# **Chun-Peng James Chen**

<b>CONTACT INFORMAT</b> 2475 Litton Reaves hal		<i>Email</i> : niche@vt.edu	
175 West Campus DriveCVirginia TechG		ORCID: 0000-0002-2018-0702	
		Google Scholar: jYRGjLgAAAAJ	
		Website: vt-ads.github.io	
SUMMARY	I am an assistant professor of animal data sciences in the School of Animal Sciences (SAS) at Virginia Tech. My research interests include deep learning, statistical genetics, and software development. I am particularly interested in developing data-driven stategies for solving real-world problems in precision livestock farming. My research also tightly integrates with indutrial applications. Currently, I am developing computer vision algorithms to model animal behaviors for better animal welfare and management.		
EDUCATION	Washington State University, Pullman, Washington, US Ph.D., Crop Science	SA 08/2016 - 05/2021	
	<ul> <li><i>Thesis</i>: A Paradigm Shift in Breeding: From Genomics to Phenomics</li> <li><i>Advisor</i>: Dr. Zhiwu Zhang</li> </ul>		
	<b>National Taiwan University,</b> Taipei, Taiwan B.S., Agronomy	09/2010 - 06/2014	
PROFESSIONAL POSITIONS	<b>Assistant Professor</b> (70% Research and 30% Teaching School of Animal Sciences, Virginia Tech Blacksburg, Virginia, USA	) 07/2022 - Present	
	Affiliated Faculty Member Center for Advanced Innovation in Agriculture (CAIA), Vir Blacksburg, Virginia USA	01/2022 - Present ginia Tech	
	<b>Assistant Professor</b> (70% Research and 30% Teaching Department of Animal and Poultry Sciences, Virginia Tec Blacksburg, Virginia, USA		
	<b>Postdoctoral Associate</b> ( <i>Advisor</i> : Dr. Hao Cheng) Department of Animal Science, University of California, D Davis, California, USA	03/2021 - 12/2021 Davis	
	Graduate Research Assistant Department of Crop and Soil Sciences, Washington State Pullman, Washington, USA	08/2016 - 12/2020 e University	
	<b>Biostatistician Intern</b> Department of Research and Development, BASF West Sacramento, California, USA	06/2019 - 08/2019	
	<b>Research Assistant</b> Institute of Plant and Microbial Biology, Academia Sinica Taipei, Taiwan	04/2016 - 06/2016	
	Data Analyst Yu-Shun International Cultural CO., LTD	10/2015 - 03/2016	

Taipei, Taiwan

Corporal

564 Armor Brigade, 8th Army Corps, Republic of China Army Taipei, Taiwan

10/2014 - 09/2015

#### **EDITORIAL** Ad Hoc Reviewer ACTIVITIES · Number of manuscripts reviewed per journal: Bioinformatics (1), Crop & Pasture Science (1), Frontier in Genetics (1), Journal of Animal Science (1), Journal of Dairy Science (3), PLoS One (1)

· Number of manuscripts reviewed per year: 2018(1), 2019(2), 2021(1), 2022(2), 2023(2)

#### **GRANTS DIRECTED OR CO-DIRECTED**

# **External Competitive Grants**

United State Department Agriculture (USDA) - NIFA - DSFAS 09/2023 - 08/2028 Principal Investigator \$649,741 2022-11638 Acoustic data as a novel trait to manage welfare and environmental impact in precision cow farming.

### Washington Wheat Foundation

11/2018 - 07/2019 Principal Investigator \$3,238 Instant and non-destructive prediction of wheat Hagberg falling number from hyperspectral imaging by using parallel computation with graphics processing units (GPU)

# Internal Competitive Grants

2023 CALS Integrated Internal Competitive Grants Program	07/2023 - 06/2024
Principal Investigator	\$30,000
Ant Detective: Automated Computer-Vision-Based Kit to Prevent the Species	Spread of Invasive

#### Center for Advanced Innovation in Agriculture (CAIA) 03/2022 - 06/2022 Principal Investigator \$4,000 Agricultural Leadership for Cyberbiosecurity: A Teaching Case Study

#### **Peer-Reviewed Research Journal Articles**

- Chen, Chun-Peng James, Y. Hu, X. Li, C. F. Morris, S. Delwiche, A. H. Carter, C. Steber, and Z. Zhang, 2023 An independent validation reveals the potential to predict Hagberg–Perten falling number using spectrometers. The Plant Phenome Journal 6: e20070
- 11. Massahiro Yassue, R., G. Galli, C.-P. James Chen, R. Fritsche-Neto, and G. Morota, 2023 Genome-wide association analysis of hyperspectral reflectance data to dissect the genetic architecture of growth-related traits in maize under plant growth-promoting bacteria inoculation. Plant Direct **7**: e492
- Chen, C. J., J. Rutkoski, J. C. Schnable, S. C. Murray, L. Wang, X. Jin, B. Stich, J. Crossa, B. J. Hayes, and Z. Zhang, 2023 Role of the Genomics-Phenomics-Agronomy Paradigm in Plant Breeding. In *Plant Breeding Reviews*, volume 46, pp. 622–668, WI-LEY
- 9. Chen, C. and G. Ferreira, 2022 Evaluation of walking activity data during pregnancy as an indicator of pregnancy loss in dairy cattle. JDS Communications p. S2666910222001466
- 8. Chen, C. P. J., G. Morota, K. Lee, Z. Zhang, and H. Cheng, 2022 VTag: a semisupervised pipeline for tracking pig activity with a single top-view camera. Journal of Animal Science 100
- Chen, C. J., D. Garrick, R. Fernando, E. Karaman, C. Stricker, M. Keehan, and H. Cheng, 2022a XSim version 2: simulation of modern breeding programs. G3 Genes|Genomes| Genetics 12
- Hu, Y., S. M. Sjoberg, Chen, C. J., A. L. Hauvermale, C. F. Morris, S. R. Delwiche, A. E. Cannon, C. M. Steber, and Z. Zhang, 2022 As the number falls, alternatives to the Hagberg–Perten falling number method: A review. Comprehensive Reviews in Food Science and Food Safety 21: 2105–2117
- 5. Tang, Z., A. Parajuli, **Chen, C. J.**, Y. Hu, S. Revolinski, C. A. Medina, S. Lin, Z. Zhang, and L. X. Yu, 2021 Validation of UAV-based alfalfa biomass predictability using photogrammetry with fully automatic plot segmentation. Scientific Reports **11**: 3336
- 4. Chen, C. J. and Z. Zhang, 2020 GRID: A Python Package for Field Plot Phenotyping Using Aerial Images. Remote Sensing **12**: 1697
- Liu, L., J. Zhou, Chen, C. J., J. Zhang, W. Wen, J. Tian, Z. Zhang, and Y. Gu, 2020 GWAS-Based Identification of New Loci for Milk Yield, Fat, and Protein in Holstein Cattle. Animals 10: 2048
- Zhou, J., L. Liu, Chen, C. J., M. Zhang, X. Lu, Z. Zhang, X. Huang, and Y. Shi, 2019 Genome-wide association study of milk and reproductive traits in dual-purpose Xinjiang Brown cattle. BMC Genomics 20: 827
- 1. Chen, C. J. and Z. Zhang, 2018b iPat: intelligent prediction and association tool for genomic research. Bioinformatics **34**: 1925–1927

#### **Peer-Reviewed Conference Proceedings**

 Chen, C. J., G. Morota, and H. Cheng, 2022b VTag: Automatic pipeline to annotate video data for pig phenomics studies. The 12th World Congress of Genetics Applied to Livestock Production, Rotterdam, The Netherlands 1. **Chen, C. J.** and Z. Zhang, 2018a GWAS and GS Are as Easy as Clicking and Dragging with iPat. The 11th World Congress of Genetics Applied to Livestock Production, Auckland, New Zealand

#### DEVELOPED SOFTWARE

- 4. VTag: a semi-supervised pipeline for tracking pig activity with a single top-view camera
  - Publication on JAS: https://doi.org/10.1093/jas/skac147
  - GitHub Repository: https://github.com/vt-ads/vtag
- 3. XSimV2: A fast and user-friendly tool to simulate sequence data and complicated pedigree structures
  - Publication on G3: https://doi.org/10.1093/g3journal/jkac032
  - GitHub Repository: https://github.com/reworkhow/XSim.jl
  - Documentation: https://reworkhow.github.io/XSim.jl/index.html
- 2. GRID: A Python Package for Aerial High-Throughput Phenotyping
  - Publication on Remote Sensing: https://doi.org/10.3390/rs12111697
  - GitHub Repository: https://github.com/Poissonfish/GRID
  - Software Page: http://zzlab.net/GRID
  - Documentation: https://poissonfish.github.io/GRID/index.html
- 1. iPat: Intelligent Tool for Prediction and Association
  - Publication on *Bioinformatics*: https://doi.org/10.1093/bioinformatics/bty015
  - GitHub Repository: https://github.com/Poissonfish/iPat
  - Software Page: http://zzlab.net/iPat
  - Documentation: https://poissonfish.github.io/iPat/index.html

#### PRESENTATIONS

# **Conference Presentations**

12.	American Dairy Science Association (ADSA) Annual Meeting Ottawa, Canada	06/2023
	A computer vision strategy to alleviate cow mastitis and improve dairy farm sustainability	
11.	The 12th World Congress on Genetics Applied to Livestock Production (WCGALP) Rotterdam, The Netherlands VTag: Automatic pipeline to annotate video data for pig phenomics studies	07/2022
10.	American Dairy Science Association (ADSA) Annual Meeting Kansas City, Missouri, USA Evaluation of Walking Activity Data During Pregnancy as an Indicator of Pregnancy Loss in Dairy Cattle	06/2022
9.	National Animal Genome Research Program (NRSP8) San Diego, California, USA VTag: a Semi-Supervised Pipeline for Tracking Pig Activity with a Single Top-View Camera	04/2022
8.	Plant and Animal Genome (PAG) XXVIII San Diego, California, USA GRID: a Python Package for Aerial High-Throughput Phenotyping	01/2020
7.	<b>Wheat Quality Council</b> Spokane, Washington, USA Toward Instant, Non-Destructive Prediction of Wheat Hagberg-Perten Falling	01/2019 Number
6.	<b>Plant and Animal Genome (PAG) XXVII</b> San Diego, California, USA Toward Instant, Non-Destructive Prediction of Wheat Hagberg-Perten Falling	01/2019 Number
5.	Plant and Animal Genome (PAG) XXVII San Diego, California, USA iPat: A Genomics Analysis Tool for Everyone	01/2019
4.	The 11th World Congress on Genetics Applied to Livestock Production (WCGALP) Auckland, New Zealand GWAS and GS are as easy as clicking and dragging with iPat	02/2018
3.	Plant and Animal Genome (PAG) XXVI San Diego, California, USA iPat: Intelligent Prediction and Association Tool for Genomic Research	01/2018

2. Plant and Animal Genome (PAG) Asia Seoul. korea	05/2017
iPat, a Versatile Tool for Genomics Studies	
1. <b>Plant and Animal Genome (PAG) Asia</b> Seoul, korea Segregation Analysis and Its Implementation in iPat	05/2017
Intramural Seminars	
Virginia Tech	
6. Animal and Poultry Sciences Seminar Leveraging Activity Data to Improve Pregnancy Diagnoses and Herd Assessments	03/2022
UC Davis	
5. <b>Animal Science Seminar</b> VTag: a Semi-Supervised Pipeline for Tracking Pig Activity with a Single Top-View Camera	12/2021
Washington State University	
<ol> <li>Crop Sciences Ph.D. Exit Seminar A Paradigm Shift in Breeding: From Genomics to Phenomics</li> </ol>	04/2021
3. <b>Plant Sciences Retreat</b> GRID: a Python Package for Aerial High-Throughput Phenotyping	02/2021
2. <b>Plant Sciences Retreat</b> GWAS and GS are as easy as clicking and dragging with iPat	03/2018
1. Crop Sciences Ph.D. Proposal Seminar Application of Random Forest in Genomics Selection	11/2017

### TEACHING

# Lead Lectures

APSC-5984 Special Study: Agriculture Data Science Virginia Tech	01/2023 - 05/2023
Short Courses and Workshop	
<b>Modeling Methods: Three types of risks in model validation</b> Co-instructors: Dr. Robert Tempelman and Dr. Mark Hanigan National Animal Nutrition Program (NANP): Journal of Dairy Science, and JDS Communications Joint Workshop	06/2023
Modern Programming in Genome to Phenome Co-instructors: Dr. Rohan Fernando and Dr. Hao Cheng University of California, Davis	08/2022
Guest Lectures	
<b>CropS 545 Statistical Genomics</b> Introduction to Machine Learning and Ensemble Methods Instructor: Dr. Zhiwu Zhang Washington State University	05/2018
<b>CropS 545 Statistical Genomics</b> Principal Component Analysis Instructor: Dr. Zhiwu Zhang Washington State University	02/2017