Chunpeng James Chen

Department of Animal and Poultry Sciences

Virginia Tech

2080 Litton Reaves Hall, Blacksburg, VA 24061, USA Email: niche@vt.edu | Lab website: vt-ads.github.io

EDUCATION

2016-2021 Washington State University, Pullman, WA, USA

Crop Science, Ph.D.

Dissertation: A Paradigm Shift in Breeding: From Genomics to Phenomics

2010-2014 National Taiwan University, Taipei, Taiwan

Agronomy, B.S.

PROFESSIONAL POSITIONS

01/2022 ~ Present	Assistant Professor of Animal Data Sciences
	Department of Animal and Poultry Sciences, Virginia Tech,
	Blacksburg, VA, USA
$03/2021 \sim 12/2021$	Postdoctoral Associate
	Department of Animal Science, University of California, Davis,
	Davis, CA, USA
$08/2016 \sim 12/2020$	Graduate Research Assistant
	Department of Crop and Soil Sciences, Washington State University,
	Pullman, WA, USA
$06/2019 \sim 08/2019$	Biostatistician Intern
	Department of Research and Development, BASF,
	West Sacramento, CA, USA
$04/2016 \sim 06/2016$	Research Assistant
	Institute of Plant and Microbial Biology, Academia Sinica, Taiwan
$10/2015 \sim 03/2016$	Data Analyst
	Yu-Shun International Cultural CO., LTD, Taipei, Taiwan
$10/2014 \sim 09/2015$	Corporal
	564 Armor Brigade, 8th Army Corps, Republic of China Army,

TEACHING EXPERIENCE

Taiwan

$06/2021 \sim 12/2021$	Undergrad/Graduate Student Training
	Department of Animal Science, University of California Davis
	Davis, CA, USA
05/2018	Guest Lecture
	Introduction to Machine Learning and Ensemble Methods
	Statistical Genomics (CropS 545) at Washington State University
02/2017	Guest Lecture
	Principal Component Analysis

Statistical Genomics (CropS 545) at Washington State University

Spring 2017 **Teaching Assistant**

Statistical Genomics (CropS 545) at Washington State University

PROFESSIONAL	LACTIVITIES
Oral Presentation	
Apr 2021	WSU Crop Sciences Exit Seminar
•	A Paradigm Shift in Breeding: From Genomics to Phenomics
Feb 2021	WSU Plant Sciences Retreat 2021
	GRID: a Python Package for Aerial High-Throughput Phenotyping
Jan 2020	Plant and Animal Genome 2020
J	GRID: a Python Package for Aerial High-Throughput Phenotyping
Jan 2019	Wheat Quality Council 2019
•	Toward Instant, Non-Destructive Prediction of Wheat Hagberg-Perten
	Falling Number
Jan 2019	Plant and Animal Genome 2019
	iPat: A Genomics Analysis Tool for Everyone
Jan 2019	Plant and Animal Genome 2019
	Toward Instant, Non-Destructive Prediction of Wheat Hagberg-Perten
	Falling Number
Mar 2018	WSU Plant Sciences Retreat 2018
	GWAS and GS are as easy as clicking and dragging with iPat.
Feb 2018	World Congress on Genetics Applied to Livestock Production 2018
	GWAS and GS are as easy as clicking and dragging with iPat.
Jan 2018	Plant and Animal Genome 2018
	iPat: Intelligent Prediction and Association Tool for Genomic Research
Nov 2017	WSU Crop Sciences Proposal Seminar
	Application of Random Forest in Genomics Selection
May 2017	Plant and Animal Genome Asia 2017
	iPat, a Versatile Tool for Genomics Studies
May 2017	Plant and Animal Genome Asia 2017
	Segregation Analysis and Its Implementation in iPat
Poster Presentation	
Mar 2018	WSU Plant Science Symposium 2018
	iPat: intelligent prediction and association tool for genomic research
Jan 2018	Plant and Animal Genome 2018
	iPat: Intelligent Prediction and Association Tool for Genomic Research
May 2017	Plant and Animal Genome Asia 2017
	Intelligent prediction and association tool for genomic research
Jan 2017	Plant and Animal Genome 2017
	iPat: Interface of Prediction and Association for Genomics

Ad Hoc Reviewer (3)

Frontier in Genetics (1), Bioinformatics (1), Crop & Pasture Science (1), and PLoS One (1)

SOFTWARE DEVELOPED

- XSimV2: A fast and user-friendly tool to simulate sequence data and complicated pedigree structures (https://github.com/reworkhow/XSim.jl)
- GRID: A Python Package for Aerial High-Throughput Phenotyping (https://zzlab.net/GRID and https://poissonfish.github.io/GRID/index.html)
- iPat: Intelligent Tool for Prediction and Association (https://zzlab.net/iPat and https://zzlab.net/iPat and https://zzlab.net/iPat and https://zzlab.net/iPat and https://zzlab.net/iPat and <a href="https://zzlab.net/iPat</

FOUNDING RECEIVED

- Washington Wheat Foundation, \$3,238, Principal Investigator, "Instant and non-destructive prediction of wheat Hagberg falling number from hyperspectral imaging by using parallel computation with graphics processing units (GPU)", November 8, 2018 November 7, 2019,
- Travel grant of International Conference of Plant and Animal Genome, 2018, \$1,300

PUBLICATIONS (10)

First / Co-first Authorship (5)

- 1. **Chen, C.J.** and Zhang, Z. (2020) GRID: A Python Package for Field Plot Phenotyping Using Aerial Images. Remote Sensing, 12, 1697.
- 2. Liu, L., Zhou, J., **Chen, C.J.** et al. (2020) GWAS-based identification of New Loci for Milk Yield, Fat, and Protein in Holstein Cattle. <u>Animals 10, 2048</u>.
- 3. Zhou, J., Liu, L., **Chen, C.J.** et al. (2019) Genome-wide association study of milk and reproductive traits in dual-purpose Xinjiang Brown cattle. <u>BMC Genomics 20, 827</u>.
- 4. **Chen C.J.**, Zhang, Z. (2018) iPat: intelligent prediction and association tool for genomic research. *Bioinformatics*, 34, 1925-1927.
- 5. **Chen C.J.**, Zhang, Z. (2018) GWAS and GS are as easy as clicking and dragging with iPat. The World Congress on Genetics Applied to Livestock Production.

Co-Authorship (1)

1. Tang, Z., Parajuli, A., **Chen, C.J.,** Hu, Y., Revolinski, S., Medina, C.A., Lin, S., Zhang, Z., and Yu, L.-X. (2021) Validation of UAV-based alfalfa biomass predictability using photogrammetry with fully automatic plot segmentation. <u>Scientific Reports 11, 3336.</u>

Under Publication (4)

- 1. **Chen, C.J.**, Garrick, D., Fernando, R., Karaman, E., and Cheng, H. (2021). XSim Version 2: Simulation of Modern Breeding Programs. (**Under review by G3**)
- 2. **Chen, C.J.,** Rutkoski, J., Schnable, J., Murray, S., Wang, L., Jin, X., Stich, B., Crossa, J., Hayes, B, and Zhang, Z. (2021). Harnessing Agronomics Through Genomics and Phenomics in Plant Breeding: A Review. (**Accepted by Plant Breeding Review**)
- 3. **Chen, C.J.**, Morota, G., and Cheng, H. (2021). Tracking Livestock Behaviors with Labor-Free Approach.
- 4. **Chen, C.J.** and Cheng, H. (2021). LMMonBoard: An Interactive Dashboard for Visualizing Mixed Models in Quantitative Genetics

PROGRAMMING PROFESSION

Python, R, Julia, Java, C, C++

AWARDS

2018-2021	Roscoe and Frances Cox Scholarship
2018-2020	Vogel Washington State Crop Improvement Association Scholarship
Oct, 2017	2 nd place for the 3 rd Annual WSU Hardware Hackathon
Jun, 2017	Winner of Student Workshop Competition at PAG Asia