Trevor Bedford

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Faculty positions

Assistant Member, Fred Hutchinson Cancer Research Center
Vaccine and Infectious Disease Division, Computational Biology Program

Postgraduate training

2012–2013 Newton International Fellow, University of Edinburgh

Institute of Evolutionary Biology

EMBO Fellow, University of Edinburgh

Institute of Evolutionary Biology

2008–2011 **HHMI Associate**, University of Michigan

Dept. of Ecology and Evolutionary Biology

2008 Intern, Wolfram Research Inc.

Education

2003–2008 **PhD in Biology**, Harvard University

1999–2002 **BA in Biological Sciences** (with honors), University of Chicago

Grants, honors & awards

2012–2014 Marie Curie International Incoming Fellowship, European Commission.

2012–2013 Newton International Fellowship, The Royal Society.

2011–2013 Long Term Fellowship, European Molecular Biology Organization.

2005–2008 NSF Graduate Research Fellowship, National Science Foundation.

2003–2005 NIH Genetics and Genomics Training Program, National Institutes of Health.

2001 HHMI Undergraduate Fellowship, Howard Hughes Medical Institute.

1999–2002 University Scholarship, University of Chicago.

Teaching & mentoring

Courses and workshops

2013	Presented phylodynamic inference practical at GenEpi workshop, London School of Hygiene and
	Tropical Medicine.

- Lecturer, Structured Population Models in Epidemiology, École Normale Supérieure.
- Completed an intensive 7-week "Postdoctoral Short-Course on College Teaching in Science and Engineering" at the University of Michigan.
- Teaching fellow, Population Genetics, Harvard University.
- Teaching fellow, Genetics and Genomics, Harvard University.

Supervised trainees

2012	Michael Landis,	Google Summer of	Code student, NESCent.

2010–2011 Daniel Zinder, PhD candidate, University of Michigan.

2009–2011 Ed Baskerville, PhD candidate, University of Michigan.

2007–2008 Rebekah Rogers, PhD candidate, Harvard University.

2004–2005 Jehee Choi, undergraduate, Harvard University.

2004–2005 Michelle White, undergraduate, Harvard University.

Professional service

External PhD examiner: Paul Wikramaratna, Dept. of Zoology, Oxford University.

Ad-hoc referee: American Naturalist, Bioinformatics, Complex Systems, Emerging Infectious Diseases, Genome Biology and Evolution, Genome Research, Journal of Biology, Journal of the Royal Society Interface, Molecular Biology and Evolution, Molecular Ecology, National Science Foundation, Phil Trans R Soc B, PLoS One, PLoS Pathogens, Proc Natl Acad USA, Virology, Virulence.

Publications & talks

JOURNAL ARTICLES

- Volz EM, Koelle K, **Bedford T**. Viral phylodynamics. PLoS Comput Biol 9: e1002947. (PDF)
- Zinder D, **Bedford T**, Gupta S, Pascual M. The roles of competition and mutation in shaping antigenic and genetic diversity in influenza. PLoS Pathog 9: e1003104. (PDF)
- Bedford T, Rambaut A, Pascual M. Canalization of the evolutionary trajectory of the human influenza virus. BMC Biol 10: 38. (PDF)
- Baele G, Lemey P, **Bedford T**, Rambaut A, Suchard MA, Alekseyenko AV. Improving the accuracy of demographic and molecular clock model comparison while accommodating phylogenetic uncertainty. Mol Biol Evol 29: 2157–2167. (PDF)
- Baskerville EB, Dobson AP, **Bedford T**, Allesina S, Anderson TM, Pascual M. Spatial guilds in the Serengeti food web revealed by a Bayesian group model. PLoS Comput Biol 7: e1002321. (PDF)
- Bedford T, Cobey S, Pascual M. Strength and tempo of selection revealed in viral gene genealogies. BMC Evol Biol 11: 220. (PDF)
- Rogers RL, **Bedford T**, Lyons AM, Hartl DL. Adaptive impact of the chimeric gene *Quetzalcoatl* in *Drosophila melanogaster*. Proc Natl Acad Sci USA 107: 10943–10948. (PDF)
- Bedford T, Cobey S, Beerli P, Pascual M. Global migration dynamics underlie evolution and persistence of human influenza A (H₃N₂). PLoS Pathog 6: e1000918. (PDF) (Faculty of 1000 "Must Read")
- Bedford T, Hartl DL. Optimization of gene expression by natural selection. Proc Natl Acad Sci USA 106: 1133–1138. (PDF)
- Rogers RL, **Bedford T**, Hartl DL. Formation and longevity of chimeric and duplicate genes in *Drosophila melanogaster*. Genetics 181: 313–322. (PDF)
- Bedford T, Hartl DL. Overdispersion of the molecular clock: temporal variation of gene-specific substitution rates in *Drosophila*. Mol Biol Evol 25: 1631–1638. (PDF)
- Bedford T, Wapinski I, Hartl DL. Overdispersion of the molecular clock varies between yeast, *Drosophila* and mammals. Genetics 179: 977–984. (PDF)
- Volkman SK, Lozovsky E, Barry AE, **Bedford T**, Bethke L, Myrick A, Day KP, Hartl DL, Wirth DF, Sawyer SA. Genomic heterogeneity in the density of noncoding single-nucleotide and microsatellite polymorphisms in *Plasmodium falciparum*. Gene 387: 1–6. (PDF)
- Castillo-Davis CI, **Bedford TB**, Hartl DL. Accelerated rates of intron gain/loss and protein evolution in duplicate genes in human and mouse malaria parasites. Mol Biol Evol 21: 1422–1427. (PDF)
- Nielsen KM, Kasper J, Choi M, **Bedford T**, Kristiansen K, Wirth DF, Volkman SK, Lozovsky ER, Hartl DL. Gene conversion as a source of nucleotide diversity in *Plasmodium falciparum*. Mol Biol

- Evol 20: 726-734. (PDF)
- Feder ME, **Bedford T**, Albright DR, Michalak P. Evolvability of Hsp70 expression under artificial selection for inducible thermotolerance in independent populations of *Drosophila melanogaster*. Phys Biochem Zool 75: 325–334. (PDF)

Conference abstracts

- 2013 Mathematical and Computational Evolutionary Biology, Montpellier, France.
- 2013 HIV Dynamics and Evolution, Utrecht, Netherlands.
- Society for Molecular Biology and Evolution Annual Meeting, Dublin, Ireland.
- Population Genetics Group, Nottingham, UK.
- Epidemics³, Boston, MA. (Award for Best Oral Presentation by a Young Investigator.)
- Ecology and Evolution of Infectious Diseases, Cornell University, Ithaca, NY.
- Society for Molecular Biology and Evolution Annual Meeting, Barcelona, Spain. Poster presentation.
- Society for Molecular Biology and Evolution Annual Meeting, Halifax, Canada. Poster presentation. (Graduate student poster award)

Invited presentations

- Infectious Disease Dynamics Workshop, Isaac Newton Institute for Mathematical Sciences, Cambridge University, Cambridge, UK.
- Spring Symposium, Centre for Immunity, Infection and Evolution, University of Edinburgh, Edinburgh, UK.
- Institute of Biologie Seminar, École Normale Supérieure, Paris, France.
- Infectious Disease Epidemiology Seminar, School of Public Health, Imperial College, London, UK.
- Ecology and Evolution Seminar, Centre d'Ecologie Fonctionnelle et Evolutive, Université Montpellier, Montpellier, France.
- Institute Seminar, Institute of Medical Virology and Epidemiology of Viral Diseases, Universitäts Klinikum Tübingen, Tübingen, Germany.
- Virology Seminar, MRC National Institute for Medical Research, London, UK.
- Next-Generation Molecular and Evolutionary Epidemiology of Infectious Disease, The Royal Society, Kavli International Centre, Buckinghamshire, UK.
- VIDD Seminar Series, Fred Hutchinson Cancer Research Center, Seattle, WA.
- Scottish Phylogeny Discussion Group, University of Glasgow, Glasgow, UK.

2012	Viral Evolution: Linking Genetics to Epidemics, University of Cologne, Cologne, Germany.
2011	Thematic Seminar, Institute of Biodiversity, Animal Health and Comparative Medicine, University of Glasgow, Glasgow, UK.
2011	RAPIDD Phylodynamics Workshop, NESCent, Durham, NC.
2011	Infectious Disease Epidemiology Seminar, Harvard School of Public Health, Boston, MA.
2010	Departmental Seminar, Department of Ecology and Evolutionary Biology, University of Michigan, Ann Arbor, MI.
2010	Phyloseminar.org, webcast.
2010	Departmental Seminar, Department of Ecology and Evolutionary Biology, University of Michigan, Ann Arbor, MI.
2009	Departmental Seminar, School of Computational Science, Florida State University, Tallahassee, FL.
2006	NKS Wolfram Science Conference, Washington, DC.

Last updated: August 27, 2013