Curriculum Vitae

Tracy A. Heath

October 4, 2014

University of California, Berkeley Department of Integrative Biology 3060 Valley Life Science Building Mail Code 3140 Berkeley, CA 94720-3140 USA

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UPCOMING POSITION

Assistant Professor

Iowa State University

Department of Ecology, Evolution, & Organismal Biology

Ames, IA Starting January 1, 2015

Current Positions

Associate Research Specialist Department of Integrative Biology University of California, Berkeley

Berkeley, CA

December 2010 - December 2014

Postdoctoral Researcher

University of Kansas

Department of Ecology and Evolutionary Biology

Lawrence, KS June 2013 - December 2014

EDUCATION

Ph.D. in Ecology, Evolution, and Behavior

University of Texas

August 2008

B.A. in Biology

Austin, TX

Advisor: David Hillis

Boston University

May 2000

Boston, MA

Research Positions

Postdoctoral Research Fellow

University of Kansas

NSF Fellowship in Biological Informatics

University of California, Berkeley

Supervisors: Mark Holder & John Huelsenbeck

September 2008 - November 2010

Graduate Research Trainee

University of Texas

Integrative Graduate Education and Research Traineeship (IGERT)

Austin, TX

Computational Phylogenetics and Applications to Biology

2003 - 2005 & 2007 - 2008

Visiting Researcher

University of Canterbury

Biomathematics Research Centre

Christchurch, NZ

The Allan Wilson Centre for Molecular Ecology and Evolution

2007

| Graduate Research Assistant | University of Texas |
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| School of Biological Sciences | Austin, TX |
| CIPRES: Modeling and Simulation | 2005 - 2007 |
| Tree Set Visualization: Program Documentation and Web Page | 2004 |
| Tree Set Visualization: Analysis and Visualization of Tree Space | 2003 |
| Genetic Algorithms for Maximum Likelihood Inference (GAML) $$ | 2002 |
| Laboratory Manager | Boston University |
| Department of Biology, DNA Sequencing Facility | Boston, MA |
| Supervisor: Michael Sorenson | 2000 - 2001 |
| Research Assistant | Boston University |
| Department of Biology, Evolutionary Genetics Lab | Boston, MA |
| Mentor: Christopher Schneider | 2000 |

TEACHING

Workshop Instructor

| Wellcome Trust-EMBL-EBI-EMBO Course: Computational Molecular Evolution | 2013 - 2015 |
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| Workshop on Molecular Evolution, Woods Hole, MA | 2012 - 2015 |
| Bodega Applied Phylogenetics Workshop, Bodega Bay, CA | 2011 - 2015 |
| NESCent Academy Course: Phylogenetic Analysis Using RevBayes | 2014 |
| Workshop on Molecular Evolution, Český Krumlov, Czech Republic | 2013 |
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^{*}For teaching materials see: http://treethinkers.org/tracy-heath-workshop-materials/

Teaching Assistant/Laboratory InstructorUniversity of TexasSchool of Biological SciencesAustin, TXAdvanced Systematics (computer lab course)2002 & 2007Vertebrate Natural History (lab/field course)2002Limnology and Oceanography (lab/field course)2001

Publications and Manuscripts

- Cook, J.A., S.V. Edwards, E.A. Lacey, R.P. Guralnick, P.S. Soltis, D.E. Soltis, C.K. Welch, K.C. Bell, K.E. Galbreath, C. Himes, J.M. Allen, T.A. Heath, A.C. Carnival, K.L. Cooper, M. Liu, J. Hanken, and S. Ickert-Bond. 2014. Natural history collections as emerging resources for innovative education. *Bioscience* 64:725–734.
- **Heath, T.A.**, J.P. Huelsenbeck, and T. Stadler. 2014. The fossilized birth-death process for coherent calibration of divergence-time estimates. *Proceedings of the National Academy of Sciences* 111(29):E2957–E2966.
- Höhna, S., **T.A. Heath**, B. Boussau, M.J. Landis, F. Ronquist, and J.P. Huelsenbeck. 2014. Probabilistic graphical model representation in phylogenetics. *Systematic Biology*, 63:753–771
- **Heath, T.A.** and B.R. Moore. 2014. Bayesian inference of species divergence times. In: Chen, M.-H., L. Kuo, and P.O. Lewis (editors). *Bayesian Phylogenetics: Methods, Algorithms, and Applications*. Chapman & Hall/CRC. (invited book chapter; http://www.crcpress.com/product/isbn/9781466500792)
- Stoltzfus, A., H. Lapp, N. Matasci, H. Deus, B. Sidlauskas, C.M. Zmasek, G. Vaidya, E. Pontelli, K. Cranston, R. Vos, C.O. Webb, L.J. Harmon, M. Pirrung, B. O'Meara, M.W. Pennell, S. Mirarab,

- M.S. Rosenberg, J.P. Balhoff, H.M. Bik, **T.A. Heath**, P.E. Midford, J.W. Brown, E.J. McTavish, J. Sukumaran, M. Westneat, M.E. Alfaro, A. Steele, and G. Jordan 2013. Phylotastic! Making tree-of-life knowledge accessible, reusable and convenient. *BMC Bioinformatics*, 14:158.
- Darriba, D., A. Aberer, T. Flouri, **T.A. Heath**, F. Izquierdo-Carrasco, and A. Stamatakis. 2013. Boosting the performance of Bayesian divergence time estimation with the Phylogenetic Likelihood Library. *IEEE 27th International Symposium on Parallel & Distributed Processing* doi:10.1109/IPDPSW.2013.267. (peer-reviewed conference proceedings)
- **Heath, T.A.** 2012. A hierarchical Bayesian model for calibrating estimates of species divergence times. *Systematic Biology*, 61:793-809.
- **Heath, T.A.**, M.T. Holder, and J.P. Huelsenbeck. 2012. A Dirichlet process prior for estimating lineage-specific substitution rates. *Molecular Biology and Evolution* 29:939-955.
- **Heath, T.A.**, S.M. Hedtke, and D.M. Hillis. 2008. Taxon sampling and the accuracy of phylogenetic analyses. *Journal of Systematics and Evolution* 46:239-257.
- **Heath, T.A.**, D.J. Zwickl, J. Kim, and D.M. Hillis. 2008. Taxon sampling affects inferences of macroevolutionary processes from phylogenetic trees. *Systematic Biology* 57:160-166.
- Hillis, D.M., **T.A. Heath**, and K. St. John. 2005. Analysis and visualization of tree space. *Systematic Biology* 54(3):471-482. (cover article)
- Wilcox, T.P., D.J. Zwickl, **T.A. Heath**, and D.M. Hillis. 2002. Phylogenetic relationships of the dwarf boas and a comparison of Bayesian and bootstrap measures of phylogenetic support. *Molecular Phylogenetics and Evolution* 25(2):361-371.

SOFTWARE

- RevBayes (C/C++) Bayesian phylogenetic inference using Markov chain Monte Carlo-relying on probabilistic graphical models and an interpreted programming language interface (development team member; https://github.com/revbayes/revbayes)
- DPPDiv (C++) Bayesian estimation of species divergence times under a Dirichlet process prior (developer; https://github.com/trayc7/FDPPDIV)
- FossilGen (C++) A program for simulating phylogenies and fossil occurrence times under the serial-sampled birth-death process (developer; https://github.com/trayc7/FossilGen)
- datelife (R/Python) A web service that synthesizes divergence time estimates from published studies—a Phylotastic service (co-developer; http://www.datelife.org, http://phylotastic.org)
- TreeSetViz (Java) Multidimensional scaling for visualizing and summarizing large sets of phylogenetic trees (contributor; http://comet.lehman.cuny.edu/treeviz)

Invited Talks

| The European Bioinformatics Institute, EMBL-EBI External Seminar Series | Oct. 2014 |
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| University of California, Berkeley, UCMP Fossil Coffee Seminar Series | 2014 |
| University of Connecticut, Ecology & Evolutionary Biology Seminar Series | 2014 |
| Smithsonian Institution National Museum of Natural History, PhyloPizza | 2014 |
| Brown University, Center for Computational Molecular Biology Seminar Series | 2013 |
| Phyloseminar.org Series on Integrating Fossils into Phylogenies (video link) | 2013 |
| Brigham Young University, Biology Department Seminar Series | 2013 |

| Cornell University, Ecology & Evolutionary Biology Weekly Seminar Series | 2013 |
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| France-Berkeley Fund Symposium on Fossils & Phylogenetics, Berkeley, CA | 2013 |
| Bowdoin College, Biology Department Seminar Series | 2013 |
| California Academy of Sciences, Research Seminar Series | 2012 |
| University of California, Davis Center for Population Biology Seminar Series | 2012 |
| Symposium on Molecular Dating, Hennig XXXI, Riverside, CA | 2012 |
| Symposium on Bayesian Inference of Phylogeny, Berkeley, CA | 2011 |
| SSB Symposium, Evolution 2010, Portland, OR | 2010 |
| CIPRES Symposium & Workshop on Evolutionary Simulation, Philadelphia, PA, USA | 2006 |

Conference Presentations

| Society for Vertebrate Paleontology (education/outreach poster), Berlin, Germany | Nov. 2014 |
|--|-----------|
| SACNAS 2014 (professional development session), Los Angeles, CA, USA | Oct. 2014 |
| iEvoBio 2014, Raleigh, NC, USA* | 2014 |
| Evolution 2013, Snowbird, UT, USA* | 2013 |
| Evolution 2012, Ottawa, Ontario, Canada* | 2012 |
| Evolution 2011, Norman, OK, USA | 2011 |
| Evolution 2009, Moscow, ID, USA | 2009 |
| Evolution 2007, Christchurch, New Zealand | 2007 |
| The Annual New Zealand Phylogenetics Meeting – Doom07, Whakapapa, NZ | 2007 |
| Evolution 2006, Stony Brook, NY, USA | 2006 |
| NSF IGERT Project Meeting (poster), Arlington, VA, USA | 2006 |
| Evolution 2004, Fort Collins, CO, USA | 2004 |
| *slides available at http://www.slideshare.net/travc7 | |

Grants and Awards

NESCent Academy Course: Phylogenetic Analysis Using RevBayes — PI: T. Heath; Co-PIs:

B. Boussau, S. Höhna, J. Huelsenbeck, M. Landis, B. Moore, F. Ronquist, T. Stadler (Summer 2014)

NSF DEB-1256993: "Integrating Fossil Data into Likelihood-based Phylogenetic Analyses with Trilobites as a Model System" (http://phylo.bio.ku.edu/fossil) — PI: M. Holder, Co-PI:

B. Lieberman, Postdoc/coauthor: T. Heath (2013 – 2016)

NSF Postdoctoral Fellowship in Biological Informatics (2008 – 2010)

University of Kansas/Haskell Indian Nations University NIH Institutional Research and Academic Career Development Award (IRACDA) Postdoctoral Fellowship (2008; declined)

University of Texas Ecology, Evolution, and Behavior Program Hartman Merit Fellowship (2008)

NSF, Integrative Graduate Education and Research Traineeship (IGERT) for graduate training in computational phylogenetics and applications to biology (2003 - 2008)

NSF, Integrative Graduate Education and Research Traineeship (IGERT) Travel Award (2007)

Allan Wilson Centre for Molecular Ecology & Evolution Summer Studentship, Biomathematics Research Centre, University of Canterbury, Christchurch, NZ (2007)

University of Texas Ecology, Evolution, and Behavior Program Hamilton Fellowship (2006)

Summer Institute in Statistical Genetics Scholarship, North Carolina State University (2004)

Workshop and Hackathon Participation

AIM-UP! Workshop on "Museums and Climate Change", Pacific Grove, CA (February 2014)

NESCent BEAST Developers Working Group; Durham, NC (December 2013)

France-Berkeley Fund Workshop on Integrating Molecular Phylogenies and the Fossil Record; Organizers: H. Morlon, T. Quental, C. Marshall; Berkeley, CA (2013)

AdaCamp Unconference for Women in Open Technology, San Francisco, CA (2013)

AIM-UP! All Hands Meeting (www.aim-up.org), Cambridge, MA (2013)

NESCent Hackathon: Phylotastic! Making the Tree of Life Accessible for Automated Re-use, Durham, NC (2012)

Workshop on Molecular Evolution, Marine Biological Laboratory, Woods Hole, MA (2007)

Cass Workshop, "Random models in phylogenetics and resolving ancient divergences", Biomathematics Research Centre, University of Canterbury, Arthur's Pass, New Zealand (2007)

Summer Institute in Statistical Genetics, North Carolina State University, Raleigh, NC (2004)

Professional Service & Outreach

Evolution 2015 SSB Symposium Organizer – "Breaking Barriers: Empirical, Theoretical, and Gender Issues in Phylogenetics"; with C. Moreau, B. O'Meara, & S. Perkins (Brazil, June 2015)

SACNAS Professional Development Session Organizer – Cracking the (bio)Code: How to Start a Research Career in Computational Biology – 2014 SACNAS National Conference, Los Angeles, CA; with F. Zapata, E. Huerta-Sanchez, & C. Sandefur (October 18, 2014)

Blog contributor: http://treethinkers.org/blog (March 2013 – present)

Society of Systematic Biologists Elected Council Member (2013 – 2015)

iEvoBio 2013 Organizing Committee Member (2012 - 2013)

Mentor for the Undergraduate Diversity at SSE/SSB Program (Evolution 2011 and Evolution 2012) Reviewer:

- Systematic Biology Editorial Board Member (2011 present)
- Bioinformatics, BMC Evolutionary Biology, Evolution, Gene, Journal of Systematics & Evolution, Molecular Biology & Evolution, Proceedings B, Systematic Biology, Trends in Ecology & Evolution
- The Graduate Women in Science 2012 National Fellowships Program
- The Society of Systematic Biologists Graduate Student Research Award Competition (2012, 2014)

Invited Panelist/Speaker:

- Girls Who Code Summer Program, @Twitter, San Francisco, CA (www.girlswhocode.com; 2013)
- The Women in Science and Engineering Residential Program Seminar, UC-Berkeley (2011)
- The Initiative for Maximizing Student Development (IMSD) Graduate Fellows Program Workshop, UC-Berkeley (2011)

Society/Organization Membership: Society of Systematic Biologists, Society for Advancement of Chicanos and Native Americans in Science (SACNAS), Society of Vertebrate Paleontology