LI WANG

(806) 319-0182 \$\dilpisorus@gmail.com

Office: 353B Bessey Hall, Iowa State University, Ames, IA 50011 United States

Website: https://lepisorus.github.io/

EDUCATION

Institute of Botany, Chinese Academy of Sciences, Beijing, China

2011

PhD Plant Evolution

Thesis: Phylogeny, diversification and phylogeography of the derived fern genus Lepisorus

Wuhan University, Wuhan, Hubei Province, China

2006

BS Biotechnology

PROFESSIONAL EXPERIENCE

Iowa State University

April 2014 - present

Postdoctoral Research Associate on Bioinformatics

- Principal Investigator: Dr. Matt Hufford
- Investigated demography and its effect on mutation load in maize and teosinte using high-depth whole genome re-sequencing data
- Explored parallel adaptation to highland environments in maize populations
- Conducting comparative genomic analyses of four de-novo assembled genomes of maize landraces
- Mentored graduate students in molecular and bioinformatic techniques

Iowa State University

April 2014 - December 2015

Postdoctoral Research Associate on Bioinformatics

- Principal Investigator: Dr. Andrew Severin
- Provided services at Genomic Informatics Center
- Detected selection during soybean improvement

Texas Tech University

Feburary 2012 - March 2014

Postdoctoral Research Associate on Bioinformatics

- Studied gene expression and co-expression network differences resulting from latitude adaptation in balsam poplar with transcriptome data
- Evaluated gene expression pattern and enriched pathways owing to sex dimorphism in dioecious poplars
- Assessed DNA and protein evolutionary rates and detected signatures of selection
- Mentored graduate students in molecular and bioinformatic techniques

Institute of Botany, Chinese Academy of Sciences

June 2007 - July 2011

Research Assistant

- Investigated phylogeny, phylogeography, polyploidization and reticulate evolution in ferns
- Mentored multiple graduate students in experimental and analyses techniques

Goettingen University, Germany

February 2009 - November 2010

Research Assistant

• Examined phylogeny, phylogeography, polyploidization and reticulate evolution in ferns

Natural History Museum, London, UK

December 2009, April 2010

Research Assistant

• Inspected phylogeny, phylogeography, polyploidization and reticulate evolution in ferns

Institute of Hydrobiology, Chinese Academy of Sciences Undergraduate Research

June 2005 - December 2005

• Collected data of fish diversity to determine the effect of constructing the three gorges dam on Yangtze River

SKILLS

Scripting languages

• Proficient in Unix scripting/bash, python, perl, R, markdown and latex

Software proficiencies

- Assembly: Trinity, Falcon, MaSuRCA, Platanus, GenomeScope, Redundans
- Annotation: GeneMark, Augustus, Busco, CEGMA, Repeatmasker, BLAST, Trinotate, Blobtools
- Aligners: Bowtie, BWA, Gmap, Gsnap, Hisat2
- Transcriptomics: Trinity, Rsem, edgeR, WGCNA, mapMan
- Comparative genomics: GATK, Bedtools, vcftools, PLINK, TASSEL, MEGA
- Machine learning: scikit learn (regression, clustering algorithms)
- Population genomics: PopGenome, ANGSD, NGSadmix, MSMC, PSMC, fastSimcoal, ms
- Phylogenetics: Mafft, ClustalW, Muscle, RaxML, Phagon
- Visualization: Circos, Cytoscape, Photoshop
- Other: Seqtk, Aspera, Bamtools, Bioawk, Cdbfasta, Fastqc, Gawk, Jellyfish, Picard Tools, Samtools, SRA-Toolkit, Tabix, Trimmomatic, Sickle, Vcftools, BLAT, Emboss etc

Lab proficiencies

• DNA and RNA extraction, bioNano, PCR, clone etc.

CODING ACCOMPLISHMENTS

Publicly available pipelines and scripts

- https://github.com/HuffordLab/Wang_et_al_Demography computational pipeline for demography and its effect on mutation in maize
- https://github.com/ISUgenomics/commonscripts Common genomic analyses

AWARDS AND HONORS

2011

• DEAN Award in Chinese Academy of Sciences, Beijing, China

2011

• The excellent graduate student in Chinese Academy of Sciences, Beijing, China

2010-2011

• BHPB-GUCAS Scholarship for excellent Ph.D. candidates, Beijing, China; \$2,000

2010

• The second prize for giving talks at students' symposium in Natural History Museum, London, UK

2008 - 2010

• DAAD-CAS Joint Scholarship Program – 1,000 Euro per month for two years, Goettingen, Germany

2009 - 2010

• Merit Student by the Graduate University of Chinese Academy of Sciences, Beijing, China

CONFERENCE PARTICIPATION

Oral Presentation

- Li Wang, Harald Schneider, Xianchun Zhang. (July 2011). The rise of the Himalaya enforced the diversification of SE Asian ferns by altering the monsoon regimes. International Botany Conference, Melbourne, Australia
- Li Wang, Harald Schneider, Qiaoping Xiang, Xianchun Zhang. (November 2010). Phylogeography of the alpine fern *Lepisorus clathratus* on "the roof of the world". The Fifth Asian Fern Symposium, Shenzhen, China
- Li Wang, Xianchun Zhang. (December 2008). Phylogeny of the paleotropical fern genus *Lepisorus* (Polypodiaceae, Polypodiopsida) inferred from four chloroplast genome regions. Symposium on Academic study of Chinese ferns, Shenzhen, China

Poster Presentation

- Li Wang, Emily Josephs, Lucas M. Roberts. Timothy Mathes Beissinger, Jeffrey Ross-Ibarra, Matthew Hufford. (March 2017). Detection of Convergent Highland Adaptation in maize landrace populations. 59th maize genetic conference, St. Luis, Missouri, USA
- Li Wang, Timothy Mathes Beissinger, Jeffrey Ross-Ibarra, Matthew Hufford. (March 2016). Evolution of maize during post-domestication expansion across the Americas. 58th maize genetic conference, Jacksonville, Florida, USA
- Li Wang, Timothy Mathes Beissinger, Jeffrey Ross-Ibarra, Matthew Hufford. (March 2015). Inference of Maize Population History during Migration to Highland Habitats. 57th maize genetic conference, St. Charles, Illinois, USA
- Li Wang, Timothy Mathes Beissinger, Jeffrey Ross-Ibarra, Matthew Hufford. (January 2015). Inference of Maize Population History during Migration to Highland Habitats. Plant and Animal Genome Conference, San Diego, CA, USA
- Li Wang, Matt Olson. (June 2013). Timing for success: exprssion phenotype and local adaptation related to latitude in the boreal forest tree. Evolution, Snowbird, Utah, USA

Attended

- July 2009, Niche Evolution Conference, Zurich, Switzerland
- August 2009, Systematics 2009, Leiden, Netherland
- May 2006, Symposium on Academic study of Chinese ferns, Shangri-La, China

LEADERSHIP SKILLS

Social activity manager of Postdoctor Association at Iowa State University

 $Participated \ in \ the \ Self-expression \ and \ Leadership \ Program \ at \ Landmark \ (\texttt{http://www.landmarkworldwide.com/})$

Will take a leadership program for women scientists on a three-week expedition to Antarctica in February 2018

PUBLICATIONS

Published

- Li Wang, Timothy Mathes Beissinger, Anne Lorant, Claudia Ross-Ibarra, Jeffrey Ross-Ibarra, Matthew Hufford. 2017. The interplay of demography and selection during maize domestication and expansion. bioRxiv. doi: https://doi.org/10.1101/114579.
- Timothy Beissinger, **Li Wang**, Kate Crosby, Arun Durvasula, Matthew B Hufford, Jeffrey Ross-Ibarra. 2016. Recent demography drives changes in linked selection across the maize genome. **Nature plants** 2: 16084.
- Li Wang, Peter Tiffin, Matthew Olson. 2014. Timing for success: exprssion phenotype and local adaptation related to latitude in the boreal forest tree, Populus balsamifera. Tree Genetics and Genomes. DOI: 10.1007/s11295-014-0731-3.
- Harald Schneider, Lijuan He, Jeannine Marquardt, Li Wang, Jochen Heinrichs, Sabine Hennequin, Xianchun Zhang. 2013. Exploring the origin of the latitudinal diversity gradient: contrasting the sister fern genera *Phegopteris* and *Pseudophegopteris*. Journal of Systematics and Evolution 51: 61-70.
- Xianchun Zhang, Ran Wei, Hongmei Liu, Lijuan He, **Li Wang**, Gangming Zhang. 2013. Phylogeny and classification of the extant lycophytes and ferns from China. **Chinese Bulletin of Botany** 48: 119-137.
- Li Wang, Harald Schneider, Xianchun Zhang, Qiaoping Xiang. 2012. The rise of the Himalaya enforced the diversification of SE Asian ferns by altering the monsoon regimes. BMC Plant Biology 12:210. DOI: 10.1186/1471-2229-12-210.
- Li Wang, Harald Schneider, Zhiqiang Wu, Lijuan He, Xianchun Zhang, Qiaoping Xiang. 2012. Indehiscent sporangia enable the accumulation of local fern diversity at the Qinghai-Tibetan Plateau. BMC Evolutionary Biology 12:158. DOI:10.1186/1471-2148-12-158.
- Li Wang, Zhiqiang Wu, Nadia Bystriakova, Stephen W. Ansell, Qiaoping Xiang, Jochen Heinrichs, Harald Schneider, Xianchun Zhang. 2011. Phylogeography of the alpine fern *Lepisorus clathratus* on "the roof of the world". PloS One 6: e25896. DOI: 10.1371/journal.pone.0025896.
- Li Wang, Xinping Qi, Qiaoping Xiang, Jochen Heinrichs, Harald Schneider, Xianchun Zhang. 2010. Phylogeny of the paleotropical fern genus *Lepisorus* (Polypodiaceae, Polypodiopsida) inferred from four chloroplast genome regions. Molecular Phylogenetics and Evolution 54: 211-225.
- Li Wang, Zhiqiang Wu, Qiaoping Xiang, Jochen Heinrichs, Harald Schneider, Xianchun Zhang. 2010. A molecular phylogeny and a revised classification of tribe Lepisoreae (Polypodiaceae) based on an analysis of four plastid DNA regions. Botanical Journal of the Linnean Society 162: 28-38.
- Hongmei Liu, Li Wang, Hui Zeng, Xianchun Zhang. 2008. Advances in the studies of lycophytes and monilophytes with reference to systematic arrangement of families distributed in China. Journal of Systematics and Evolution 46: 808-829.
- Xianchun Zhang, Li Wang, Xinping Qi. 2008. Additions to the pteridophyte flora of Xizang (IV). Newsletter of Himalayan Botany 41: 20-27.
- Li Wang, Xianchun Zhang. 2007. Pteridophyta. in Zhenyu Li, Lei Shi (editors), Flora of Emei Mountain. Beijing Science and Technology Press, Beijing, pp. 153-224.
- Li Wang, Xianchun Zhang. 2007. Chinese aquatic ferns. China Nature 138: 54-55.

In Preparation

- Purushottam R. Lomate, Veena Dewangan, Neha S. Mahajan, Yashwant Kumar, Abhijeet Kulkarni,
 Li Wang, Smita Saxena, Vidya S. Gupta, and Ashok P. Giri. 2017. Integrated transcriptomic and proteomic analyses revealed the participation of endogenous protease inhibitors in the regulation of protease gene expression in Helicoverpa armigera (submitted)
- Garrett M. Janzen, **Li Wang**, Matthew B. Hufford. Review: Adaptive Introgression Expanded the Genetic Base of Crops during post-Domestication Spread. (**Expected 2017**)
- Brian Sanderson, **Li Wang**, Peter Tiffin, Zhi-qiang Wu, Matthew S. Olson. Males and females of *Populus balsamifera* differ primarily in the expression of energy-related genes. (**Expected 2017**)

RESEARCH FIELD WORK

Nov 2015

Mexican Central Plateau, Mexico

- Collected teosinte hybrid populations
- Team work with collaborators from both University of California Davis and Mexican national laboratory of genomics for biodiversity

Jun-Jul 2012 Alaska, US

- Collected leaves for RNA extraction in the common garden for balsam poplar at University of Alaska,
 Fairbanks
- Cooperated with researchers in University of Alaska, Fairbanks

Jul 2010 Altai Mountains, Russia

- Studied fern biodiversity in the region of Altai Mountains
- Worked jointly with Russian collaborators

Jun 2010 Alps, Switzerland and France

- Explored fern biodiversity in the Alps and collected populations of several fern species
- Worked closely with collaborators from Natural History Museum, UK

Nov 2008 Tropical and subtropical regions, China

• Examined fern biodiversity in Hainan and Guangdong Province

- \bullet Worked together with researchers from Zhongshan University, China

May 2008 Central Inner Mongolia, China

• Collected populations of *Lepisorus clathratus* in the northern desert region of China

Dec 2007 Southeast of Yunnan, China

- Investigated plant biodiversity at border of China and Vietnam
- Discovered several new species
- Collaborated with researches from Kunming Botanical Garden, China

Oct 2007 The Himalayas and Tibetan Plateau, China

- Investigated fern biodiversity at southeast of Tibet as a member of expedition team organized by World Wildlife Foundation
- Collaborated with many researches from different discipline

Aug 2007 Hengduan Mountains, China

• Explored fern biodiversity

May 2005 The three Gorges region, China

- Studied fish biodiversity
- Cooperated with local fishermen and administrative officers

REFERENCES

Dr. Matt Hufford

Assisstant Professor of Biological Sciences Ecology, Evolution, & Organismal Biology Iowa State University Ames, Iowa 50011

Tel: (515) 294-8511

Email: mhufford@iastate.edu

Dr. Andrew Severin

Scientist I/Facility Manager Genome Informatics Facility Iowa State University Ames, Iowa 50011

Tel: (515) 294-1320

Email: severin@iastate.edu

Dr. Jeffrey Ross-Ibarra

Professor of Biological Sciences Department of Plant Sciences University of California, Davis Davis, CA 95616

Tel: (530) 752-1152

Email: rossibarra@ucdavis.edu

Dr. Matt Olson

Associate Professor Department of Biological Sciences Texas Tech University Lubbock, TX 79409 Tel: (806) 834-7252

Email: matt.olson@ttu.edu