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EDUCATION

Ph.D. in Ecology and Evolutionary Biology, University of Illinois 2005
Program in Ecology and Evolutionary Biology, Urbana-Champaign, Illinois
B.A. in Biology, Northwestern University, Evanston, Illinois 1995

PROFESSIONAL EXPERIENCE

Professor, Department of Biological Sciences 2023–present
Northern Arizona University, Flagstaff, Arizona
President, International Brucellosis Society 2023–2024
Associate Chair & Director of Graduate Studies, Department of Biological Sciences 2022–2024
Northern Arizona University, Flagstaff, Arizona
Associate Professor, Department of Biological Sciences 2017–2023
Northern Arizona University, Flagstaff, Arizona
Assistant Professor, Department of Molecular, Cellular, and Biomedical Sciences 2014–2017
University of New Hampshire, Durham, New Hampshire
Assistant Research Professor, Department of Biological Sciences 2009–2014
Northern Arizona University, Flagstaff, Arizona
Postdoctoral Research Associate, Center for Microbial Genetics & Genomics 2007–2009
PI: Dr. Paul Keim, Northern Arizona University, Flagstaff, Arizona
Postdoctoral Researcher, Center for Conservation & Evolutionary Genetics 2005–2006
PI: Dr. Rob Fleischer, Smithsonian Institution, Washington, DC
Ornithologist, Exotic Bird Species Importation Database, USGS-BRD, Hawaii 2000
Field Biologist, Endangered Puaiohi Recovery Program, USGS-BRD, Hawaii 1998–2000
Biological Technician, Avian Productivity Research, USFWS-King Salmon, Alaska 1998
Field Technician, Nightingale Reed-warbler Study, USGS-BRD, Saipan, CNMI 1997–1998
Biological Technician, Hawk and Bat Inventory, USFS-Kaibab Nat'l Forest, Arizona 1997
Field Assistant, Hawaiian Honeycreeper Research, USGS-BRD, Hakalau, Hawaii 1997
Biological Technician, Wildlife Inventory, USFS-Tongass Nat'l Forest, Alaska 1996
Field Assistant, Cascades Black-tailed Deer Study, OR Dept. F&W, Oregon 1995–1996
Field Assistant, Fisheries Program, USFS-Tongass Nat'l Forest, Alaska 1995

PUBLICATIONS

Olson, S. H., C. M. Davy, C. L. Lausen, Y. A. Dzal, **J. T. Foster**, W. F. Frick, N. W. Fuller, C. G. Haase, C. R. Hranac, H. E. Kretser, S. Lieberman, C. McCann, M. McClure, C. L. Mormorunni, J. C. Ray, C. Walzer, C. K. R. Willis, and L. P. McGuire. 2025. Facilitating evolutionary rescue from a wildlife health threat with cross-sectoral strategies: a case study on white-nose syndrome. Ecology and Society 30(3):14. doi: 10.5751/ES-16200-300314

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heterogeneities in wildlife health and conservation priorities. Integrative and Comparative Biology. icaf123, doi: 10.1093/icb/icaf123

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Kyriazis, C.C., M. Venkatraman, B. Masuda, C.C. Steiner, L. Cassin-Sackett, L.H. Crampton, A.M. Flanagan, **J.T. Foster**, M.L. Houck, A.C. Misuraca, E.H. Paxton, J.A. Robinson, R.C. Fleischer, O.A. Ryder, M.G. Campana, and A.P. Wilder. 2025. Population genomics of recovery and extinction in Hawaiian honeycreepers. Current Biology 35: 1–12. doi: 10.1016/j.cub.2025.04.078

Grider, J, B. Udell, B. Reichert, **J.T. Foster**, W. Kendall, T. Cheng, and W.F. Frick. A novel method for estimating pathogen presence, prevalence, load, and dynamics at multiple scales. 2025. Scientific Reports 15: 9423. doi: 10.1038/s41598-025-93865-x

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Case, S., H.K. Kawelo, J. Hoh, D. O'Hearn, J.H. Sperry, **J.T. Foster**, D. Drake, J. Vizentin-Bugoni, J.P. Kelley, C. Tarwater. 2024. Effects of fruit novelty on feeding preference in four globally invasive frugivorous birds. Biological Invasions 26: 4179–4198. doi: 10.1007/s10530-024-03436-w

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reveals new infection patterns in Hawai'i. International Journal for Parasitology 54: 123-130 doi: 10.1016/j.ijpara.2023.10.001

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Case, S.B., K. Postelli, D.R. Drake, J. Vizentin-Bugoni, **J.T. Foster**, J.H. Sperry, J.P. Kelley, and C.E. Tarwater. 2022. Introduced galliforms as seed predators and dispersers in Hawaiian forests. Biological Invasions 24: 3083–3097. doi: 10.1007/s10530-022-02830-6

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Yang, Y., **J.T. Foster**, M. Yi, L. Zhan, Y. Zhang, B. Zhou, J. Jiang, and L. Mei. 2021. Phenotypic homogeneity of emetic *Bacillus cereus* isolates in China. Letters in Applied Microbiology 73: 646-651. doi: 10.1111/lam.13527

Bechtel, M.J., K.K. Drake, T.C. Esque, N.C. Nieto, **J.T. Foster**, and M.B. Teglas. 2021. Borreliosis transmission from ticks associated with desert tortoise burrows: examples of tick-borne relapsing

fever in the Mojave Desert. Vector-borne and Zoonotic Diseases 21: 635-637. doi: 10.1089/vbz.2021.0005

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Ridenour, C.L., J. Cocking, S. Poidmore, D. Erickson, B. Brock, M. Valentine, C. Roe, S Young, J. Henke, K.Y Hung, J. Wittie, E. Stefanakos, C. Sumner, M. Rueda, V. Raman, N. Seaton, W. Bendik, H.M. Hornstra O'Neill, K. Sheridan, H. Centner, D. Lemmer, V. Fofanov, K. Smith, J. Will, J. Townsend, **J.T. Foster**, P.S. Keim, D.M. Engelthaler, and C.M. Hepp. 2021. St. Louis Encephalitis Virus in the Southwestern United States: a phylogeographic case for a multi-variant introduction event. Frontiers in Genetics. 12:837 doi: 10.3389/fgene.2021.667895

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Whatmore, A.M. and **J.T. Foster**. 2021. Emerging genetic diversity of *Brucella*: implications for taxonomy, typing and molecular epidemiology. Infection, Genetics and Evolution 92:104865. doi: 10.1016/j.meegid.2021.104865

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Knight, E.C., A-L. Harrison, A.L. Scarpignato, S.L. Van Wilgenburg, E.M. Bayne, J. Ng, E. Angell, R. Bowman, M. Brigham, B. Drolet, W. Easton, T. Forrester, **J.T. Foster**, S. Haché, K. Hannah, J. Ibarzabal, T. Imlay, S. MacKenzie, L. McGuire, G. Newberry, D. Newstead, A. Sidler, D. Swanson, P. Sinclair, J. Stephens, P. Thomas, J. Tremblay, P.P. Marra. 2021. A comprehensive approach to estimating migratory connectivity across the annual cycle reveals peaks in spatial and temporal connectivity that can direct conservation efforts. Ecography. 44:1-15. doi: 10.1111/ecog.05111

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Rouzic, N., L. Desmier, M-E. Cariou, E. Gay, **J.T. Foster**, C.H.D. Williamson, F. Schmitt, M. Le Henaff, A. Le Coz, A. Lorléac'h, J-P. Lavigne, D. O'Callaghan and A. Keriel. 2020. First case of human brucellosis caused by an amphibian-type *Brucella* sp. Clinical Infectious Diseases. doi: 10.1093/cid/ciaa1082

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Foster, J.T., J.M. Scott, and P.W. Sykes. 2000. 'Akikiki (*Oreomystis bairdi*) *In* The Birds of North America, No. 552 (A. Poole and F. Gill, eds.). The Birds of North America Inc., Philadelphia, PA.

OTHER PUBLICATIONS (Book Chapters, Proceedings, etc.)

Foster, J.T., T. Pearson, P. Keim. 2023. Genomic microbiology as applied to animal forensics. The Microbiologist 563. Applied Microbiology International Feature of Forensic Microbiology Issue

Caswell, C., A. Arenas-Gamboa, and **J.T. Foster**. 2023. Chapter 16 *Brucella in* Pathogenesis of Bacterial Infections in Animals: 5th edition. (Prescott, J. F., MacInnes, J. I., Van Immerseel, F., Boyce, J. D., Rycroft, A. N, J. Vázquez-Boland, eds.) Wiley-Blackwell.

Keim, P., J.W. Sahl, T. Pearson, A. Vogler, C.H. Williamson, D. Birdsell, R.T. Okinaka, **J.T. Foster**, D.M. Wagner. 2020. Forensic analysis in bacterial pathogens, Chapter 9. In Microbial Forensics, 3rd edition (B. Budowle, S. A. Morse and S. Schutzer, eds). Elsevier Press. Pp. 123-140. doi: 10.1016/B978-0-12-815379-6.00009-X

Foster, J.T., R.L. Bull, and P. Keim. 2020. Chapter 16. Ricin Forensics: Comparisons to Microbial Forensics. In Microbial Forensics, 3rd edition (B. Budowle, S. A. Morse and S. Schutzer, eds). Elsevier Press. Pp. 241-250. doi: 10.1016/B978-0-12-815379-6.00016-7

Warren, C.C., L.K. Berthold, H.L. Mounce, **J.T. Foster**, L.C. Sackett. 2019. Evaluating the risk of avian disease in reintroducing the endangered Kiwikiu (Maui Parrotbill: *Pseudonestor xanthophrys*) to Nakula NAR, Maui, Hawai'i. Pacific Cooperative Studies Unit Technical Report #201. University of Hawai`i at Mānoa, Department of Botany. Honolulu, HI. 50 pp. PSCU 10125/68734

Roberto, F., J. Arce, L. C. Beck, T. R. Blank, R. Cahall, K. Damer, T. Ficht, **J. Foster**, K. Kiss, M. Nikolich, S. Olsen, R. Ozanich, D. Rozak, F. Schaefer, S. Sozhamannan, R. Tiller, and S. G. Coates. 2019. AOAC SMPR 2016.009:Standard Method Performance Requirements (SMPRs) for DNA-Based Methods of Detecting *Brucella suis* in Field-Deployable, Department of Defense Aerosol Collection Devices. Journal of AOAC International 100:255-260.

Foster, J.T., T. Pearson, P. Keim. 2015. Genomic microbiology as applied to animal forensics. The Microbiologist 16:14-17. September issue Society for Applied Microbiology

Kunz, T.H., **J.T. Foster**, W.F. Frick, A.M. Kilpatrick, G.F. McCracken, M.S. Moore, J.D. Reichard, D. Reeder, and A.H. Robbins. 2011. White-nose syndrome: An overview of ongoing and future research needs. Pp. 189-203. *In* Proceedings of Protection of Threatened Bats at Coal Mines (K.C. Vories A.H. Caswell, and T.M. Price, eds). Office of Surface Mining and Coal Research Center, Southern Illinois University, Carbondale, IL.

Okinaka, R.T., K.K. Hill, T. Pearson, **J.T. Foster**, A. J. Vogler, A. Tuanyok, D. M. Wagner, and P. Keim. 2011. Forensic Analysis in Bacterial Pathogens, Chapter 16. In Microbial Forensics, 2nd Edition. (B. Budowle, S.E. Schutzer, R.G. Breeze, P.S. Keim and S.A. Morse, eds). Elsevier Press. Pp. 259-276.

Foster, J.T., R.L. Bull, and P. Keim. 2011. Ricin Forensics, Chapter 19. In Microbial Forensics, 2nd Edition (B. Budowle, S.E. Schutzer, R.G. Breeze, P.S. Keim and S.A. Morse, eds). Elsevier Press. Pp. 315-326.

Foster, J.T. 2009. The history and impacts of exotic birds in the Hawaiian Islands. Chapter 13 *in* T.K. Pratt, C.T. Atkinson, P.C. Banko, J.D. Jacobi, and B.L. Woodworth, eds. Conservation biology of Hawaiian Forest Birds. Yale University Press, New Haven, CT.

Runde, D. E., W.C. Pitt, and **J.T. Foster**. 2007. Population ecology and potential impacts of emerging populations of exotic parrots. Managing Vertebrate Invasive Species: Proceedings of an International Symposium (G.W. Witmer, W.C. Pitt, K. A. Fagerstone, Eds). USDA/APHIS/WS, National Wildlife Research Center, Fort Collins, CO.

SELECTED FUNDING

Strategic Environmental Research & Development Program (DoD-SERDP) 2025–2028
A predictive model for invasive terrestrial arthropod species in the Indo-Pacific
Role: Co-Investigator, PI: M. Johnson (US Army) & N. Graham (U Hawaii-Hilo), RC25-C1-0201

US Fish & Wildlife Service, C-SWG 2024–2026
Collection of Anianiau from the wild to develop husbandry techniques
Role: Co-Investigator, PI: L. Nietmann (Hawaii DOFAW), F24AS00061

Strategic Environmental Research & Development Program (DoD-SERDP) 2024–2027
Predicting impacts of species loss on ecosystem resilience.
Role: Co-Investigator, PI: J. Sperry (US Army), RC24-4510

US Fish & Wildlife Service 2022–2024
Disease Dynamics of White-Nose Syndrome in the Southwest
Role: PI, F22AS00291

Department of Defense–Defense Threats Reduction Agency 2022–2025
Phylogeography of Brucellosis in Iraq
Role: PI, HDTRA-12210013

Department of Defense-ESTCP 2022–2026
Metabarcoding for monitoring bird species habitat quality on DoD installations
Role: PI, RC22-B5-7440

US Geological Survey–North American Bat Monitoring Program 2021–2024
NABat and One Health: Expanding the North American Bat Monitoring
Program to monitor bat health in the COVID era
Role: Co-Investigator, PI: W. Frick (BCI), G21AC10783

Department of Defense–Defense Threats Reduction Agency 2020–2025
Biosurveillance of Brucellosis in Azerbaijan, Georgia, and Turkey
Role: PI, HDTRA1-20-1-0028

Department of Defense–Defense Threats Reduction Agency 2020–2024
A collaborative interdisciplinary approach to an endemic zoonotic
disease in Cameroon
Role: Co-Investigator, PI: A. Arenas (Texas A&M)

Pacific Southwest Regional Center of Excellence for Vector-Borne Diseases 2020–2021
Tortoises, Ticks, and Tick-borne Pathogens of the Mojave Desert

Training Grant Program from the U.S. CDC (Agreement 1U01CK000516)
Role: PI, Student Fellow: Molly Bechtel (actual lead on award)

Flinn Foundation and NAU VPR's Office 2020–2021
Identifying the geographic link between enzootic cycling of arboviruses and human clinical cases
Role: Co-Investigator, PI: C. Hepp (NAU)

US Fish & Wildlife Service, Competitive State Wildlife Grants (C-SWG) 2020–2024
Improving rodent control to benefit endangered Puaiohi and other species of greatest conservation need in Hawaii
Role: Co-Investigator, PI: L. Berry (Hawaii DOFAW), F19AS00191

National Science Foundation, Ecology & Evolution of Infectious Diseases 2019–2024
Lessons from the past: synthesizing drivers of host persistence…
Role: Co-Investigator, PI: K. Langwig (Virginia Tech), DEB-1911853

Strategic Environmental Research & Development Program (DoD-SERDP) 2019–2021
Predicting impacts of species loss on ecosystem resilience
Role: Co-Investigator, PI: J. Sperry (US Army), RC20-C1-1420

US Fish & Wildlife Service 2019–2020
Investigating survival mechanisms of remnant northern long-eared bat (*Myotis septentrionalis*) populations
Role: Co-Investigator, PI: W. Turner (SUNY Albany), F18AS00119

US Fish & Wildlife Service 2019–2020
Genomics of white-nose syndrome resistance
Role: PI, F18AS00119

National Institutes of Health-NCI, Arizona Cancer Evolution Center 2019–2020
Evolution of TP53 Functional Homologs in Long-lived Vesper Bats
Role: Co-Investigator, PI: M. Tollis (NAU)

National Fish and Wildlife Foundation–Bats for the Future Fund 2018–2019
Breaking the transmission cycle: testing approaches for reducing pathogen contamination on hibernaculum substrates
Role: Co-Investigator, PI: W. Frick (BCI)

Department of Homeland Security-S&T Directorate 2017–2020
Creation of Bacterial Select Agent Repository and Test-Bed for DHS Assays
Role: PI, HSHQDC-17-C-B0021

National Science Foundation, Ecology & Evolution of Infectious Diseases 2017–2022
Predicting the evolution of vector-borne disease dynamics in a changing world
Role: Co-Investigator, PI: D. Fonseca (Rutgers), DEB- 2001213

US Fish & Wildlife Service 2017–2019
Demography of a recovery: tracking the rebound of little brown bat populations
Role: PI, F17AP00588

US Fish & Wildlife Service 2017–2019
Determining survival strategies for multiple bat species affected by WNS
Role: Co-Investigator, PI: K. Langwig (Virginia Tech), FWS-R5-ES-XX

US Fish & Wildlife Service 2017–2018
Genomics of white-nose syndrome resistance
Role: PI, F17AP00589

New England Transportation Consortium 2016–2018
Use of Forested Habitat Adjacent to Highways by Northern Long Ear(ed) Bats
Role: PI, NETC 15-1

The Nature Conservancy & Bat Conservation International 2016–2017
Reducing white-nose syndrome through treatment of the environmental reservoir
Role: PI.

US Department of Agriculture-NRCS-NJ-CIG 2016–2018
Bats as Tools for the Early Detection of Agricultural Pests
Role: Co-Investigator, PI: B. Maslo (Rutgers University)

New Hampshire Agricultural Experiment Station, McIntire-Stennis 2015–2018
Novel molecular identification of forest insect pests
Role: PI, USDA-NH00080-M, 1006866

US Fish & Wildlife Service 2015–2017
Disease dynamics of white-nose syndrome in an endemic region.
Role: Co-Investigator, PI: W. Frick (UC Santa Cruz)

US Fish & Wildlife Service 2015–2017
Field trial of a probiotic to protect bats from white-nose syndrome.
Role: Co-Investigator, PI: M. Kilpatrick (UC Santa Cruz)

Strategic Environmental Research & Development Program (DoD-SERDP) 2014–2018
Seed Dispersal Networks and Novel Ecosystem Functioning in Hawaii
Role: PI, RC-2434

US Fish & Wildlife Service 2014–2016
Genomic differences between *Pseudogymnoascus destructans* and closely related fungi from bat hibernacula: insights into fungal pathogenicity…
Role: PI, #F14AP00644

National Institutes of Health-NIAID 2013–2015
Environmental sampling of *Leptospira* species to link to human leptospirosis cases
Role: Co-Investigator, PI: T. Pearson, 1R15AI101913-01A1

National Park Service 2014–2015
Comprehensive Assessment of Risk of Caves to WNS in U.S. National Parks
Role: Co-Investigator, PI: C. Chambers, P14AC01179

Department of Homeland Security–International S & T 2013
Genomic Sequencing and Signatures in *Clostridium botulinum* Strains from Argentina
Role: PI, DHS-2010-ST-108-000019, Add-on funding to prior award

Technology Research Initiative Fund (TRIF): State of Arizona 2012–2014
Support for Post-Doctoral Associates (SPA) Program:
Genomics of *Clostridium botulinum* for human health
Role: PI

National Institutes of Health-NIAID 2010–2014
NIH-R01, The Role of Plasmids in Rickettsia Biology
Role: Co-Investigator, PI: U. Munderloh (U Minnesota), 5R01AI081690-03

Department of Interior-National Park Service 2012
Genetic Diversity and Population Status of Bats in Northern Arizona
Role: Co-Investigator, PI: C. Chambers, Task #P12AC11177

Technology Research Initiative Fund (TRIF): State of Arizona 2012–2014
Support for Post-Doctoral Associates (SPA) Program: Bridging the Gap
between Renewable Energy & Bat Conservation in Arizona
Role: Co-Investigator, PI: C. Chambers

Technology Research Initiative Fund (TRIF): State of Arizona 2012
Equipment grant: Acquisition of an Illumina MiSeq for DNA Sequencing
Role: Co-Investigator, PI: G. Allan

Bat Conservation International 2012–2013
Evaluating Sensitive Molecular Detection of *Geomyces destructans*
Role: PI

US Fish & Wildlife Service 2012–2014
Antifungal skin microbes as tools for WNS management
Role: Co-Investigator, PI: W. Frick, UC Santa Cruz, FWS-R5-ES-12-001

National Science Foundation- Ecology & Evolution of Infectious Diseases 2011–2016
The effect of sociality on transmission and spread of a multi-host pathogen
Role: Co-Investigator, PI: M. Kilpatrick, UC Santa Cruz, DEB-1336290

Department of Defense- US Army Medical Research & Materiel Command 2011–2013
Telemedicine & Advanced Technology Research Center
Next-Generation Sequencing for Clinical Diagnostics
Role: PI, W81XWH-11-1-0690

Department of Homeland Security–International S & T 2011–2013
Genomic Sequencing and Signatures in *Clostridium botulinum* Strains from Argentina
Role: PI, DHS-2010-ST-108-000019

Department of Homeland Security 2011–2013
Statistical confidence in evidentiary material based on bacterial population genetics
Population genetics of *C. botulinum*, plague, tularemia, anthrax and other pathogens.

Role: Co-Investigator, PI: P. Keim, HSHQDC-10-C-00139, Phase II

Department of Defense- Defense Threats Reduction Agency 2011–2013
Genomics, Proteomics, and Biochemical Profiles of *Bacillus anthracis* and *Yersinia pestis*
Role: NAU PI, Lead PI: H. Kreuzer, PNNL, HDTRA1-08-10-BRCWMD

National Institutes of Health-Institute of Dental & Craniofacial Research 2010–2013
Molecular Ecology of Chronic Rhinosinusitis
Role: PI, Prior PI: C. Liu, transfer of PI on project, R15DE021194-01

Department of Homeland Security 2010–2013
Technology and Data Integration with the Bundeswehr Institute of Microbiology
Role: Co-Investigator, PI: D. Wagner, DHS-2010-ST-108-00015

US Fish & Wildlife Service 2010–2013
DNA-based detection techniques capable of differentiating *Geomyces destructans*
Role: Co-Investigator, PI: D. Lindner, US Forest Service, FWS-R5-ES-10-049

Department of Homeland Security 2010–2013
Standards and analysis tools for next-generation Sequencing Approaches
for Detecting Rare Variants in *Bacillus anthracis* cultures
Role: PI of NAU subcontract, PI: P. Keim, TGen, HSHQDC-10-C-00152

Department of Homeland Security 2010–2012
Bacterial Population Genetics in a Forensics Context: Phase II,
Genetics of plague, brucellosis, tularemia, anthrax and other pathogens.
Role: Co-Investigator, PI: P. Keim, HSHQDC-10-C-00139

US Fish & Wildlife Service 2010–2011
Fine-scale population structure in *Geomyces destructans*, Fungal genetics WNS
Role: PI, FWS-R5-ES-10-049

Department of Defense 2010
Transformational Medical Technologies Initiative Program
Dangerous Pathogen Whole Genome Sequencing
Role: PI, DOD-81661-001-10

USGS-National Wildlife Health Center 2010
Phylogeography of White-Nose Syndrome in Eastern US Bats
Role: PI, USGS-10-ERPA-1000

USGS-Colorado Plateau Field Station 2010
Blood parasites in Southwestern Willow Flycatchers
Role: PI

Department of Homeland Security 2009–2010
Design and validation of pathogen and toxin-specific TaqMan PCR assays
Role: NAU PI, Lead PI: K. Hill (LANL), HSHQPM-09-X-00015

Department of Homeland Security 2008–2009

Bacterial Population Genetics in a Forensics Context: Phase I
Genetics of plague, brucellosis, tularemia, anthrax and other pathogens.
Role: Co-Investigator, PI: P. Keim, HSHQDC-08-C-00158

Department of Homeland Security 2008–2009
Population Genetics of Castor Beans (*Ricinus communis*)
Role: Co-Investigator, PI: P. Jackson

Environment Canada, Science & Technology: STAGE 2009
Molecular ecology and evolution of *Pasteurella multocida* from large-scale avian cholera outbreaks across Canada
Role: Co-Investigator, PI: C. Soos, Environment Canada

Arizona Bat Conservation Partnership 2009
Developing microsatellite markers for spotted bats (*Euderma maculatum*)
Role: Co-Investigator, PI: C. Chambers, Funding cut by State of Arizona

USGS-National Wildlife Health Center 2009
Population Genetics of Avian Cholera from Waterfowl
Role: PI

Research Funding—Environment Canada, Science & Technology 2009
Diagnostic Assays for Avian Cholera in Waterbirds
Role: PI

EPA STAR Fellowship—US Environmental Protection Agency, 3 years 2001–2004
Graduate Fellowship—University of Illinois, Fall 2004 tuition and stipend 2004
Budweiser Conservation Scholarship—National Fish & Wildlife Foundation 2004
Graduate College Dissertation Travel Grant—University of Illinois 2004

PROFESSIONAL AFFILIATIONS

American Ornithological Society
American Society for Microbiology
International Brucellosis Society
The Wildlife Society
Wildlife Disease Association