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 Research Technician II; Fred Hutch Cancer Research Center 06/23 - 09/23• Project with Matsen group on evolutionary dynamics of antibody affinity maturation Graduate Research Assistant; University of Washington 09/20 - Present Developing methods to study recent evolution in human populations Studying patterns of identity-by-descent in families affected by Alzheimer's (ADSP FUS) 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 Graduate Research Assistant; Los Alamos National Laboratory 06/20 - 09/20Constructed spatiotemporal occupancy models for vector epidemiology Applied maximum entropy modeling for mosquito species distribution mapping Teaching Assistant; University of Washington 09/19 - 06/20, 07/22Actuarial Assistant; Liberty Mutual Insurance 07/18 - 08/19Performed reserving analyses for the leading global surety Developed SQL/SAS code to query claims database Passed actuarial exams (MAS I, P, and FM) Research Assistant; University of Oregon 02/18 - 06/18Built neural nets in Python to predict punctuation for audio recordings Trained keras models with graphical processing units

PAPERS

Teaching Assistant, Math Tutor; University of Oregon

Actuarial Intern; Liberty Mutual Insurance

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

09/14 - 04/1806/17 - 09/17

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.

SKILLS

Software: Python, R, Excel, Unix, SQL, qsub (experienced); C++, Java, SAS, slurm (intermediate) Languages: English (first), German (moderate fluency), Spanish (limited proficiency)