Jean Feng

Position

July 2020– **Assistant Professor In-Residence**, Department of Epidemiology and Biostatistics, Present *University of California, San Francisco*

Education

Sept 2015– PhD, Biostatistics, University of Washington, Seattle, WA.

June 2020 Advisors: Noah Simon, Frederick Albert Matsen IV

2012–2013 MS, Computer Science, Stanford University, Stanford, CA.

2009–2013 BS, Computer Science, Stanford University, Stanford, CA.

Publications

<u>Jean Feng</u> and Noah Simon. Ensembled sparse-input hierarchical networks for high-dimensional datasets. *Statistical Analysis and Data Mining*, 2022.

<u>Jean Feng</u>, Alexej Gossmann, Berkman Sahiner, and Romain Pirracchio. Bayesian logistic regression for online recalibration and revision of risk prediction models with performance guarantees. *Journal of the American Medical Informatics Association*, 2022.

Daniel Lazzareschi, Ravindra L Mehta, Laura M Dember, Juliane Bernholz, Alparslan Turan, Amit Sharma, Sachin Kheterpal, Chirag R Parikh, Omar Ali, Ivonne H Schulman, Abigail Ryan, Jean Feng, Noah Simon, Romain Pirracchio, Patrick Rossignol, and Matthieu Legrand. Overcoming barriers in the design and implementation of clinical trials for acute kidney injury: a report from the 2020 kidney disease clinical trialists meeting. *Nephrol. Dial. Transplant*, 2022.

<u>Jean Feng</u>, Arjun Sondhi, Jessica Perry, and Noah Simon. Selective prediction-set models with coverage guarantees. *Biometrics*, 2021.

<u>Jean Feng, Scott Emerson, and Noah Simon.</u> Approval policies for modifications to machine Learning-Based software as a medical device: A study of bio-creep. *Biometrics*, 2021.

<u>Jean Feng</u>, William S DeWitt, Aaron McKenna, Noah Simon, Amy Willis, and Frederick A Matsen. Estimation of cell lineage trees by maximum-likelihood phylogenetics. *Annals of Applied Statistics*, 2021.

<u>Jean Feng.</u> Learning to safely approve updates to machine learning algorithms. <u>Proceedings of the Conference on Health, Inference, and Learning</u>, 2021.

Brian D Williamson and Jean Feng. Efficient nonparametric statistical inference on population feature importance using shapley values. *International Conference on Machine Learning*, 2020.

<u>Jean Feng</u> and Noah Simon. An analysis of the cost of hyper-parameter selection via split-sample validation, with applications to penalized regression. *Statistica Sinica*, 2020.

Jean Feng, David A Shaw, Vladimir N Minin, Noah Simon, and Frederick A Matsen, IV. Survival analysis of DNA mutation motifs with penalized proportional hazards. *Ann. Appl. Stat.*, 2019.

Kristian Davidsen, Branden J Olson, William S DeWitt, 3rd, <u>Jean Feng</u>, Elias Harkins, Philip Bradley, and Frederick A Matsen, 4th. Deep generative models for T cell receptor protein sequences. *Elife*, 2019.

Jean Feng, Brian Williamson, Noah Simon, and Marco Carone. Nonparametric variable importance using an augmented neural network with multi-task learning. *International Conference on Machine Learning*, 2018.

<u>Jean Feng</u> and Noah Simon. Gradient-based regularization parameter selection for problems with nonsmooth penalty functions. *J. Comput. Graph. Stat.*, 2018.

Pre-prints

<u>Jean Feng</u> and Noah Simon. Sparse-Input neural networks for high-dimensional nonparametric regression and classification. *arXiv*, 2019.

Funding History

2020-2022 UCSF-Stanford CERSI Program; Role: PI

"Safe algorithmic change protocols for modifications to AI/ML-based Software as a Medical Device."

\$100,946 in direct costs

Presentations

Invited Oral Presentations and Seminars

- 2022 Approval policies for modifications to Machine Learning-Based Software as a Medical Device: A study of bio-creep, International Biometrics Society Journal Club
- 2021 Safe approval policies for continual learning systems in healthcare, Brown University
- 2021 Bayesian logistic regression for online recalibration and revision of risk prediction models with guarantees, ASA Biopharmaceutical Section Regulatory-Industry Statistics Workshop
- 2021 Variable Selection and Architecture Search for Neural Networks, ASA Statistical Learning and Data Science Webinar
- 2021 Bayesian logistic regression for online recalibration and revision of risk prediction models with guarantees, Western North American Region (WNAR) Annual Meeting
- 2021 Learning to safely approve updates to machine learning algorithms, ACM Conference on Health, Inference, and Learning
- 2021 Safe approval policies for continual learning systems in healthcare, University of Waterloo

- 2020 Efficient nonparametric statistical inference on population feature importance using Shapley values, International Conference on Machine Learning
- 2020 Training Procedures and Regulatory Policies for Safe Machine Learning Models in Healthcare, University of California, San Francisco
- 2020 Training Procedures and Regulatory Policies for Safe Machine Learning Models in Healthcare, The University of Texas, MD Anderson Cancer Center
- 2020 Training Procedures and Regulatory Policies for Safe Machine Learning Models in Healthcare, University of California, Irvine
- 2020 Approval policies for modifications to Machine Learning-Based Software as a Medical Device: A study of Bio-creep, International Conference on Health Policy Statistics
- 2019 Sparse-Input Neural Networks for High-dimensional Nonparametric Regression and Classification, Western North American Region (WNAR) Annual Meeting
- 2018 Nonparametric variable importance using an augmented neural network with multitask learning, International Conference on Machine Learning
- 2018 Sparse-Input Neural Networks for High-dimensional Nonparametric Regression and Classification, University of Washington Biostatistics Colloquium
- 2018 Sparse-Input Neural Networks for High-dimensional Nonparametric Regression and Classification, Joint Statistical Meetings
- 2017 Sparse-Input Neural Networks for High-dimensional Nonparametric Regression, ICML Workshop on Principled Approaches to Deep Learning
- 2011 Haptic Belt with Pedestrian Detection, Neural Information Processing Systems
 Contributed Oral Presentations
- 2019 Uncertainty-Aware Black-Box Predictors with Coverage Guarantees, Joint Statistical Meetings

Awards

- 2020 University of Washington Thomas R. Fleming Excellence in Biostatistics Award
- 2020 International Conference on Health Policy Statistics, Student Travel Award For manuscript: Approval policies for modifications to Machine Learning-Based Software as a Medical Device
- 2018 Joint Statistical Meetings Section on Statistical Learning and Data Science, Student Paper Award
 For manuscript: Sparse-input neural networks for high-dimensional nonparametric regression and classification
- 2018 University of Washington Biostatistics Donovan J. Thompson Award for Best Combined Performance on Ph.D. Theory and Applied Qualifying Examinations
- 2015–2017 Big Data for Genomics and Neuroscience Training Grant

Software

EASIER-Net Python and R packages for fitting neural networks for high-dimensional data

https://github.com/jjfeng/easier_net https://github.com/jjfeng/easier_net_R

GapML Python package for analyzing cell-lineage tracing data from GESTALT

https://github.com/matsengrp/gestaltamania

SPINN Python package for estimating sparse-input neural networks

http://github.com/jjfeng/spinn

samm Python package for estimating somatic hypermutation rates of nucleotide motifs

http://github.com/matsengrp/samm

Teaching

Winter 2021 Instructor, Biostat 216: Machine Learning in R for the Biomedical Sciences, UCSF

2020-2021 Instructor, Machine Learning Boot Camp: Analyzing Biomedical and Health Data, Columbia University

2020 Instructor, Supervised statistical learning, 6th Seattle Symposium in Biostatistics

Service

Referee Service

- Nature Medicine
- International Conference on Machine Learning
- International Conference on Learning Representations
- Journal of Computational and Graphical Statistics
- Statistics in Biopharmaceutical Research
- Annals of Applied Statistics
- Annals of Statistics
- Statistics in Medicine
- Neural Networks
- o ASA Section on Statistical Learning and Data Science Student Paper Committee

Conferences

- Joint Statistical Meetings, Session Chair
- Western North American Region (WNAR) Annual Meeting, Session Chair
- Eastern North American Region (ENAR) Annual Meeting, Roundtable leader

University Service

- UCSF Initiative for Digital Transformation in Computational Biology & Health Grant Review Committee, September 2021
- Department of Epidemiology and Biostatistics, Digital Health Initiative Steering Committee, 2020–Present

Work Experience

2019 **Research Intern**, *Insitro*, South San Francisco, CA. Developed statistical models of genomic data.

2012–2015 **Software engineer**, *Coursera*, Mountain View, CA.

Built the professional certificate program and payment system. Technical lead on projects with 3-5 people. Mentored interns and junior engineers.