SONGPENG ZU

Department of Statistics, Harvard University

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EDUCATION

PhD on Control Science and Engineer Tsinghua University 2011.09 - 2017.01 Bachelor on Biology Tsinghua University 2007.09 - 2011.07

RESEARCH EXPERIENCE

Post-Doctoral Fellow Department of Statistics, Harvard University 2019, 09 - Now

Advisor: Jun Liu, PhD

• Algorithm design and data analysis for single-cell sequencing and tumor immunology.

PhD Department of Automation, Tsinghua University 2011.07 - 2017.01

Advisor: Shao Li, PhD

Studied the compound-protein interactions from the machine learning perspective.

- Predicted latent chemogenomic features from drug-target interactions by the global-optimized probabilistic model under EM framework.
- Inferred the binding affinities of compound-protein interactions by the hierarchical Bayesian model under multi-task learning view.
- Analyzed the molecule-based evaluation of clinical drug responses in cancer by ensemble method to solve the high-dimensional small and imbalanced data.

Visiting Scholar Department of Statistics, Harvard University 2014.03 - 2014.09 Advisor: Jun Liu, PhD

• Applied the Bayesian Nonparametric algorithm via sliced inverse model to detect the non-linear relationships on cis-eQTLs.

Undergraduate research training School of Life Science, Tsinghua University 2009 - 2010 Advisor: Li Yu, PhD

• Molecular mechanisms about cell autophagy.

FUNDING

Harvard Data Science Initiative Postdoctoral Fellow Research Fund
 Principle Investigator: Songpeng Zu \$6,933
 Title: Learning Peptide-specific T Cell Receptors in Human Cancers by Deep Neural Network and Structural Modeling

PUBLICATIONS

- Zu S.*, Li S.* (equal contribution) et al., REPlY: representation learning of B cell receptor sequences. Under preparation, 2021
- Zu S.*, Sahu A.* (equal contribution) et al., MSSC: multiple-sample differential expression analysis of single-cell RNA sequencing data. Under preparation, 2021
- Hu Z.*, **Zu S.*** (equal contribution), and Liu J.S. SIMPLEs: single-cell RNA sequencing imputation and cell clustering methods by modeling gene module variation. Accepted by **NAR Genomics and Bioinformatics**, 2020
- Ding Q., Hou S., **Zu S.**, et al., VISAR: an interactive tool for dissecting chemical feature learned by deep neural network QSAR models. **Bioinformatics**, 36 (11), pp3610-3612., 2020.
- Xu X., **Zu S.**, et al., Modeling Attention Flow on Graphs. **NeurIPS 2018** Relational Representation Learning **Workshop**.

- Xu X., Chen L., **Zu S.**, et al., Hulu video recommendation: from relevance to reasoning. **RecSys'18** Proceedings of the 12th **ACM Conference** on Recommender Systems, pages 482-482, 2018
- Xu X., Zu S., et al., Backprop-Q: Generalized Backpropagation for Stochastic Computation Graphs. NeurIPS 2018 Deep Reinforcement Learning Workshop.
- Ding Z., **Zu S.**, and Gu J. Evaluating the molecule-based prediction of clinical drug response in cancer. **Bioinformatics**, 32(19), 2891-2895, 2016.
- **Zu S.**, Chen T., Li S. Global optimization-based inference of chemogenomic features from drug-target interactions. **Bioinformatics**, 31 (15), 2523 2529, 2015.

SOFTWARE

- SIMPLEs: an R package for single cell RNASeq data imputation based on cell similarities and gene correlations.
- GIFT: A C++ package to infer the chemogenomic information based on drug-protein interactions.

INDUSTRIAL EXPERIENCE

Algorithm Expert (P7) City-Brain Lab, DAMO Academy, Alibaba, China 2018.12 - 2019.08

- Studied the deep Poisson-Gamma probabilistic model for traffic prediction.
- Developed one graphical neural network based online pipeline for transportation system with C++ language.

Researcher (IC3) Recommendation Team, HULU Beijing, China 2017.01 - 2018.11

- Applied the recurrent neural networks for online real-time recommendation
- Lead the development of the multimodal algorithm for personalized recommendation by integrating user sequential watch behavior and shows' metadata.
- Studied the stochastic computational graph optimization with reinforcement learning strategies like Q-learning.

TEACHING EXPERIENCE

•	Guest Speaker of Stat 221 at Depeartment of Statistics, Harvard	2020.10.	2020.11
	 Introduction to variational auto-encoder with pytorch. 		
	 Introduction to STAN, a probabilistic programming language. 		
•	Teaching Assistant of Probabilistic Graphical Models for graduate students	2013.09 -	2014.01
•	Teaching Assistant of Systems Biology for undergraduate students	2014.09 -	2015.01

OTHER EXPERIENCE

- Undergraduate Affair Counselor (for scholarship and financial aid assessment) 2011.08 2013.01
- Completed the Full Marathon 2009.10, 2010.10, 2011.10
- The volunteer of 2008 Beijing Olympic Games 2008.08

AWARDS

• National Scholarship for Graduate Students	2015
• Tsinghua Scholarship for Overseas Graduate Studies	2014
• Tsinghua Excellent Undergraduate Affair Counselor	2013
Tsinghua Zhongying Tang Scholarship	2008, 2009, 2010