Ziqiao Wang – Curriculum Vitae

Contact

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Education

The University of Texas MD Anderson UTHealth Graduate School (GSBS), Houston, TX   2016–2021
 Ph.D. in Biostatistics   GPA 4.0/4.0

Stony Brook University, Stony Brook, NY   2013-2015
 B. S., Double Majored in Mathematics **&** Applied Mathematics and Statistics   GPA 3.9/4.0
 Honors: Summa Cum Laude

Nanjing University, Nanjing, China   2011-2013
 B.S. in Mathematics   GPA 4.1/5.0

Professional Experience

Johns Hopkins University

- Postdoctoral Fellow, Department of Biostatistics   9/2022-

Eli Lilly and Company

- Advisor - Statistics, Statistics Data and Analytics, Early Phase Immunology   3/2022-8/2022
- Research Scientist - Statistician, Statistics Data and Analytics, Early Phase Immunology   6/2021-3/2022

The University of Texas MD Anderson Cancer Center

- Graduate Research Assistant, Department of Biostatistics   2016-2021
- Graduate Research Assistant, Department of Pathology   2018-2021
- Teaching Assistant for Modern Nonparametrics, Department of Biostatistics   Spring 2021

Merck & Co.

Summer Intern, Methodology Group, Biostatistics and Research Decision Science (BARDS)   5/2018-8/2018

Stony Brook University

- Teaching Assistant for Finite Mathematics Structures, Department of Applied Mathematics and Statistics
    Fall 2015
- Teaching Assistant for Data Analysis, Department of Applied Mathematics and Statistics   Summer 2015
- Peer tutor for college-level math courses, Department of Mathematics   Fall 2014-Spring 2015

Honors & Awards

- 2024-2029 K99/R00 NIH Pathway to Independence Award   NIH NHGRI
- 2024-2025 Trans-Omics for Precision Medicine (TOPMed) Fellow (Award amount: $76,650)   NIH NHLBI
- 2020-2021 Andrew Sowell-Wade Huggins Scholarship in Cancer Research   GSBS
- 2020 Student Virtual Conference Award (*The American Society of Human Genetics* (**ASHG**) 2020)   GSBS
- 2020 Student Travel Award (*the Joint Statistical Meetings* (**JSM**) 2020)   QS program, GSBS
- 2019 Scholarship for Summer Institute in Statistical Genetics   University of Washington
- 2019 Student Travel Award (**JSM** 2019)   GSBS
- 2019 Student Travel Award (The International Biometric Society Eastern North American Region (**ENAR**) 2019 Spring Meeting)   QS program, GSBS
- 2018 Student Travel Award (Conference on Statistical Learning and Data Science / Nonparametric Statistics (**SLDS**) at Columbia University)   QS program, GSBS
- 2015 Outstanding Academic Achievement Award   Stony Brook University
- 2013-2015 Dean's List (Every Semester)   Stony Brook University
- 2013 Outstanding Leadership Honor   Nanjing University
- 2012 Ren Min Scholarship Academic Award   Nanjing University
- 2012 Scholarship for Outstanding Performance in Extra Curriculum Activities   Nanjing University

Peer Reviewed Publications

∗: Co-first Author.

1. **Wang, Z.**, Shi, W., Carroll, R., Chatterjee, N. (2024). Joint Modeling of Gene-Environment Correlations and Interactions Using Polygenic Risk Scores in Case-Control Studies. *American Journal of Epidemiology*. doi: 10.1093/aje/kwae081.

2. **Wang, Z.**, Lu, Y., Fornage, M., Jiao, L., Li, D., Wei, P. (2022). Identification of novel susceptibility methylation loci for pancreatic cancer in a two-phase epigenome-wide association study, *Epigenetics*. doi: 10.1080/15592294.2022.2026591.
   (*Platform presentation, IGES 2019*).

3. **Wang, Z.**, Wei, P. (2020). IMIX: A Multivariate Mixture Model Approach to Association Analysis Through Multi-Omics Data Integration, *Bioinformatics*. doi:10.1093/bioinformatics/btaa1001.
   (*Platform presentation, ASHG 2020*).

4. **Wang, Z.**, Chen, J. (2020). Testing for Trend in Benefit-Risk Analysis with Prioritized Multiple Outcomes. *Statistics in Biopharmaceutical Research*. doi: 10.1080/19466315.2019.1690037.

5. Xu, Y.*, **Wang, Z.***(*co-first author*), Wei, P., Gairola, R., Kelsey, K., Sikora, A., Li, G., Gu, J. (2022). Hypermethylation of nc886 in HPV-positive oropharyngeal cancer and its clinical implications: an epigenome DNA methylation profiling study. *Molecular Therapy-Nucleic Acids*, 30:596-605. doi:10.1016/j.omtn.2022.11.012.

6. Bondaruk, J.*, Jaksik, R.*, **Wang, Z.***(*co-first author*), Cogdell, D.*, Lee, S., Chen, Y., Dinh, K., Majewski, T., Zhang, L., Cao, S., Yao, H., Weinstein, J., Navai, N., Dinney, C., Gao, J., Theodorescu, D., Logothetis, C., Guo, C., Wang, W., McConkey, D., Wei, P., Kimmel, M., Czerniak, B. (2022). The origin of bladder cancer from mucosal field effects. *iScience*. doi: 10.1016/j.isci.2022.104551.

7. Yang, G.*, Bondaruk, J.*, Cogdell, D.*, **Wang, Z.***(*co-first author*), Lee, S., Lee, J., Zhang, S., Choi, W., Wang, Y., Liang, Y., Wang, G., Wang, Y., Yao, H., Dadhania, V., Logothetis, C., Siefker-Radtke, A., Kamat, A., Dinney, C., Theodorescu, D., Kimmel, M., Wei, P., Guo, C., Weinstein, J., McConkey, D., Czerniak, B. (2020). Urothelial-to-Neural Plasticity Drives Progression to Small Cell Bladder Cancer. *iScience*. doi: 10.1016/j.isci.2020.101201.

8. Zhang, M., Ru, M., Zhang, J., **Wang, Z.**, et al. (2024). Alcohol consumption does not modify the polygenic risk score-based genetic risk of breast cancer in postmenopausal women: Atherosclerosis Risk in Communities study. *Cancer Prevention Research*. doi: 10.1158/1940-6207.CAPR-24-0208.

9. Dzaye, O., ..., **Wang, Z.**, Chatterjee, N., Matsushita, K., Blaha, M. (2024). Polygenic Risk Scores and Extreme Coronary Artery Calcium Phenotypes in Adults > 75 Years Old: The Atherosclerosis Risk in Communities Study. *Circulation: Cardiovascular Imaging* 17.11 (2024): e017531. doi: 10.1161/CIRCIMAGING.123.016377.

10. Lee, S., Bondaruk, J., Wang, Y., Chen, H., Lee, J., Majewski, T., Mullen, R., Cogdell, D., Chen, J., **Wang, Z.**, Yao, H., Kus, P., Jeong, J., Lee, I., Choi, W., Navai, N., Guo, C., Dinney, C., Baggerly, K., Mendelsohn, C., McConkey, D., Behringer, R., Kimmel, K., Wei, P., Czerniak, B. (2024). Loss of LPAR6 and CAB39L dysregulates the basal-to-luminal urothelial differentiation program, contributing to bladder carcinogenesis. *Cell reports.* doi: 10.1016/j.celrep.2024.114146.

11. Guo, C. C., Bondaruk, J., Yao, H., **Wang, Z.**, Zhang, L., Lee, S., Lee, J. G., Cogdell, D., Zhang, M., Yang, G., Dadhania, V., Choi, W., Wei, P., Gao, J., Theodorescu, D., Logothetis, C., Dinney, C., Kimmel, M., Weinstein, J. N., McConkey, D. J., ... Czerniak, B. (2020). Assessment of Luminal and Basal Phenotypes in Bladder Cancer. *Scientific reports*, 10(1), 9743. doi:10.1038/s41598-020-66747-7.

12. Guo, C., Majewski, T., Zhang, L., Yao, H., Bondaruk, J., Wang, Y., Zhang, S., **Wang, Z.**, Lee, J., Lee, S., Cogdell, D., Zhang, M., Wei, P., Grossman, H., Kamat, A., Duplisea, J., Ferguson, J., Huang, H., Dadhania, V., Dinney, C., Weinstein, J., Baggerly, K., McConkey, D., Czerniak, B. (2019). Dysregulation of EMT Drives the Progression to Clinically Aggressive Sarcomatoid Bladder Cancer. *Cell reports*, 27(6), 1781–1793.e4. doi:10.1016/j.celrep.2019.04.048.

13. Zhang, Y., Sturgis, E.M., Wei, P., Liu, H., **Wang, Z.**, Ma, Y., Liu, C., Gu, K.J., Wei, Q., and Li, G. (2019). A Genetic Variant within MDM4 3'UTR miRNA Binding Site is Associated with HPV16-positive Tumors and Survival of Oropharyngeal Cancer. *Molecular carcinogenesis*, 58(12), 2276–2285. doi:10.1002/mc.23116.

14. Yang, Z., Zhang, Y., Wang, X., Huang, J., Guo, W., Wei, P., Li, G., **Wang, Z.**, Huang, Z., and Zhang, L. (2019). Putative Biomarkers of Malignant Transformation of Sinonasal Inverted Papilloma into Squamous Cell Carcinoma. *The Journal of international medical research*, 47(6), 2371–2380. doi:10.1177/0300060519838385.

Preprints and Papers under Review

1. **Wang, Z.**, Czerniak, B., Wei, P. (2023). Spatial IMIX: A Mixture Model Approach to Spatially Correlated Multi-Omics Data Integration. *Preprint.* doi:10.1101/2023.07.15.549148.

2. **Wang, Z.**, Grosvenor, L., Ray, D., Ruczinski, I., Beaty, T., Volk, H., Ladd-Acosta, C., Chatterjee, N. (2024+). Estimation of Direct and Indirect Polygenic Effects and Gene-Environment Interactions using Polygenic Scores in Case-Parent Trio Studies. *Preprint.* doi: 10.1101/2024.10.08.24315066.

3. Mathews, L., Hu, X., **Wang, Z.**, et al. (2024+). MMP7 as a Causal Factor for Incident Hypertension and its Complications Proteomic Analysis from the Atherosclerosis Risk in Communities (ARIC) Study and Mendelian Randomization Study.

4. Alonso, D., Patel, D., Schmitz, C., **Wang, Z.**, Knorr, J., Manner, D., Datta, G., Koester, A., Na, S., Klekotka, P., Nirula, A. (2024+). Safety and efficacy of LY3454738, a novel CD200R checkpoint agonist, in a 12-week phase 1 study in atopic dermatitis.

Work in Progress

1. **Wang, Z.**, Burk, V., Chatterjee, N., Platz, E. (2024+). Discovering plasma proteins associated with common solid cancer incidence in ARIC.

2. **Wang, Z.**, Dutta, D., Chatterjee, N. (2024+). Canonical correlation analysis and partitioned polygenic scores reveal interesting biological pathways in diseases.

3. Johansen, M., Chen, J.,**Wang, Z.**, et al. (2024+). Proteomics and the Risk of Incident Embolic and Thrombotic Stroke.

4. Syleouni, M., Joshu, C., ..., **Wang, Z.**, Chatterjee, N., Rohrmann, S., Platz, E. (2024+). Discovering plasma proteins associated with breast cancer incidence in post-menopausal women in the prospective ARIC study.

Presentations

Invited Talks

1. "Estimation of Direct and Indirect Polygenic Effects and Gene-Environment Interactions using Polygenic Scores in Case-Parent Trio Studies." Department of Biostatistics, Johns Hopkins University. Baltimore, MD. September 23, 2024.

2. "Novel Methods for Estimating Risk Parameters Associated with Polygenic Scores using Case-Parent Trio Designs." STATGEN 2024: Conference on Statistics in Genomics and Genetics. The American Statistical Association (ASA) Section on Statistics in Genomics and Genetics (SSGG). Pittsburgh, PA. May 1-3, 2024.

3. "Risk Parameter Estimation under Polygenic Models from Case-Parent Trios and Applications on GEARs Study." GEARs seminar, Department of Epidemiology, Johns Hopkins University, Baltimore, MD. April 26, 2023.

4. "IMIX: A multivariate mixture model framework for integrative analysis of multiple types of omics data." Discovery statisticians and biologists meeting, Eli Lilly and Company, Indianapolis, IN. August 5, 2022.

5. "Testing for Trend in Benefit-Risk Analysis with Prioritized Multiple Outcomes." Dr. Anil Korkut's lab, Department of Bioinformatics & Computational Biology, The University of Texas MD Anderson Cancer Center, Houston, TX. October 12, 2018.

6. "Internship Experience at Merck." GSBS Quantitative Sciences Department Retreat, NASA Space Center, Houston, TX. September 29, 2018.

Selected Conference Talks

1. "Estimation of Direct and Indirect Polygenic Effects and Gene-Environment Interactions using Polygenic Scores in Case-Parent Trio Studies." PRIMED Consortium Fall 2024 Meeting. Bethesda, MD. October 24-25, 2024.

2. "Joint Modeling of Gene-Environment Correlations and Interactions Using Polygenic Risk Scores in Case-Control Studies." **ENAR** 2024, Baltimore, MD. March 10-13, 2024.

3. "Spatial IMIX: A Mixture Model Approach to Spatially Correlated Multi-Omics Data Integration." The Joint Statistical Meetings (**JSM**) 2022, Washington, D.C. August 7-11, 2022.

4. "IMIX: A multivariate mixture model framework for integrative analysis of multiple types of omics data." **Platform Presentation** (top 8% submitted abstracts selected), The American Society of Human Genetics (**ASHG**) 2020 Annual Meeting, virtual. October 27-30, 2020.

5. "IMIX: A multivariate mixture model framework for integrative analysis of multiple types of omics data." The Joint Statistical Meetings (**JSM**) 2020, virtual. August 2-6, 2020.

6. "A two-stage epigenome-wide association study identifies novel pancreatic cancer susceptibility loci by leveraging public controls." **Platform Presentation**, International Genetic Epidemiology Society (**IGES**) Annual Meeting, Houston, TX. October 12 - 14, 2019.

7. "Statistical methods for leveraging public controls in a two-stage epigenome-wide association study." The Joint Statistical Meetings (**JSM**) 2019, Denver, CO. July 27 - August 1, 2019.

8. "Robust and Powerful New Methods for Trend Testing of Composite Endpoints in Benefit-Risk Analysis." Merck & Co., Bards, Upper Gwynedd, PA. August 9, 2018.

Selected Conference Posters

1. "Novel Methods for Estimating Risk Parameters Associated with Polygenic Scores using Case-Parent Trio Designs." The American Society of Human Genetics (**ASHG**) 2023 Annual Meeting, Washington, D.C. November 1-5, 2023.

2. "A multivariate mixture model approach to integrative analysis of omics data." Integrative Biostatistics Research for Imaging, Genomics, & High-throughput Technologies in Precision Medicine (**iBRIGHT**) 2019, Houston, TX. November 11 - 13, 2019.

3. "A two-stage epigenome wide association study identifies novel pancreatic cancer susceptibility loci by leveraging public controls." The American Society of Human Genetics (**ASHG**) 2019 Annual Meeting, Houston, TX. October 15 - 19, 2019.

4. "Statistical Methods for Leveraging Public Controls in Epigenome-Wide Association Study." Conference on Statistical Learning and Data Science / Nonparametric Statistics (**SLDS**), Columbia University, New York, NY. June 4 - 6, 2018.

Academic Service and Outreach

1. Peer Reviewer: BMC Bioinformatics, Scientific Reports, Statistics in Biopharmaceutical Research, Statistics in Biosciences, Wires Computational Statistics, Human Genetics and Genomics (HGG) Advances

2. Organizer and Chair, Invited Session: Advanced Methods for Data Integration in Genetics. STATGEN 2025: Conference on Statistics in Genomics and Genetics. ASA SSGG. 05/2025

3. Awards Committee, ENAR of International Biometrics Society 08/2024 - present

4. Co-founder, ASA Postdoctoral Association (ASAPA) 01/2024 - present

5. Organizer, Bi-Weekly Statistical Genetics Working Group Seminar Series, Department of Biostatistics, Johns Hopkins University 01/2023 - present

6. Executive Member, Council of Emerging and New Statisticians (CENS), ENAR of International Biometrics Society 05/2022 - 05/2024

7. Organizer, Invited Session: Paths to Success: Perspectives from Recruiters and Hiring Committees with a Focus on Fostering Inclusivity and Equity, ENAR 2024 03/2024

8. Session Chair, Contributed Papers: Genome Wide Association Studies, ENAR 2024 03/11/2024

9. Session Chair, Contributed Papers: High Dimensional Data Analysis, ENAR 2021 03/15/2021

10. Peer Mentor, GSBS Peer Mentoring Program for First-Year GSBS PhD student 08/2019 - 05/2021

11. Student Representative, GSBS Standing Committees Curriculum Committee 10/2019 - 09/2020

12. Committee Member, GSBS Quantitative Sciences Program Curriculum Committee 09/2018 - 08/2020

13. Member, Stony Brook University Environmental Club 2014-2015

14. Volunteer Teacher, AIESEC, Dare To Dream Volunteer Teaching Program, An Hui, China Summer 2014

Research Mentorship

- Charissa Luk, Undergraduate Student in Biomedical Engineering at JHU. July 2023 - present. Project: Data preprocessing and normalization of OLINK proteomics data in the UK Biobank.

- Bipasha Akhter and Radiah Azmyne Khan, Visiting Scholars from Bangladesh. Aug 2024 - present. Project: Genome-wide association studies on negative birth outcomes in South Asian women.

- Tryggvi McDonald, First-year PhD student from the Department of Human Genetics and Genomics, JHU. Jan - April, 2024. Project: Identifying incident cardiovascular disease and type 2 diabetes patients and PGSxE interactions in the UK Biobank.

- Sameer Gabbita, Undergraduate Student in Biomedical Engineering at JHU. Summer Internship, Bloomberg Distinguished Professor (BDP) Summer Research Program at JHU, 2024. Project: Mendelian Randomization to identify the causal relationship between LDL and Breast Cancer Survival Rates.

- Zoey Hall, Undergraduate Student from Howard University, major in Computer Science, minor in Maternal & Child health. Vivien Thomas Scholars Initiative Summer Fellow, Diversity Summer Internship Program at Johns Hopkins Bloomberg School of Public Health, 2024. Project: Polygenic scores of Attention-deficit/hyperactivity disorder (ADHD) in the Black population from the Boston Birth Cohort.

Funding Support

Current:

- 1K99HG013674-01 (Wang, Ziqiao) 09/19/2024 - 08/30/2026
  National Human Genome Research Institute/NIH
  Enhancing the Interpretability and Applicability of Polygenic Scores through Multi-Omics Integration and Analysis of Family-Based Studies
  Role: Principal Investigator

Completed:

- 1R01HG010480 (Chatterjee, Nilanjan)   05/01/2019 – 09/15/2024
  National Human Genome Research Institute/NIH
  Robust Methods for Polygenic Analysis to Inform Disease Etiology and Enhance Risk Prediction
  Role: Postdoctoral Fellow

- 5R01HL116720 (Wei, Peng)   8/24/2017-7/31/2023
  National Heart, Lung, and Blood Institute/NIH
  Association Analysis of Rare Variants with Sequencing Data
  Role: Graduate Research Assistant

- 5R01CA169122 (Wei, Peng)   4/20/2017-5/31/2019
  National Cancer Institute/NIH
  Genetic Susceptibility and Risk Model for Pancreatic Cancer
  Role: Graduate Research Assistant

- 600657-30-120536-28 (Wei, Peng)   12/1/2017-11/30/2019
  The University of Texas MD Anderson Cancer Center/IRG
  Epigenetic Susceptibility to Pancreatic Cancer
  Role: Graduate Research Assistant

Short Courses and Workshops

1. "Everything you ever wanted to know about applying for NIH Grants (but were afraid to ask)." NHGRI's ASHG workshop. Washington, DC. Nov, 2023.

2. "Guidance and Strategies for Crafting Competitive Funding Proposals." JHU. Oct 19, 2023.

3. "Writing a Winning NIH Grant Proposal Seminar." JHU. Oct 9, 2023.

4. "Mendelian randomisation." Short course, Imperial College London. May 22-23, 2023.

5. "Integrative clustering of multiple genomic data types using a joint latent variable model with application to breast and lung cancer subtype analysis." **Presentation** at Statistical Genetics Journal Club, Houston, TX. April 8, 2020.

6. "Mini Course: Academic Writing Style." Writing Support Services, UTHealth SPH, Houston, TX. March 2 - 23, 2020.

7. "PubMed for Advanced Searchers." Scientific Publications, Research Medical Library, MD Anderson Cancer Center, Houston, TX. November 14, 2019.

8. "Interactive Invited Workshop: The Michigan Imputation Server: Data Preparation, Genotype Imputation, and Data Analysis." **ASHG**, Houston, TX. October 17, 2019.

9. "Education Workshop on Mendelian Randomization." **IGES**, Houston, TX. October 12, 2019.

10. "Conversation 2: Workplace Conversations." Scientific Publications, Research Medical Library, MD Anderson Cancer Center, Houston, TX. October 2 - November 20, 2019.

11. "Making Presentations." Scientific Publications, Research Medical Library, MD Anderson Cancer Center, Houston, TX. July 29 - September 16, 2019.

12. "Literature Reviews." Scientific Publications, Research Medical Library, MD Anderson Cancer Center, Houston, TX. September 26, 2019.

13. "PubMed: The Basics." Scientific Publications, Research Medical Library, MD Anderson Cancer Center, Houston, TX. September 24, 2019.

14. "Mixed Models in Quantitative Genetics"; "Association Mapping: GWAS and Sequencing Data"; "Computational Pipeline for WGS Data." University of Washington, Seattle, WA. July 17 - 26, 2019. (**Scholarship**).

15. "Writing 2: Making your sentences flow." Scientific Publications, Research Medical Library, MD Anderson Cancer Center, Houston, TX. May 13 - June 27, 2019.

16. "An Introduction to Causal Effect Estimation with Examples Using SAS Software." **ENAR**, Philadelphia, PA. March 25 , 2019. (**Travel award**).

17. "Writing 1: Writing coherent paragraphs." Scientific Publications, Research Medical Library, MD Anderson Cancer Center, Houston, TX. March 12 - April 25, 2019.

18. "Common SNPs explain a large proportion of the heritability for human height." **Presentation** at Statistical Genetics Journal Club, Houston, TX. February 8, 2019.

19. "Writing 3: Editing your own writing." Scientific Publications, Research Medical Library, MD Anderson Cancer Center, Houston, TX. January 7 - February 21, 2019.

Professional Memberships

- PRIMED Consortium (2024 -)
- American Society of Human Genetics (2019 -)
- International Genetic Epidemiology Society (2019 -)
- International Chinese Statistical Association (2019 -)
- Eastern North American Region (ENAR) / International Biometric Society (2019 -)
- American Statistical Association (2017 -)
- Institute of Mathematical Statistics (2017 -)

Research Interests

- Statistical Genetics
- Cancer Genomics
- Mixture Model
- Machine Learning and Nonparametric Methods

Last updated: December 18, 2024