, University at Buffalo	2017
Perry Poster Day 2016 Winner, Biostatistics, University at Buffalo	2016
Outstanding Graduate, The Second Institute of Oceanography	2013

SELECTED PUBLICATIONS

Published:

Chen, Z., Sun, Q., Chen, Q., Sun, J., Zeng, J. (2013). Effect of acid stress on the early life stages of *Sargassum horneri*. *Asian Journal of Ecotoxicology*, 8(6): 864-870.

Chen, Z., Shou, L., Liao, Y., Gao, A., Zeng, J., Chen, Q. (2013). Community structure of benthic algae and its seasonal variation in the rocky intertidal zone of Sanya. *Acta Ecologica Sinica*, 33(11): 3370- 3382.

Castilla-Earls, A. P., Restrepo, M. A., Perez-Leroux, A. T., Gray, S., Holmes, P., Gaile, D. P., **Chen, Z.** (2016). Interactions between bilingual effects and language impairment: Exploring grammatical markers in Spanish-speaking bilingual children. *Applied psycholinguistics*, 37(5), 1147-1173.

Castilla-Earls, A., Pérez-Leroux, A. T., Restrepo, M. A., Gaile, D. P., **Chen, Z.** (2018). The complexity of the Spanish subjunctive in bilingual children with SLI. *Language acquisition*, 25(1), 72-84.

Norris, H. L., Friedman, J., **Chen, Z.**, Puri, S., Wilding, G. E., Edgerton, M. (2018). Salivary metals, age, and gender correlate with cultivable oral *Candida* carriage levels. *Journal of oral microbiology*, 10(1), 1447216.

Perez, A. C., Johnson, A., **Chen, Z.**, Wilding, G. E., Malkowski, M. G., Murphy, T. F. (2018). Mapping protective regions on a three-dimensional model of the *Moraxella catarrhalis* vaccine antigen oligopeptide permease A. *Infection and immunity*, 86(3), e00652-17.

Yu, J., **Chen, Z.**, Wang, K., Tezal, M. (2019). Suggestion of the confidence interval of the Cronbach alpha in application to complex survey data. *Survey Methodology*, 45(3), 465-484.

Submitted:

He, Z., **Chen, Z.**, Wu, Y and Gaile, D. P. (2019). A novel and quick method to power pilot studies for the comparison of assay platforms and under controlled specificity. (submitted)

Chen, Z., Koestler, D., Miecznikowski, J. C., Ren, X. Gaile, D. P. (2019). A novel exact method for Top-K family-wise error control for small sample non-small scale inference. (submitted)

In Progress:

Chen, Z., Wilding, G. E. (2020). Unconditional exact tests based on multiple comparisons for Poisson rate.

Chen, Z., Wilding, G. E. (2020). ROC analysis for multistage diagnostic testing procedures in the presence of intermediate results.

PRESENTATIONS

Paper presentation, Joint Statistical Meetings, Denver, CO

Jul 2019

- ROC analysis for multistage diagnostic testing procedures in the presence of intermediate results.

Poster and oral, Sixth Annual Conference of The Upstate Chapter of American Statistical Association, Rochester, NY

Apr 2017

- On The Strong Control of Top-K Error Rates.

Poster, Perry Poster Day, University at Buffalo, Buffalo, NY

Apr 2016

- A Novel Exact Method for Top-K Family-wise Error Control.