

SHANE CHU

555 Melville Ave, Apt 3N , St. Louis, MO 63130
+1 917-640-7301 ♦ skchu@wustl.edu

EDUCATION

- Ph.D., Computer Science** Aug 2018 - Present
Washington University in St. Louis
Advisor: Gary D. Stormo
Thesis: Sparse representations for regulatory genomics
- Bachelor of Arts in Mathematics** Aug 2014 - May 2018
University of Kansas

WORK EXPERIENCE

- Predoctoral researcher** Jan 2019- Present
Washington University in St. Louis
- Designed new models based on sparse representations for biological sequences.
 - Designed and implemented training algorithms for large-scale problems in regulatory genomics.
 - Designed and implemented fast inference techniques for
 - finding DNA sequence motifs that may exhibit patterns such as variable spacings, multimeric bindings, multiple binding domains, and alternative structural conformations.
 - finding the composition of the regulatory elements in functional DNA sequences.
 - Revealed new statistically significant motifs that show alternative binding modes in the JASPAR database.
- Graduate research assistant** Jan 2019 - Present
Washington University in St. Louis
- Created several bioinformatics packages in Julia (available from the Julia registry).
 - Developed NGS pipelines using best practices.
- Research assistant** Nov 2017 - May 2018
Center for Remote Sensing of Ice Sheets (CReSIS)
- Hyperparameter optimization for models on large-scale ice bottom detection.
 - Integrated parameter tuning subroutines to the CReSIS codebase to automate model selections.
- Research Intern** May 2017 - Aug 2017
ZOLOZ
- Constructed parallel data processing pipelines for data collections.
 - Integrated ZOLOZ's software development kit into web and mobile technologies.

TEACHING EXPERIENCE

- Assistant instructor** Jan 2021 - Jun 2022
Washington University in St. Louis
- Held weekly recitations and office hours for mathematical optimization (ESE 415).
 - Prepared lecture notes and office hours for Bayesian methods in machine learning (CSE 515).

PUBLICATIONS

In preparation:

Preprints:

Published:

- JD Paden, V Berger, M Al-Ibadi, S Chu, M Xu, D Crandall, G Fox, “Subglacial bed topography using machine learning and geostatistical analysis applied to 2D and 3D radar sounding”, Advancing Earth and Space Science, Lawrence, Kansas, KS, USA 2018.

- V Berger, M Xu, S Chu, D Crandall, J Paden, GC Fox, “Automated Tracking of 2D and 3D Ice Radar Imagery Using Viterbi and TRW-S,” *IEEE International Geoscience and Remote Sensing Symposium*, Lawrence, Kansas, KS, USA 2018. doi: 10.1109/IGARSS.2018.8519411.

CONFERENCE PRESENTATIONS

- Poster, *Machine Learning in Computational and Systems Biology*, International Society for Computational Biology (ISMB) July 2022

SOFTWARE ENGINEERING SKILLS

- Languages: Python, Julia, C/C++ , Matlab, R, JavaScript, \LaTeX .
- High-performance computing: CUDA.jl, Slurm
- Pipeline Development: Snakemake
- Environments: Linux (Ubuntu)
- Mathematical Tools: FFTW
- Deep Learning: Flux.jl, PyTorch
- Visualizations: Makie.jl, Luxor.jl, D3.js, HTML

OPEN SOURCE CONTRIBUTIONS

Github: <https://github.com/kchu25>

Packages:

- **uCDL.jl** Motif discovery with deep unfolded convolutional dictionary learning.
- **SeqShuffle.jl** Shuffle strings such that it preserves the k -mer frequency in each string.
- **MotifPvalue.jl** Implemented an approximation algorithm that estimates the score thresholds of position weight matrices.
- **SimDNA.jl** Create DNA strings that simulate motifs with variable spacing, multimeric bindings, and multiple binding domains.

KEY COURSES

- Washington University in St. Louis: Large-scale optimization for data science, machine learning, Bayesian methods for machine learning, Bayesian inference, advanced algorithms.
- University of Kansas: Operating systems, parallel computing, numerical analysis, mathematical analysis, genetics, microbiology.

LANGUAGES

English and Mandarin.