NICHOLAS G. CRAWFORD

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AREAS OF SPECIALIZATION

 Bioinformatics, evolutionary biology, computational biology, population genomics, phylogenomics, speciation, ultra conserved elements, herpetology and entomology

EDUCATION

• 2013, Postdoctoral Fellow, California Academy of Sciences

Advisors: Durrell Kapan and Brian Simison

• 2007/13, Ph.D. Biology, Boston University

Advisor: Christopher Schneider

Thesis Title: Anolis Lizard Genomics and Phylogenetics

Additional Projects:

- o Mammal, Reptile, and Avian Phylogenomics, with Travis Glenn
- o Butterfly Speciation Genomics, with Sean Mullen

2004/07, M.S. Biology, San Diego State University

Advisor: Tod Reeder

Thesis: Population structure within the Plateau Striped Whiptail (Aspidoscelis velox complex) a parthenogenetic lizard

Additional Project:

Phylogenetic relationships among Australian skinks of the genus Glaphyromorphus

1997/01, B.S. Biology, Union College

Graduated Cum Laude with Academic and Departmental Honors

SKILLS AND TECHNIQUES

- Programming Languages/Frameworks: Python (Modules: Numpy, Scipy, Matplotlib, Biopython, Pysam, Multiprocessing, iPython, MrJob, Pandas, Flask, and Django), MySQL/Sqlite3, Git/Github, Unix/Linux, LSF/SGE/AWS Clusters, Hadoop/MapReduce, also some R, Perl, Html, and CSS
- Software: GATK, ABySS, Trinity, Breakdancer, Samtools, VCFtools, Tophat, Cufflinks, Bowtie2, Stampy, GO Elite, Blast/NCBI, MrBayes, Sequencher, Oligo (primer design), Genemapper, GenAlEx, Arlequin
- Laboratory: Genomic, RAD-tag, and RNA-seq Illumina library construction, Primer Design, DNA purification, PCR, qPCR (Taqman and Kappa assays), Sanger Sequencing (both ABI3130 and ABI377 genetic analyzers), Microsatellite Genotyping, Bioanalyzer, and constructing cDNA libraries
- Miscellaneous Skills: Protein Purification (FPLC), Mammalian Tissue Culture, Ordering and Stocking, Maintaining Laboratory Compliance with Safety Regulations, Training Students and Techs, and Catching lizards

GRANTS AND SCHOLARSHIPS

- 2012 Grand Challenges Award, Smithsonian Institute Consortia, Next Generation Phylogenetics, MJ Braun, K Wurdack, W Wcislo, J Maldonado, K Helgen, S Brady, M Cummings, TC Glenn, BC Faircloth, RT Brumeld, E Braun, JC McCormack, NG Crawford, N White. \$100,000.
- 2011 Next-generation Sequencing Small Grant. Phylogenetic utility of ultra- conserved elements for the avian tree of life. MJ Braun, ND White, TC Glenn, BC Faircloth, RT Brumeld, EL Braun, JE McCormack, NG Crawford. \$10,000
- 2011 Amazon Education Research Grant (aws.amazon.com), Computational Resources. BC Faircloth, NG Crawford, JE McCormack. \$10,000.
- 2010 Amazon Education Research Grant (aws.amazon.com), Computational Resources. NG Crawford, BC Faircloth, TC Glenn. \$7,500.
- 2010 Doctoral Dissertation Improvement Grant (NSF). NG Crawford, C Schneider. \$15,000.
- 2009 Theodore Roosevelt Memorial Fund (American Museum of Natural History) labwork/fieldwork. \$1,500.

- 2005 Theodore Roosevelt Memorial Fund (American Museum of Natural History) labwork/fieldwork. \$1,960.
- 2005 Harry E. Hamber Memorial Scholarship, tuition. \$1,400.
- 2000 IEF Grant, thesis research. \$100.
- 2000 NYSEP Grant, summer stipend. \$1,700.
- 1999 Booth Ferris Grant, summer stipend. \$1,700.

PUBLICATIONS

- Marcus R. Kronforst, Matthew E. B. Hansen, Nicholas G. Crawford, Jason R. Gallant, Rob J. Kulathinal, Durrell D. Kapan, Sean P. Mullen. 2013. Hybridization reveals the evolving genomic architecture of speciation. Cell Reports, 5, 1 –12.
- John E. McCormack, Michael G. Harvey, Brant C. Faircloth, **Nicholas G. Crawford**, Travis C. Glenn, Robb T. Brumfield. 2013. Aphylogeny of birds based on over 1,500 loci collected by target enrichment and high-throughput sequencing. PLoS ONE. 8(1): e54848.
- Martha M. Muñoz, Nicholas G. Crawford, Thomas J. McGreevy, Rebecca D. Tarvin, Nicholas J. Messana, Liam J. Revell, Rosanne M. Zandvliet, Juanita M. Hopwood, Elbert Mock, André L. Schneider, and Christopher J. Schneider. 2013.
 Divergence in coloration and the evolution of reproductive isolation in the Anolis marmoratus species complex.
 Molecular Ecology. 22(10), 2668–2682.
- John E. McCormack, Michael G. Harvey, Brant C. Faircloth, **Nicholas G. Crawford**, Travis C. Glenn, Robb T. Brumfield. 2013. *A phylogeny of birds based on over 1,500 loci collected by target enrichment and high-throughput sequencing*. PLoS ONE. 8(1), e54848.
- John A. St. John, Edward L. Braun, Sally R. Isberg, Lee G. Miles, Amanda Y. Chong, Jamie Gongora, Pauline Dalzell, Christopher Moran, Taisen Iguchi, Bertrand Bed'Hom, Shane C. Burgess, Amanda M. Cooksey, Todd A. Castoe, Arkhat Abzhanov, Llewellyn D. Densmore, Miryam Venegas-Anya, Matthew J. Greenwold, Roger H. Sawyer, Federico G. Hoffmann, Nicholas G. Crawford, Jennifer C. Drew, Scott V. Edwards, Matthew K. Fujita, Jonathan M. Howard, Brant C. Faircloth, Daniel E. Janes, Shahid Yar Khan, Satomi Kohno, A.P. Jason de Koning, Stacey L. Lance, Fiona M. McCarthy, John E. McCormack, Mark E. Merchant, Daniel G. Peterson, David D. Pollock, Nader Pourmand, Brian J. Raney, Kyria A. Roessler, Jeremy R. Sanford, Carl J. Schmidt, Eric W. Triplett, Tracey D. Tuberville, Erich D. Jarvis, Louis J. Guillette Jr, Travis C. Glenn, Richard E. Green and David A. Ray. 2012. Sequencing three crocodilian genomes to illuminate the evolution of archosaurs and amniotes. Genome Biology. 13, 415.
- Kenro Kusumi, Rob J. Kulathinal, Arhat Abzhanov, Stephane Boissinot, Nicholas G. Crawford, Brant C. Faircloth, Travis C. Glenn, Daniel E. Janes, Jonathan B. Losos, Douglas B. Menke, Steven Poe, Thomas J. Sanger, Christopher J. Schneider, Jessica Stapley, Juli Wade, Jeanne Wilson-Rawls. 2012. Developing a community-based genetic nomenclature for anole lizards. BMC genomics, 12(1), 554.
- **Nicholas G. Crawford**, Brant C. Faircloth, John E. McCormack, Robb T. Brumfield, Kevin Winker, Travis C. Glenn. 2012. *More than 1000 ultraconserved elements provide evidence that turtles are the sister group of archosaurs*. Biology Letters. 8(5), 783-786.
- Brant C. Faircloth, John E. McCormack, **Nicholas G. Crawford**, Michael Harvey, Robb T. Brumfield, Travis C. Glenn. 2012. *Ultraconserved elements anchor thousands of genetic markers for target enrichment spanning multiple evolutionary timescales*. Systematic Biology. 61(5), 713-715.
- John E. McCormack, Brant C. Faircloth, **Nicholas G. Crawford**, Patricia Adair Gowaty, Robb T. Brumfield, Travis C. Glenn. 2011. *Ultraconserved Elements Are Novel Phylogenomic Markers that Resolve Placental Mammal Phylogeny when Combined with Species Tree Analysis*. Genome Research. 22(4), 746-54.
- **Nicholas G. Crawford**. 2010. *SMOGD: Software for the Measurement of Genetic Diversity*. Molecular Ecology Resources, 10: 556-557.
- Nicholas G. Crawford, Jaime Zaldvar-Rae, Cris Hagen, Amanda Schable, Erica Bree Rosenblum, Jeff A. Graves, Tod W. Reeder, Michael G. Ritchie, Travis C. Glenn. 2007. Thirteen polymorphic microsatellite DNA loci from whiptails of the genus Aspidoscelis (Teiidae: Squamata) and related cnemidophorine lizards. Molecular Ecology Resources. 8: 219-223
- Nicholas G. Crawford, Cris Hagen, Heather F. Sahli, Elizabeth A. Stacy, Travis C. Glenn. 2007. Fifteen polymorphic microsatellite loci from Hawaiis Metrosideros polymorpha Myrtaceae: Myrtales), a model species for ecology and evolution. Molecular Ecology Resources, 8, 308-310.
- Caleb R. Hickman, Maureen B. Peters, **Nicholas G. Crawford**, Cris Hagen, Travis C. Glenn, Christopher M. Sommers. 2008. *Development and characterization of microsatellite loci in the American white pelican (Pelecanus*

erythrorhynchos). Molecular Ecology Resources, 8, 1439-1441.

- Nicholas G. Crawford, Maureen B. Peters, Cris Hagen, Travis C. Glenn, Stephen K. Davis, Christopher M. Somers.
 2007. Twelve polymorphic microsatellite loci from Spragues pipit, Anthus spragueii (Motacillidae:Passeriformes), a threatened grassland endemic songbird. Molecular Ecology Resources, 9, 315-317.
- Nicholas G. Crawford. 2007. Microsatellites in cnemidophorine lizards: their utility in investigating the landscape genetics of the plateau striped whiptail (Aspidoscelis velox Complex). Masters Thesis: San Diego State University.
- Olga V. Tsyusko, Tracey D. Tuberville, Maureen B. Peters, Nicholas G. Crawford, Cris Hagen, Steve Weller, Ann Sakai, and Travis C. Glenn. 2007. Microsatellite markers isolated from polyploid woodsorrell (Oxalis alpina). Molecular Ecology Notes, 7, 1284-1286.

PROFESSIONAL EXPERIENCE

- 2007-2011 Teaching Fellow, Boston University: Introductory Biology, Genetics, Evolution, Animal Behavior
- 2007 South Carolina, Summer, 6 months, Research Technician, supervised by Travis Glenn
- 2006 South Carolina, Summer, 6 weeks, Microsatellite loci preparation at the Savannah River Ecology Laboratory, supervised by Travis Glenn
- 2004/2005 Graduate Teaching Assistant, SDSU, Introductory Biology and Introductory Zoology
- 2004 Research Assistant, SDSU, Advisor: Tod Reeder
- 2002/03 Research Associate, Boston Biochem
- 2001 Research Associate, Pfizer Pharmaceuticals
- 2001 Research Associate, Harvard Medical School

FIELD EXPERIENCE

- 2012 Guantanamo Bay, Cuba, Summer, 1 week, collecting lizards
- 2009 Guadeloupe, French West Indies, Summer, 5 days, collecting lizards
- 2007 New Mexico, Summer, 1 week, collecting lizards
- 2005 New Mexico, Summer, 1 week, collecting lizards
- 2005 Arizona, Summer, 2 weeks, collecting lizards
- 2004 Arizona, Summer, 2 weeks, collecting lizards
- 2001 New York, Summer, 1 week, collecting treefrogs
- 1999 New York, Summer, 1 week, collecting treefrogs

CONFERENCES, MEETINGS, AND TALKS

- 2013 Joint annual meeting of the Society for the Study of Evolution and the Society of Systematic Biologists Presentation Title: Genomics of local adaptation and colorful pigmentation in Anolis lizards
- 2013 UMass Lowell Seminar Series: Invited Speaker
 Presentation Title: From Archosaurs to Anoles: Genomic Approaches to Studying Reptile Evolution
- 2012 Union College Seminar Series: Invited Speaker Union College.
 Presentation Title: Genomic approaches to understanding reptile evolution.
- 2012 World Congress of Herpetology, Vancouver.
 Presentation Title: The Genetics of Colorful Pigmentation in Anolis Lizards.
- 2012 Evolutionary Genomics Super Group, Broad Institute *Presentation Title:* Ultraconserved Elements as Phylogenomic Markers.
- 2012 MCZ Lunchtime Seminar, Harvard University
 Presentation Title: Thousands of ultraconserved elements in combination with cloud computing and species-tree methods help resolve deep divergences in reptiles, mammals, and birds
- 2010 The Genetics and Evolution of Animal Coloration, Radcliffe Workshop Presentation Title: Identifying coloration genes when you can't easily do QTL mapping
- 2009 Anole Symposium Harvard University
 Presentation Title: Anolis carolinensis: Pigmentation Genetics Poster Title: Genome scan identifies two loci associated with color polymorphism in Anolis marmoratus
- 2009 Gordon Conference: Evolutionary & Ecological Functional Genomics Tilton, New Hampshire.
 Poster Title: Genetics of Colorful Pigmentation in Anolis Lizards
- 2009 Society for Integrative and Comparative Biology Annual Meeting, Boston Massachusetts.
 Presentation Title: Evolution of Dewlap Pigmentation in Anoline lizards
- 2008 Union College Seminar Series: Invited Speaker Union College

Presentation Title: Population Structure of the Plateau Striped Whiptail a Parthenogenetic Species of Lizard

- 2007 Island Biogeography Symposium, Harvard University, Cambridge Massachusetts
- 2005 American Society of Ichthyologists and Herpetologists 85th Annual Meeting, Tampa, Florida *Presentation Title:* Phylogenetic relationships among Australian skinks of the genus Glaphyromorphus
- 2004 American Society of Ichthyologists and Herpetologists 84th Annual Meeting, Norman, Oklahoma
- 2004 Evolution Conference, Fort Collins, Colorado

SOCIETY AFFILIATIONS

- American Society of Ichthyologists and Herpetologists (not presently affiliated)
- Society for Comparative and Integrative Biology (not presently affiliated)