# **JONGHYUN YUN**

Data Scientist, PhD in Statistics

@ jonghyun.yun@gmail.com

% jyun.netlify.app

github.com/jonghyun-yun

### **EMPLOYEMENT HISTORY**

#### **Data Scientist**

#### Institute of Statistical Data Intelligence

## 09/2019 - Present

Mansfield, TX, USA

- Developing and/or applying cutting edge ML for prediction modeling, ensemble methods, Bayesian models, time series, causal inference, visualization, segmentation for big data.
- Applying NLP and survival model to analyze timestamped sequence of action data (log data) using TensorFlow,
   Keras and Numpy.
- Developing network modeling frameworks to discover dynamic interaction b/w customers and merchandise.
- Parallel programming using C/C++ for complex Bayesian inference.
- Processing, cleansing and validating the integrity of data using SQL and Pandas.
- Presenting analysis and visualization using R and Python, and developing software packages.

**Assistant Professor of Statistics** 

Department of Mathematics, University of Texas at Arlington

**1** 09/2016 - 08/2019

Arlington, TX, USA

Assistant Professor of Statistics

Department of Mathematical Sciences, University of Texas at El Paso

**(1)** 08/2015 - 06/2016

**♀** El Paso, TX, USA

Postdoctoral Researcher

Quantitative Biomedical Research Center, University of Texas Southwestern Medical Center

**(1)** 09/2012 - 07/2015

Dallas, TX, USA

# **EDUCATIONAL HISTORY**

PhD in Statistics

Department of Statistics, University of Illinois at Urbana-Champaign

**1** 09/2006 - 08/2012

♦ Champaign, IL, USA

Dissertation: Ensemble Filtering of State Space Models

Minor in Mathematics

### **STRENGTHS**

Project leadership Presentaton Interdisciplinary collaboration Mentorship		
Advanced statistical analysis   Machine learning   Predictive modeling   Dimension reduction   Visualization		
Time Series Hidden Markov model Natural language processing Sparse models Bayesian inference		
Causal inference   Monte Carlo method   Multiple hypothesis testing   Biostatistics   Bioinformatics		
Next generation sequencing Smart infrastructure Item response theory Network model Process data model		
R Python Git SQL C/C++ MATLAB Cloud computing Parallel programming SPSS SAS		
Shell script FTEX Markdown Hugo MS Office		

# FEATURED ON-GOING PROJECTS

### Customer behavioral analysis (Github repo)

- Developing a novel machine learning approach to analyze timestamped sequence of action data (a.k.a. process data) leveraging natural language processing and survival models.
- Identifying behavioral differences in consumer buying decision-making processes.
- Presenting analysis and visualization using R and python, and developing software packages

### Network dependence analysis using time to events (Github repo)

- Developing Cox model equipped with latent space to discover patterns between connection time and outcome in bipartite network models.
- Implementing developed model has great potential applications including identifying relationship between customer shopping time and outcome (buy or not) and modeling test-taker's proficiency and time used to solve times in educational assessments such as Duolingo.

# **PUBLISHED INTELLECTUAL CONTRIBUTIONS**

- 1. Yun, J., Ryu, K. R. & Ham, S. Spatial Analysis Leveraging Machine Learning and GIS of Socio-Geographic Factors Affecting Cost Overrun Occurrence in Roadway Projects. *Automation in Construction* **133**, 104007 (2022).
- 2. Yun, J., Kang, S., Tehrani, A. D. & Ham, S. Image Analysis and Functional Data Clustering for Random Shape Aggregate Models. *Mathematics* **8**, 1903 (2020).
- 3. Yun, J., Shin, M., Jin, I. H. & Liang, F. Stochastic Approximation Hamiltonian Monte Carlo. *Journal of Statistical Computation and Simulation* **90**, 3135–3156 (2020).
- 4. Nam, J. H., Yun, J., Jin, I. H. & Chung, D. hubViz: A Novel Tool for Hub-Centric Visualization. *Chemometrics and Intelligent Laboratory Systems* **203**, 104071 (2020).
- 5. Cai, L., Li, Q., Du, Y., Yun, J., Xie, Y., DeBerardinis, R. J. & Xiao, G. Genomic Regression Analysis of Coordinated Expression. *Nat Commun* **8**, 2187 (2017).
- 6. Yun, J., Yang, F. & Chen, Y. Augmented Particle Filters. *Journal of the American Statistical Association* **112**, 300–313 (2017).
- 7. Chen, B., Yun, J., Kim, M. S., Mendell, J. T. & Xie, Y. PIPE-CLIP: A Comprehensive Online Tool for CLIP-seq Data Analysis. *Genome Biol* **15**, R18 (2014).
- 8. Kwon, I., Xiang, S., Kato, M., Wu, L., Theodoropoulos, P., Wang, T., Kim, J., Yun, J., Xie, Y. & McKnight, S. L. Poly-Dipeptides Encoded by the C9orf72 Repeats Bind Nucleoli, Impede RNA Biogenesis, and Kill Cells. *Science* **345**, 1139–45 (2014).
- 9. Yun, J., Wang, T. & Xiao, G. Bayesian Hidden Markov Models to Identify RNA-Protein Interaction Sites in PAR-CLIP. *Biometrics* **70**, 430–440 (2014).

#### **Non-Refereed Articles**

1. Yun, J. & Chen, Y. Comments on "Particle Markov Chain Monte Carlo Methods" by C. Andrieu, A. Doucet, and R. Hollenstein. *Journal of the Royal Statistical Society Series B-Statistical Methodology* **72**, 332–333 (2010).

#### **Book Sections**

1. Wang, T., Yun, J., Xie, Y. & Xiao, G. in Hidden Markov Models 177–184 (Humana Press, New York, NY, 2017).

#### **Software**

- 1. Yun, J. Statistical Data Intelligence Tools for Cost-Overrun Analysis of Roadway Construction Projects 2021. github.com/jonghyun-yun/dico.
- 2. Yun, J. TEMPEST: Latent Space Competing Risk Model for Accuarcy and Reponse Time Data https://github.com/ Jonghyun-Yun/TEMPEST.
- 3. Yun, J. Process Data Modeling for PIACC Data 2021+. https://github.com/Jonghyun-Yun/proda.
- 4. Alvarez, H. & Yun, J. Baseball Statistics Collecting Functions from HTML Tables 2017. https://github.com/jonghyun-yusbrscrap.git.
- 5. Yun, J. A MATLAB Toolbox to Identify RNA-protein Binding Sites in HITS-CLIP 2013. https://qbrc.swmed.edu/labs/xiaoxie/download/README1.pdf.
- 6. Yun, J. R Package for PAR-CLIP Analysis 2013. https://qbrc.swmed.edu/labs/xiaoxie/download/README2.pdf.

### **Working Papers**

- 1. Jin, I. H., Jeon, M., Yun, J., Schweinberger, M. & Lin, L. Hierarchical Network Item Response Modeling for Discovering Differences Between Innovation and Regular School Systems in Korea. *Journal of the Royal Statistical Society: Series C (Applied Statistics)* (2020+). Invited for revision.
- 2. Yun, J., Jin, I. H. & Jeon, M. Latent Space Competing Risk Modeling for Accuacy and Response Time Based on Tests. *Journal of the American Statistical Association* (2021+). To be submitted.
- 3. Yun, J., Ick Hoon, J. & Minjeong, J. Analysis of Time-Stamped Action Sequences (2021+).
- 4. Yun, J., Wang, T., Wang, X. & Xiao, G. Identification of RNA-protein Binding Sites in HITS-CLIP Using Heterogeneous Logit Models via Semi-Supervised Learning (2021+).
- 5. Yun, J. & Chen, Y. Localized Agumented Particle Filters (2021+).
- 6. Yun, J., Wang, T., Wang, X. & Xiao, G. The Identification of Differential Binding Sites in CLIP-seq.

### **PRESENTATIONS**

#### **Invited Talks**

- 11/2021 "Latent Space Accumulator Model for Analyzing Bipartite Networks with Connection Times and Its Applications to Item Response Data", Autumn annual conference of the Korean statistical society, virtual.
- 02/2017 "Integrative modeling approaches for next-generation sequencing data", *Colloquim Series*, Texas A&M University-Commerce.
- 06/2016 "Model based identification of RNA-protein binding sites", Bioinformatics Session, *International Workshop on Applied Probability*, Toronto, ON, Canada.
- 10/2015 "Comparative analysis of CLIP-seq under multiple experimental conditions", *Border Biomedical Research Center Seminar*, *UT El Paso*, El Paso, TX, USA.
- 08/2014 "Statistical strategies for identification of the RNA-protein binding site in CLIP-seq", Biometrics Section, 2014 Joint Statistical Meetings, Boston, NY, USA.
- 10/2014 "Statistical models to identify RNA-protein binding sites from CLIP experiments", *Computational and Systems Biology Seminar*, UT Southwestern, Dallas, TX, USA.
- 10/2011 "Augmented particle filters", *Robert Bohrer Student Workshop in Statistics*, University of Illinois at Urbana-Champaign, Champaign, IL, USA.

### **Poster Presentation**

02/2014 "Identification for RNA-protein binding sites in CLIP-seq", 7th Annual Bayesian Biostatistics and Bioinformatics Conference, Houston, TX, USA.

# PROFESSIONAL AND UNIVERSITY SERVICE

#### **Professional Service**

06/2016 Co-chair, Bioinformatics session at 2016 International Workshop on Applied Probability at Toronto, ON, Canada.

### **University Service (UTA)**

09/2017 - 08/2019	Department advisory committee.
09/2016 - 08/2019	Math preliminary exam B subcommittees.
01/2017 - 05/2017	Undergraduate affairs committee.
01/2019 - 08/2019	College of Science Data science working group.
04/2018	Judge, College of Science Aces Research Symposium.

### **University Service (UTEP)**

Spring 2016 Math Club Zero committee

### Referee/Reviewer Work (Journals)

• Journal of the American Statistical Association, Journal of Computational and Graphical Statistics, Computational and Mathematical Methods in Medicine, Journal of Statistical Software, Journal of Probability and Statistics, Bayesian Analysis, International Journal of Data Science, Genes, Mathematics, International Journal of Environment Research and Public Health, Antibiotics, Axioms, Healthcare

# **TEACHING ACTIVITIES**

### **University of Texas at Arlington**

Spring 2019	MATH6312 - Data Mining (10 students)
Fall 2018	MATH3316 - Statistical Inference (57 students)
Spring 2018	MATH5358 - Regression Analysis (13 students)
Fall 2017	MATH5312 - Mathematical Statistics I (12 students)
Spring 2017	MATH5392 - Selected Topics in Mathematics (Data Mining) (12 students)
	MATH5313 - Mathematical Statistics II (6 students)
Fall 2016	MATH5312 - Mathematical Statistics I (14 students)

### University of Texas at El Paso

```
Spring 2016 STAT5474 - Introduction to Data Mining (14 students)

Fall 2015 STAT5354 - Post-genomic Analysis (5 students)

BINF5113 - Math Seminar for Bioinformatics (4 students)
```

### University of Illinois at Urbana-Champaign

Spring 2012	STAT200 - Statistical Analysis (51 students)
Summer 2011	STAT100 - Statistics (30 students)
01/2010 - 05/2011	STAT400-Statistics and Probability I (Discussion Section Leader)
	Spring 2010 (59 students), Fall 2010 (60 students), and Spring 2011 (93 students)
08/2006 - 12/2009	Teaching Assistant: STAT100-Statistics, STAT400-Statistics and Probability I, STAT410-
	Statistics and Probability II, STAT424-Analysis of Variance, STAT429-Time Series Analysis,
	STAT510- Mathematical Statistics I, and STAT511-Mathematical Statistics II.

### **Yonsei University**

12/2005	Preliminary Calculus
03/2005 - 12/2005	Discussion Section Leader: STA2101-Calculus (65 students) and STA2102-Linear Algebra (67 students).
03/2004 - 12/2004	Teaching Assistant: STA1001-Introductory Statistics, STA1001-Introductory Statistics, STA3102-Multivariate Statistical Analysis, and BC682-Statistical Methods for Behavioral Sciences.

# **DIRECTED STUDENT LEARNING**

### **Graduate Supervised Research**

09/2017 - 09/2019	Anthony Thomas (Statistics, UT Arlington)
	Project: Bayesian hierarchical dynamic factor models
09/2017 - 12/2017	Mario Garza (M.S. Statistics, UT Arlington)
	Project: Forecasting sales using a finite-state HMM: an inventory control exercise

### **5 M.S. Student Committees**

09/2016 - 08/2019	Daniel Sang Le, Nidhi Kiran Dawda, Zachary Loucks, Hongbo Yu
	Statistics, UT Arlington
09/2015 - 08/2016	Tun-Lee Ng
	Statistics, UT El Paso

### **6 Ph.D. Student Committees**

09/2016 – 08/2019 Souad Sosa, Izzet Sozucok, Geoffrey Schuette, Yi Liu, Mahmoud Jawad, Piyachart Wiangnak Statistics, UT Arlington

## **Undergraduate Supervised Research**

Spring 2018 Henry Alvarez (Mathematics, UT Arlington)

Project: Developing a software package to collect baseball statistics