Jonathan Luu

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Education

Harvard University, Graduate School of Arts and Sciences PhD, Biostatistics

University of Southern California, Keck School of Medicine MS, Biostatistics

University of Southern California, Viterbi School of Engineering BS, Computer Science & Computer Engineering

Work Experience Principal Biostatistician, Regeneron Pharmaceuticals 2024 Global Development, Inflammation and Immunology Designed and analyzed clinical trials through collaboration with medical directors, clinical scientists, regulatory affairs, statistical programming, data management, medical writing, and others Developed individual protocols and statistical analysis plans and determined appropriate statistical methodology for data analysis Intern, Bristol Myers Squibb 2023 Early Development, Oncology Explored relationship between surrogate endpoints (PFS/ORR) and overall survival in cancer trials Compared correlation coefficients to verify Prentice criteria of surrogate endpoints to justify their use in cancer trials, stratified by treatment and indication type 2023 Research Assistant, Boston College Designed an analysis plan for a multi-period cluster-randomized crossover trial to test the effectiveness of motivational interviewing on COVID-19 vaccine uptake Teaching Assistant, Harvard T.H. Chan School of Public Health 2020-2023 Served as TA for Applied Survival Analysis, Intro to Data Science, and Survival Methods in Clinical Research Consultant, Harvard Biostatistics Consulting Center 2021-2022 Advised Harvard students and post-docs on study design, analysis planning, and programming Taught R and statistics to high school students through lecture, lab, and group projects Research Experience Comparing methods for multiple-period cluster-randomized crossover trials 2024 Compared characteristics of marginal and mixed effects models for CRTs under small sample conditions Measuring performance for costs and healthcare utilization in nursing 2024 Proposed novel metrics for comparing hospitalization costs and profiling nursing homes Hierarchical models for longitudinal clustered semi-continuous data subject to mortality 2023 Developed a Bayesian discrete-time model to efficiently analyze cost data in nursing homes Determinants of social risk screening and response equity in community health centers 2022 Characterized missing data in electronic health records for Caring Health Center Prevalence, incidence, and reversal pattern of childhood stunting from birth to age 2 years in Ethiopia 2022 Analyzed newborn data with measurement error to determine relationship between malnutrition and stunting

Duration of viral shedding and culture positivity with post-vaccination breakthrough delta variant infections

Compared 3+3 algorithm for phase I RCTs with more sophisticated methods through simulation

A simulation evaluation of the effectiveness and usability of the 3+3 design for phase I clinical trials

Performed survival analysis on COVID-19 viral load/culture data to determine optimal quarantine period

2021

2019