

# Jonathan Luu

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## Education

<b>Harvard University, Graduate School of Arts and Sciences</b> PhD, Biostatistics	Boston, MA May 2024
<b>University of Southern California, Keck School of Medicine</b> MS, Biostatistics	Los Angeles, CA May 2019
<b>University of Southern California, Viterbi School of Engineering</b> BS, Computer Science & Computer Engineering	Los Angeles, CA May 2017

## Research Experience

<b>Expanding the two-part model for clustered semi-continuous data</b> <ul style="list-style-type: none"><li>Developed Bayesian model to efficiently analyze cost and healthcare utilization data in nursing homes</li><li>Created joint metrics to effectively compare nursing homes</li><li>Applied model and metrics to Medicare data consisting of 20 million nursing home residents</li></ul>	2022
<b>Duration of viral shedding and culture positivity with post-vaccination breakthrough delta variant infections</b> <ul style="list-style-type: none"><li>Collected viral load and culture samples from MGH employees who tested positive for SARS-CoV-2</li><li>Performed survival analysis and a spline predictive analysis on the data</li></ul>	2021
<b>Estimating the treatment effect in randomized trials with correlated time-to-event outcomes</b> <ul style="list-style-type: none"><li>Simulated and compared three analysis methods for cluster randomized clinical trials</li></ul>	2020
<b>LOFT-HF sample size re-estimation</b> <ul style="list-style-type: none"><li>Re-estimated sample size for the LOFT-HF trial using blinded aggregate data</li><li>Ran simulation sensitivity analyses for type-I error using sample size, power, accrual rate, and follow-up</li></ul>	2020
<b>A phase I/II study of E7389 Halichondrin B analog in metastatic urothelial tract cancer and renal insufficiency</b> <ul style="list-style-type: none"><li>Produced Kaplan-Meier plots, response and toxicity tables, and baseline statistics for DSMC report</li><li>Analyzed data using multivariate Cox regression for progression-free and overall survival</li></ul>	2019
<b>A simulation evaluation of the effectiveness and usability of the 3+3 design for phase I clinical trials</b> <ul style="list-style-type: none"><li>Compared the 3+3 algorithm for phase I RCTs with more sophisticated methods through simulation</li></ul>	2019
<b>Expanding access to home-based palliative care: a randomized controlled trial protocol</b> <ul style="list-style-type: none"><li>Initiated and monitored REDCap database to collect data for the trial</li><li>Summarized demographics, ineligibility criteria, and patient concerns to present to funding agencies</li></ul>	2018
<b>Deep-web polar insights search engine</b> <ul style="list-style-type: none"><li>Assembled search engine that crawled the deep web for polar-related research data</li><li>Built branded website with inbuilt data visualization capabilities using Banana and D3.js libraries</li></ul>	2016

## Teaching Experience

<b>Teaching Assistant</b> <ul style="list-style-type: none"><li><b>Applied Survival Analysis (BST223)</b></li><li><b>Intro to Data Science (BST260)</b></li><li><b>Survival Methods in Clinical Research (BST224)</b></li></ul>	2021-2022 2021-2022 2022
<b>Biostatistics Consulting Center</b> <ul style="list-style-type: none"><li>Consulted clients on study design, analysis planning, and programming</li><li>Assisted with research projects, grant submissions, and student dissertations</li></ul>	2021-2022
<b>StatStart</b> <ul style="list-style-type: none"><li>Taught R programming and basic statistics to high school students</li><li>Developed computational and problem solving skills by guiding students through a project</li></ul>	2021-2022

## Skills

**Programming (from most proficient to least):** C++, Java, R, Python, SAS, HTML/CSS, Stata, C, C#, Ruby, Julia, Stan  
**Software:** Microsoft Office, Adobe Suite, AutoHotkey, Terminal, Linux, Bootcamp  
**Typing WPM:** 175  
**Other:** Git/GitHub, LaTeX, Cadence, IT experience