

# Ho, Wei-Chin

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## Education

- 2011 - 2017     **Ph.D., Department of Ecology and Evolutionary Biology, University of Michigan, Ann Arbor, USA.**
- Dissertation title: The genotype-phenotype map: origins, properties, and evolutionary consequences (Advisor: Dr. Jianzhi Zhang)
- 2008 - 2010     **M.S., Institute of Zoology, National Taiwan University, Taiwan.**
- Thesis title: Inferring speciation history of *Drosophila* by massive parallel sequencing (Advisor: Dr. Chau-Ti Ting)
- 2004 - 2008     **B.S., Department of Life Science (minor in chemistry), National Taiwan University, Taiwan.**

## Professional Experiences

- 2017 -            **Postdoctoral Research Associate, Center for Mechanisms of Evolution, The Biodesign Institute, Arizona State University, USA.** (Advisor: Dr. Michael Lynch)
- 2010 - 2011     **Research Assistant, Department of Life Science, National Taiwan University, Taiwan.** (Advisor: Dr. Chau-Ti Ting)

## Research Interests

- Factors affecting evolutionary outcomes and their evolution, including (but not limited to) mutation rates, mutational effects, robustness, and plasticity.
- Relative contribution of chance and necessity in evolution.
- Predictability of evolutionary outcomes *via* systems biology approaches.

## Publications

6. [Wei-Chin Ho](#) and Jianzhi Zhang (2019) **Genetic gene expression changes during environmental adaptations tend to reverse plastic changes even after the correction for statistical nonindependence.** *Mol. Biol. Evol.* in press
5. [Wei-Chin Ho](#) and Jianzhi Zhang (2018) **Evolutionary adaptations to new environments generally reverse plastic phenotypic changes.** *Nat. Comm.* 9: 350.
4. [Wei-Chin Ho](#), Yoshikazu Ohya, and Jianzhi Zhang (2017) **Testing the neutral hypothesis of phenotypic evolution.** *Proc. Natl. Acad. Sci. U.S.A.* 114(46): 12219-12224.
3. Calum J. Maclean\*, Brian P.H. Metzger\*, Jian-Rong Yang\*, [Wei-Chin Ho](#), Bryan Moyers, and Jianzhi Zhang (2017) **Deciphering the Genic Basis of Yeast Fitness Variation by Simultaneous Forward and Reverse Genetics.** *Mol. Biol. Evol.* 34(10): 2486-2502. (\*co-first authors)

2. Wei-Chin Ho and Jianzhi Zhang (2016) **Adaptive genetic robustness of *Escherichia coli* metabolic fluxes.** *Mol. Biol. Evol.* 33(5): 1164-1176.

1. Wei-Chin Ho and Jianzhi Zhang (2014) **The genotype-phenotype map of yeast complex traits: basic parameters and the role of natural selection.** *Mol. Biol. Evol.* 31(6): 1568-1580.

#### Public Talks and Conference Oral Presentation

- **“Phenotypic changes in organismal adaptation to new environments: plasticity distorts while evolution restores”**  
Annual Meeting of SMCB, Gold Coast, Australia, July 2016
- **“Adaptive origin of the genetic robustness of metabolic fluxes”**  
Annual Meeting of SMCB, Vienna, Austria, July 2015
- **“Prevalent adaptive evolution of morphological traits in the budding yeast *Saccharomyces cerevisiae*”**  
Annual Meeting of SMCB, San Juan, Puerto Rico, June 2014
- **“Natural selection for robustness shapes the genetic architecture of yeast complex traits”**  
University of Michigan, Ann Arbor, Jan 2013
- **“Expression divergence between two behavioral races of *Drosophila melanogaster* revealed by whole transcriptome analyses”**  
Annual Meeting of SMCB, Lyon, France, July 2010

#### Conference Poster Presentation

- **“Experimental evolution of *Escherichia coli* mutators in a complex environment”**  
Annual Meeting of SMCB, Yokohama, Japan, July 2018
- **“Does genetic correlation constrain or facilitate long-term phenotypic evolution?”**  
Annual Meeting of SMCB, Austin, USA, July 2017
- **“Testing the neutral hypothesis of phenotypic evolution using 220 morphological traits in yeast”**  
Annual Meeting of SMCB, Chicago, USA, July 2013
- **“Genome-wide genetic architecture of morphological traits in yeast”**  
Annual Meeting of SMCB, Dublin, Ireland, June 2012
- **“Differential gene expression between two behavioral races of *Drosophila melanogaster*”**  
Asian-Pacific *Drosophila* Research Conference, Taipei, Taiwan, May 2011
- **“Searching candidate loci responsible for behavior differentiation between two *Drosophila melanogaster* races by genomic approaches”**  
Symposium on College of Life Science, National Taiwan University, Taipei, Taiwan, June 2010
- **“Incomplete lineage sorting in *Drosophila simulans* clade”**  
Symposium on College of Life Science, National Taiwan University, Taipei, Taiwan, June 2009

#### Teaching and Mentoring Experiences

- Graduate Student Mentor, Supervised Teaching (EEB/MCDB 801), University of Michigan, Winter 2017
- Graduate Student Instructor, Genetics (BIOLOGY 305), University of Michigan, Winter 2017, Winter 2015, Winter 2013, Winter 2012
- Graduate Student Instructor, Evolution, University of Michigan (EEB 390), Fall 2013

- Graduate Student Instructor, Introductory Biology: Ecology and Evolution (BIOLOGY 171), University of Michigan, Fall 2011
- Teaching Assistant, Population Genetics (EEB 5045), National Taiwan University, Fall 2010-2008
- Teaching Assistant, Genetics (LS 3007), National Taiwan University, Spring 2010
- Teaching Assistant, General Biology (LS 1006), National Taiwan University, Fall 2009
- Teaching Assistant, General Biology Laboratory (LS 1017), National Taiwan University, Fall 2008

#### **Awards and Fellowships**

- Rackham One-Term Dissertation Fellowship, Rackham Graduate School, University of Michigan, 2016
- Young Investigator Travel Award, Annual Meeting of the Society for Molecular Biology and Evolution, 2018-2015
- Graduate Travel Award, Annual Meeting of the Society for Molecular Biology and Evolution, 2014, 2010
- Chia-Lun Lo Fellowship (\$10,000), Rackham Graduate School, University of Michigan, 2013
- Dean's Award, College of Life Science, National Taiwan University, 2010
- Outstanding Students Conference Travel Grant, Foundation for the Advancement of Outstanding Scholarship, 2010
- Reward of Excellence, Symposium on College of Life Science, National Taiwan University, 2010

#### **Computational Skills**

- Programming languages: C/C++, Perl, Python.
- Statistical computing: R, MATLAB.
- Evolutionary analysis tools: PAML, MrBayes, Phylip, MEGA.

#### **Academic Services**

- *Ad-hoc* Journal Reviewer for *BMC Genomics*, *Genome Biol. Evol.*, *Mol. Biol. Evol.*, *PLoS Genetics*.
- Voluntary helpers in Software Carpentry Workshops at University of Michigan (Oct-17 2016, Dec-14 2016).
- Committee Representative, 13th Annual University of Michigan Early Career Scientists Symposium: Ecology and Evolutionary Biology of Phenotypic Plasticity, University of Michigan, Ann Arbor, Fall 2016 - Winter 2017
- Seminar Committee Representative, Graduate Researchers in Ecology and Evolutionary Biology, University of Michigan, Ann Arbor, Fall 2016 - Winter 2017, Fall 2013 - Winter 2014