

CURRICULUM VITAE

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I. Personal Information

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II. Education

Degrees

2004	Ph.D.	Biostatistics, Emory University, Atlanta, Georgia
2004	M.S.	Biostatistics, Emory University, Atlanta, Georgia
1998	B.S.	Applied Mathematics, South China University of Technology, China

Other Education

1999 — 2000	Master program in Applied Statistics, Worcester Polytechnic Institute, Worcester, Massachusetts
1998 — 1999	Master program in Applied Statistics, South China Univ. of Tech., China

III. Employment and Professional Experience

Aug 2022 — present	Professor, Department of Statistics, University of Georgia
July 2016 — July 2022	Associate Professor
November 2011 — June 2016	Assistant Professor Department of Biostatistics, University of Florida
November 2011 — Present	Affiliated Investigator
November 2010 — October 2011	Assistant Member
July 2006 — October 2010	Staff Scientist Fred Hutchinson Cancer Research Center
September 2004 — June 2006	Research Associate Center for Biostatistics in AIDS Research (CBAR), Harvard University.
September 2001 — August 2004	Research Assistant

IV. Scholarship

Publications in peer-reviewed journals

1. Halloran, ME, Longini, IM, Nizam, A and **Yang Y**. Containing Bioterrorist Smallpox. *Science*, 2002; 298: 1428-1432. doi.org/10.1126/science.1074674.
2. Longini IM, Halloran ME, Nizam A and **Yang Y**. Containing Pandemic Influenza with Antiviral Agents. *American Journal of Epidemiology*, 2004; 159: 623-633. doi.org/10.1093/aje/kwh092.
3. Fu, HZ, **Yang Y** and Yao, YX. A Mathematical Model in Ranking Tax Credibility of Entities. *Mathematics in Practice and Theory* (Chinese). 2004; 34: 11-15.
4. **Yang Y**, Longini, IM and Halloran, ME. Design and Evaluation of Prophylactic Intervention Using Infectious Disease Incidence Data from Close Contact Groups. *Journal of the Royal Statistical Society, Series C*. 2006; 55: 317-330. doi.org/10.1111/j.1467-9876.2006.00539.x.
5. Longini IM, Halloran ME, Nizam A, **Yang Y**, Xu S, Burke DS, Cummings DAT and Epstein J. Containing a large bioterrorist smallpox attack: A computer simulation approach. *International Journal of Infectious Disease*. 2007; 11: 98-108. doi.org/10.1016/j.ijid.2006.03.002.
6. Halloran ME, Hayden F, **Yang Y**, Logini IM and Monto A. Antiviral Effects on Influenza Viral Transmission and Pathogenicity: Observations from Household-Based Trials. *American Journal of Epidemiology*. 2007; 165: 212-221. doi.org/10.1093/aje/kwj362.
7. Mulligan K, **Yang Y**, Wininger D, Koletar S, Parker RA, Alston-Smith B, Basar M and Grinspoon S. Effects of Metformin and Rosiglitazone in HIV-Infected Patients with Hyperinsulinemia and Elevated Waist/Hip Ratio. *AIDS*. 2007; 21: 47-57. doi.org/10.1097/QAD.0b013e328011220e.
8. Shikuma CM, **Yang Y**, Meyer WA, Glesby M, Tashima KT, Ribaud H, Webb N, Bastow B, Kuritzkes DR, and Gulick RM. Metabolic Effects of Protease Inhibitor-Sparing Antiretroviral Regimens given as Initial Treatment of HIV-1 Infection (AIDS Clinical Trials Group Study – A5095). *Journal of AIDS*. 2007; 44:540-550. doi.org/10.1097/QAI.0b013e318031d5a0.
9. **Yang Y**, Longini IM and Halloran ME. A Resampling-Based Test to Detect Person-To-Person Transmission of Infectious Diseases. *Annals of Applied Statistics*. 2007; 1:211-228. doi.org/10.1214/07-AOAS105.
10. **Yang Y**, Longini IM and Halloran ME. A Data-Augmentation Method for Infectious Disease Incidence Data from Close Contact Groups. *Computational Statistics and Data Analysis*. 2007, Vol. 51:6582-6595. doi.org/10.1016/j.csda.2007.03.007.

11. **Yang Y**, Halloran ME, Sugimoto J and Longini IM. Detecting Human-To-Human Transmission of Avian A(H5,N1) Influenza. *Emerging Infectious Disease*. 2007; 13:1348-1353. <https://doi.org/10.3201/eid1309.070111>.
12. Livingston EG, Cohn SE, **Yang Y**, Watts DH, Bardeguez AD, Jones TB, Smith LM, Umbleja T and McComsey GA. Lipids & Lactate in HIV-1 Infected Pregnancies With/ Without Protease Inhibitor-Based Therapy. *Obstetrics and Gynecology*. 2007, Vol. 110:391-397. doi.org/10.1097/01.AOG.0000271210.79340.4c.
13. **Yang Y** and DeGruttola V. Resampling-based Multiple Testing Methods with Covariate Adjustment: Application to Investigation of Antiretroviral Drug Susceptibility. *Biometrics*. 2008; 64:329-336. doi.org/10.1111/j.1541-0420.2007.00883.x.
14. **Yang Y**, Gilbert, P, Longini IM and Halloran ME. A Bayesian Framework for Estimating Vaccine Efficacy per Infectious Contact. *Annals of Applied Statistics*. 2008; 2:1409-1431. doi.org/10.1214/08-AOAS193.
15. **Yang Y**, Halloran ME and Longini IM. A Bayesian Model for Evaluating Influenza Antiviral Efficacy in Household Studies with Asymptomatic Infections. *Biostatistics*. 2009; 10: 390-403. doi.org/10.1093/biostatistics/kxn045.
16. **Yang Y**, Sugimoto JD, Halloran ME, Basta NE, Chao DL, Matrajt L, Potter G, Kenah E and Longini IM. The Transmissibility and Control of Pandemic Influenza A (H1N1) Virus. *Science*. 2009; 326: 729-733. doi.org/10.1126/science.1177373.
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18. Sugimoto, JD, Borse, NN, Ta, ML, Stockman, LJ, Fischer, GE, **Yang, Y**, Halloran, ME, Duchin, JS and Longini, IM. The effect of age on transmission of clinical pandemic influenza A (H1N1) during an outbreak in a camp and households in Washington State, United States. *Epidemiology*. 2011; 22(2): 180-187. doi.org/10.1097/EDE.0b013e3182060ca5.
19. Wang Y, Feng Z, **Yang Y**, Self SG, Gao Y, Wakefield J, Wang L, Zhang J, Chen X, Yao L, Stanaway J, Wang Z, Yang W. Hand, Foot and Mouth Disease in China: Patterns of Spread during 2008-2009. *Epidemiology*. 2011; 22: 781-792. doi.org/10.1097/EDE.0b013e318231d67a. (**Corresponding Author**).
20. **Yang Y**, Longini IM, Halloran ME and Obenchain V. A hybrid EM and Monte Carlo EM Algorithm and Its Application to Analysis of Transmission of Infectious Diseases. *Biometrics*. 2012; 68: 1238-1249. doi.org/10.1111/j.1541-0420.2012.01757.x.
21. **Yang Y** and DeGruttola V. Resampling-based Methods for Testing Equality of Covariance/Correlation Matrices. *International Journal of Biostatistics*. 2012; 8(1): Article 13. doi.org/10.1515/1557-4679.1388.
22. Fang, L-Q., Li, X-L., Liu, K., Li, Y-J., Yao, H-W., Liang, S., **Yang, Y.**, Feng, Z-J., Gray, G. C. and Cao, W-C. Mapping spread and risk of avian influenza A (H7N9) in China. *Nature*

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23. Zhou Y-B, Wang Q-X, Liang S, Gong Y-H, Nie S-J, Nan L, Yang A-H, Liang Q, **Yang Y**, Song X-X, Jiang Q-W. HIV-, HCV-, and Co-Infections and Associated Risk Factors among Drug Users in Southwestern China: A Township-Level Ecological Study Incorporating Spatial Regression. *PLoS One*. 2014; doi.org/10.1371/journal.pone.0093157.

24. Tran CH, Sugimoto DJ, Pulliam JRC, Ryan KA, Myers PD, Castleman JB, Doty R, Johnson J, Stringfellow J, Kovacevich N, Brew J, Cheung LL, Caron B, Lipori G, Harle CA, Alexander C, **Yang Y**, Longini MI, Halloran ME, Morris JG and Small PA. School-located influenza vaccination reduces community risk for influenza-like illness emergence care visits. *PLoS One*. 2014; doi.org/10.1371/journal.pone.0114479.

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46. Ma MJ, Liu C, Wu M, Zhao T, Wang G, **Yang Y**, et al. Influenza A(H7N9) Virus Antibody Responses in Survivors 1 Year after Infection, China, 2017. *Emerg Infect Dis*. 2018;24(4):663-672. doi.org/10.3201/eid2404.171995.
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(corresponding author)
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60. Cheng X-J, Tan L-H, Gao Y-Y, **Yang Y**, Schwebel DC and Hu G-Q. A new method to attribute differences in total deaths between groups to population size, age structure and age-specific mortality rate. *PLoS One*. 2019; doi.org/10.1371/journal.pone.0216613.
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mortality during 1990–2017: A global decomposition analysis. *PLoS Medicine*. 2020;17(6):e1003138. doi.org/10.1371/journal.pmed.1003138.

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Wuhan: a retrospective observational study. *Lancet Infectious Diseases*, 2021; 21(5):617-628. [doi.org/10.1016/S1473-3099\(20\)30981-6](https://doi.org/10.1016/S1473-3099(20)30981-6). (co-corresponding author)

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109. Havelaar AH, Brhane M, Ahmed IA, Kediri J, Chen D, Deblais L, ..., **Yang Y**, Yimer G, Mckune S. Unravelling the reservoirs for colonisation of infants with *Campylobacter* spp. in rural Ethiopia: protocol for a longitudinal study during a global pandemic and political tensions. *BMJ Open* 2022;12:e061311. doi: 10.1136/bmjopen-2022-061311.

110. Cicala F, Wang W-C, Wang T-H, Li N-H, Bertino E, Liang F-M, and **Yang Y**. 2021. PURE: A Framework for Analyzing Proximity-based Contact Tracing Protocols. *ACM Comput. Surv.* 2023;55(1): Article 3. <https://doi.org/10.1145/3485131>.

Under Review/Revision

1. Madewell ZJ, Wang L-P, Dean NE, Zhang H-Y, Wang Y-F, Zhang X-A, Liu W, Yang W-Z, Longini IM, Gao GF, Li Z-J, Fang L-Q, **Yang Y**. Interactions among Acute Respiratory Viruses in Urban China, 2009-2019. (Corresponding author)
2. Yu L-J, Li X-L, Wang Y-H, Zhang H-Y, Ruan S-M, Jiang B-G, Xu Q, Sun Y-S, Wang L-P, Liu W, Yang Y, Fang L-Q. Short-term effects of ambient air pollution on influenza incidence in China: A multi-city study. (Co-corresponding author)
3. Li R-T, Cheng X-J, **Yang Y**, Schwebel DC, Ning P-S, Li L, Rao Z-Z, Cheng P-X, Zhao M, Hu G-Q. Global deaths associated with population ageing between 1990 and 2019 based on a new decomposition method.
4. Radhakrishnan N, Liu M-J, Idowu B, Bansari A, Rathi K, Magar S, Mundhra L, Sarmiento J, Ghaffar U, Kattan J, Jones R, George J, **Yang Y**, Southwick F. Comparison of the Clinical Characteristics of SARS-CoV-2 delta (B.1.617.2) and omicron (B.1.1.529) Infected Patients from a Single Hospitalist Service. (Co-senior author)

Work in progress

1. Tsang KL, Zhang AR, Longini IM, Cowling B, Fang LQ, **Yang Y**. Transmission risk of SARS-CoV-2 among close contacts of confirmed cases in Shandong Province, China. Submitted.
2. Liu MJ, Jing QL and **Yang Y**. Adjusting for bias induced by surveillance threshold for institutional outbreak investigation. In preparation.
3. Liu MJ, Li H, Zhang AR, Fang LQ, Liu W, **Yang Y**. Genetic determinants for clinical outcomes in patients of Severe Fever Thrombocytopenia Syndrome. In preparation.
4. Jalali N, Zhang AR, Fang LQ, Li ZJ, **Yang Y**. The role of human mobility in the transmission of COVID-19 in Hubei Province, China. In preparation.
5. Jalali N, Liu MJ, Zhang AR, Fang LQ, **Yang Y**. A competing risks framework for estimating transmissibility of infectious agents with multiple transmission modes. In preparation.
6. Sugimoto J, **Yang Y**, Hosford J, Lauzardo M, Ahmedov S, Cooker RL, Halloran EM, Longini IM and Fennelly K. Re-defining super-spreading in tuberculosis. In preparation.
7. Tsang KL, Halloran ME, Rojas DP, Longini IM, **Yang Y**. Transmissibility of Zika disease and the impact of reporting bias in Colombia. In preparation.
8. **Yang Y**, Zhu Y-F, Li X-L and Fang L-Q. Re-visit SARS: Transmissibility, Natural History, and Seasonality. In preparation.
9. Zhu Y-F, **Yang Y** and Halloran ME. Model Selection in the Presence of High-Dimensional Missing Data with Application to Infectious Disease Data. In preparation.

Book Chapters

Yang Y. (2020) Geographically weighted regression. In X. Chen & D. Chen (Eds.), Statistical Methods for Global Health and Epidemiology: Principles, Methods and Applications (pp. 277-316).

Conference Abstracts

Regular talks

1. Yang Y. and Longini I.M.: New methods for the estimation of influenza antiviral agent efficacy. *Joint Statistical Meeting*, San Francisco, CA, 2003.
2. Yang Y. and Longini I.M.: Estimation of influenza antiviral agent efficacy. *International Biometric Society/Eastern North American Region Spring Meeting*, Tampa, FL, 2003.
3. Mulligan, K., Yang, Y., Wininger, D., Koletar, S., Parker, R.A., Alston-Smith, B., Basar, M. and Grinspoon, S.: Effects of Metformin and Rosiglitazone on Body Composition in HIV-Infected Patients with Hyperinsulinemia and Elevated Waist/Hip Ratio: A Randomized, Placebo Controlled Trial. *Conference on Retroviruses and Opportunistic Infections*, Denver, CO, 2006.
4. Shikuma, C.M., Yang, Y., Meyer, W.A., Glesby, M., Tashima, K.T., Ribaud, H., Webb, N., Bastow, B., Kuritzkes, D.R., and Gulick, R.M.: Metabolic Analyses within A5095: Effect of Efavirenz against an All-Nucleoside/Nucleotide Background. *Conference on Retroviruses and Opportunistic Infections*, Denver, CO, 2006.
5. Yang, Y. and DeGruttola, V.: Resampling-based Multiple Testing Methods with Covariate Adjustment: Application to Investigation of Antiretroviral Drug Susceptibility. *Workshop on Quantitative Methods for Research on Antiviral Resistance*, Boston, MA, 2006
6. Yang, Y., Halloran, M.E., Daniels, M. and Longini, I.M.: Modeling Competing Infectious Pathogens from a Bayesian Perspective: with Application to Influenza Studies with Incomplete Laboratory Results. *MFO workshop: Design and analysis of Infectious Disease Studies*, Mathematisches Forschungsinstitut Oberwolfach, Germany, 2009
7. Yang, Y., Halloran, M.E., Daniels, M. and Longini, I.M.: A Partial Monte carlo EM Algorithm for Analyzing Transmission of Infectious Diseases. *MIDAS Network Meeting*, Atlanta, GA, 2011
8. Yang Y.: Assessing the Transmissibility of TB and Risk Factors in Close Contact Groups. *MIDAS Network Meeting*, Austin, TX, 2013.

Invited talks

1. Yang, Y., Gilbert, P., Longini, I.M. and Halloran, M.E.: A Bayesian Framework for Estimating Vaccine Efficacy per Infectious Contact. *Disease Dynamics 2008*, University of British Columbia, Vancouver, British Columbia, 2008.
2. Yang, Y.: Statistical Analysis of Epidemics with Asymptomatic Infections and/or Co-circulating Pathogens. *US CDC*, Atlanta, GA, 2011.
3. Yang Y.: Transmissibility and control of avian influenza A (H7N9) virus. *Sichuan Provincial CDC*, Sichuan, China, 2013.
4. Yang Y.: Methods for estimating the dengue vaccine efficacy in cohort studies. *Zhongshan Municipal CDC*, Guangdong, China, 2014.
5. Yang Y.: Spatial-temporal modeling for surveillance data of multiple pathogens.

Workshop on modeling hand, foot and mouth disease sponsored by China CDC, Beijing, China, 2015.

6. Yang Y. and Mingjin Liu: Statistical adjustment for reporting bias in surveillance data of infectious diseases. *2018 International Indian Statistics Association (IISA) International Conference on Statistics*, Gainesville, Florida.
7. Yang Y.: Modeling infectious diseases. School of Public Health, Shandong University, Jinan, Shandong Province, China, 2019.
8. Yang Y.: Modeling Transmission of Infectious Diseases. Workshop on Pedestrian Dynamics and Epidemic Modeling. Pensacola, FL 2020.
9. Yang Y.: Household Transmission of SARS-CoV-2: The Role of Presymptomatic and Asymptomatic Infectivity. American Society of Tropical Medicine and Hygiene 69th Annual Meeting, 2020.
10. Household Transmission of COVID-19 and Risk Factors for Susceptibility and Infectivity in Wuhan, the First Epicenter. 2021 ASA Florida Chapter Meeting, COVID-19 Panel Discussion, April 2, 2021.
11. Yang Y.: Inference on Vaccine Efficacies in the Presence of Missing Data. US CDC Statistical Advisory Group Seminar, April 29, 2021.
12. Statistical Adjustment for Reporting Bias in Surveillance Data of Infectious Diseases. Western North American Region (WNAR) of The International Biometric Society (IBS) 2021. June 15, 2021.
13. Understanding transmission dynamics of emerging infectious diseases from contact-tracing data. WebENAR. February 25, 2022.
14. Impacts of Nonpharmaceutical Interventions on Notifiable Infectious Diseases in China during the COVID-19 Pandemic. International Summer Institute for Graduate Students, Central South University, Changsha, China. July 19, 2022
15. Assessing transmissibility and associated risk modifiers of emerging infectious diseases from contact-tracing data. JSM, Washington D.C., August 11, 2022.
16. Interactions among acute respiratory viruses in urban China, 2009-2019. Georgia Statistics Day, Athens, GA, September 1, 2022.

V. Teaching and Advising

Teaching

Institute	Course Number	Course Title	Semester, Year
Univ. of Florida	PHC7066	Large sample theory	Spring, 2012—2015
Univ. of Florida	PHC6937	Applied Survival Analysis	Fall, 2013
Univ. of Florida	PHC6092	Biostatistical Theory	Fall, 2015-2021
Univ. of Georgia	STAT4230/6230	Applied Regression	Fall, 2022
Univ. of Georgia	STAT6820	Statistical Inference	Spring, 2023

Graduate Committee Activities (Selected)

Student	Topic	Home Department	Completion Date (* Expected)	Role
Mingjin Liu	Analytic Methods for Infectious Disease Transmission Dynamics Based On Surveillance and Genetic Data	Biostatistics	2022	Advisor
Neda Jalali	Infectious Disease Modeling with Limited Surveillance Data	Biostatistics	2021	Advisor
Hasibul Hasan	Efficient Sampling in Bayesian Computing for Continuous-Time Infectious Disease Transmission in Close Contact Groups	Biostatistics	2022	Advisor
Ya Meng	HIV partner study and dengue vaccine efficacy estimation	Biostatistics	2015	Advisor
Alex Kirpich	Modeling cholera transmission in Haiti	Biostatistics	2015	Co-Advisor
Yifan Zhu	Model selection in the presence of high-dimensional data	Biostatistics	2015	Co-Advisor
Samson Grebremariam	Joint model of transmission, pathogenicity and immunology for dengue epidemics	Biostatistics	2015	Co-Advisor
Keerati Ponpetch	Environmental Determinants and Spatial Distribution of Schistosomiasis in Ethiopia	Environmental and Global Health	2020	Committee Member
Helena J. Chapman	Latent TB infection in Dominican Republic	Environmental and Global Health	2017	Committee Member
Xueying Tang	Bayesian Data Analysis under Shrinkage Priors	Statistics	2017	Committee Member
Guangyu Zhu	Likelihood-Based Sparse Partial least Squares	Statistics	2016	Committee Member
Mary N. Seraphin	Tuberculosis among Haitian immigrants in Florida	Epidemiology	2016	Committee Member

VI. Services

Professional services

- Editor Board Member, Infectious Medicine, 2021—present
- Academic Editor, PLoS ONE, 2011—present
- Frontiers in Public Health, Co-editor for the special collection on the research topic: “Updates on Clinical and Molecular Epidemiology of Tuberculosis”, 01 Sep 2022—*Present*
- Associate Editor, Biometrics, 2015—2021
- Referee for Biometrics, Biostatistics, AOAS, American Journal of Epidemiology, Epidemics, PLoS Medicine, PLoS Computational Biology, British Medical Journal, Lancet series journals, Nature Communications, Science, Nature, Science Translational Medicine, PNAS, Vaccine
- ENAR Regional Advisory Board, 2021—present
- WHO working group of vaccine trial design for Marburg virus, 2021

Grants review panels

- NIH F18 Fellowships: Epidemiology and Population Sciences review panel, Nov. 3-4, 2022
- NIH F18 Fellowships: Epidemiology and Population Sciences review panel, March 24-25, 2022
- NIH IRAP study section review panel, Jul. 8-9, 2021
- NIH 2021/01 ZAI1 MSA-W (J1) 1 review panel for “Emergency Awards: Rapid Investigation of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) and Coronavirus Disease 2019”, Aug. 27, 2020
- University of Florida Research Opportunity Seed Fund Review - Biomed Basic Committee, 2017
- Wellcome Trust proposal review. 2009

Services at the University of Georgia

- Member, Colloquium Committee, 2022—
- Member, Computing Committee, 2022—
- Member, Faculty Review Committee, 2022—

Services at the University of Florida

- Member, Admission Committee, 2014—present
- Chair, Exam Committee, 2019—present

VII. Grants, Contracts and Awards

Ongoing Grants

- U01 CK000670-01 9/30/2022 – 9/29/2025
 CDC: Building mathematical modeling workforce capacity to support infectious disease and healthcare research
 Role: Contact PI
 - U01 AI169375-01A1 9/2022 – 6/2025
 NIAID: Developing TranStat: A user-friendly R package for the analysis of infectious disease transmission and control among close contacts.
 Role: Contact PI
 - R21 OH012114 (PI: Wu) 9/2/2021 – 8/31/2023
 NIOSH: Occupational Inhalation Exposure to Infectious Respiratory Virus Aerosols in medical Facilities
 Role: Co-Investigator
 - OPP1175487 (PI: Havelaar) 01/2020 - 06/2023
 Bill & Melinda Gates Foundation: Campylobacter Genomics and Environmental Enteric Dysfunction study
 Role: Co-Investigator
 - OPP1205215 (PI: Liang) 11/2018-06/2023
 Bill&Melinda gates Foundation: *Exposure Assessment of Campylobacter infections in Rural Ethiopia*
 Role: Co-Investigator
 - U54 OH011230 (PI: Morris) 9/30/2022 – 9/30/2028
 CDC: *Southeastern and Coastal Center for Agricultural Health and Safety*
 Role: Core Biostatistician

Completed Grants

- R01 AI126357 (PI: Morris) 7/1/2017 – 6/30/2022
 NIH/NIAID: *Cholera transmission and evolution in Port-au-Prince, Haiti*
 Role: Co-Investigator
 - R01 AI139761 (PI: Dean) 07/01/2018-06/30/2023
 NIH/NIAID: *Design and Analysis of Vaccine Trials for Emerging Infectious Disease Threats*
 Role: Co-Investigator
 - R56 AI148284 (PI: Longini) 09/2020 - 08/2021
 NIH: Mathematical and Statistical Methods for the Control of Global Infectious Disease Threats
 Role: Co-Investigator
 - RAPID 2034364 (PI: Li, Ninghui) 7/1/2020 – 6/30/2021
 NSF: *Collaborative: PPSRC: Privacy-Preserving Self-Reporting for COVID-19*
 Role: Co-PI
 - R01 AI120997 (PI: Gordon) 01/2017 - 12/2021
 NIH: Dynamics of Influenza Transmission in Nicaraguan Households
 Role: Co-Investigator
 - R21 AI119773 7/2015 – 6/2017

NIH/NIAID : *Spatial-Temporal Modeling for Surveillance Data of Multiple Pathogens*.
 Role: PI
 ▪ R01 AI116770 (PI: Kenah) 1/2016 - 12/2020
 NIH/NIAID: *Regression, Phylogenetics, and Study Design in Infectious Disease Epidemiology*
 ▪ R37 AI32042-19 Halloran (PI) 4/1992 – 5/2020
 NIH/NIAID: *Methods for Evaluating Vaccine Efficacy*.
 Role: Co-Investigator
 ▪ U54 GM111274 Halloran (PI) 9/2014 – 6/2019
 NIH/NIGMS: *Modeling of Infectious Disease Agent Study Centers for Excellence*.
 Role: Co-Investigator
 ▪ U01 GM070749-07 Halloran Longini (MPI) 5/2004 – 4/2014
 NIH/NIGMS: *Containing Bioterrorist and Emerging Infectious Diseases*
 Role: Co-Investigator
 ▪ R01 A0I51164 07A1 DeGruttola (PI) 12/2008 – 11/2011
 NIH/NIAID: *Methods for Long-Term Follow-Up of HIV-Infected Patients*
 Role: PI of subcontract
 ▪ Corporation-funded research Halloran (PI) 11/2011 – 8/2012
 Medimmune Inc.: *Efficacy of the Live Attenuated Influenza Vaccine in a cohort study*
 Role: PI of subcontract
 ▪ Contract Longini (PI) 7/2014 - 6/2015
 County of Los Angeles, Department of Public Health
Using TranStat for Emerging Infectious Diseases
 Role: Co-Investigator

VIII. Awards and Honors

- 2017 Term Professorship, School of PHHP, Univ. of Florida
- 2012 Best paper of the year, School of PHHP, Univ. of Florida
- 2000 — 2004 Graduate fellowship, Emory University
- 1999 — 2000 Graduate fellowship, Worcester Polytechnic Institute
- 1997 Excellent college student with outstanding scholarship
- 1997 Outstanding college student in Guangdong Province, China
- 1997 P&G Scholarship
- 1997 Honorable mention in the international Mathematical Contest in Modeling (MCM)
- 1996 Excellent college student with first-place scholarship
- 1995 Excellent college student with second-place scholarship