Seth D. Temple

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EDUCATION

PhD, MS, Statistics, University of Washington 09/19 - Present Dissertation Committee: Sharon Browning (head), Elizabeth Thompson, Amy Willis, Kelley Harris NDSEG Fellow, NIH Predoctoral Trainee in Statistical Genetics BS, Mathematics, University of Oregon 09/14 - 06/18Summa cum laude, Phi Beta Kappa, Departmental Honors, Presidential Scholar Thesis Committee: Chris Sinclair, Peter Ralph, Samantha Hopkins **WORK EXPERIENCE** Graduate Student Researcher, Fred Hutchinson Cancer Research Center 06/23 - 12/23Project with Matsen group on evolutionary dynamics of SARS-CoV-2 Developed two new methods for anomaly detection and time series clustering Graduate Student Researcher, University of Washington 09/20 - Present Developing methods to study recent evolution in human populations Studying how genetic relatedness affects Alzheimer's dementia Extending FLARE method for local ancestry inference Instructor of Record, University of Washington BIOST 550 (Sp22): Statistical Genetics I: Mendelian Traits BIOST 581 (W23): Statistical Genetics Journal Club Teaching assistant roles for senior undergraduates (3 terms) Graduate Student Researcher, Los Alamos National Laboratory 06/20 - 09/20Constructed spatiotemporal occupancy models for vector epidemiology Applied machine learning to map geographic distribution of mosquito species Actuarial Assistant, Liberty Mutual Insurance 07/18 - 08/19Performed reserving analyses for the leading global surety Developed SQL/SAS code to query claims database Passed advanced statistics and finance actuarial exams 02/18 - 06/18Research Assistant, University of Oregon Built neural nets in Python to predict punctuation for audio recordings Trained keras models with graphical processing units Actuarial Intern, Liberty Mutual Insurance 06/17 - 07/17

PAPERS

Temple, S.D., Waples, R.K., & Browning, S.R. Recent and strong selective sweeps. Manuscript in progress.

Temple, S.D., Browning, S.R. & Thompson, E.A. A large sample coalescent theory. Manuscript in progress.

Temple, S.D., Manore, C.A. & Kaufeld, K.A. *Bayesian time-varying occupancy model for West Nile virus in Ontario, Canada*. Stoch Environ Res Risk Assess (2022). https://doi.org/10.1007/s00477-022-02257-4

Gorris, M.E., Bartlow, A.W., **Temple, S.D.**, et al. *Updated distribution maps of predominant Culex mosquitoes across the Americas*. Parasites & Vectors 14, 547 (2021). https://doi.org/10.1186/s13071-021-05051-3

Horimoto, Andrea R. V. R., Lisa A. Boyken, Elizabeth E. Blue, Kelsey E. Grinde, Rafael A. Nafikov, Harkirat K. Sohi, Alejandro Q. Nato Jr, et al. 2023. *Admixture Mapping Implicates 13q33.3 as Ancestry-of-Origin Locus for Alzheimer Disease in Hispanic and Latino Populations.* HGG Advances 4 (3): 100207

SOFTWARE: Python, R, Snakemake, cluster computing, Excel, Unix, SQL (experienced); C++, Java (intermediate)