Haotian Teng

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RESEARCH EXPERIENCE

Ph.D. Research Project, RNA methylation basecalling in third generation sequencing platform using Non-homogeneous HMM and Deep Learning.

2019-10 - present

Advisor: Prof. Carl Kingsford and Prof. Ziv Bar-Joseph, Machine Learning & Computational Biology department, SCS, CMU

- Designed a novel HMM structure to do the signal segmentation by constructing a time-dependent transition matrix from neural network output, which combine the interpretability of the HMM model and the strong expression power of the NN model
- Conducted data augmentation with random walk monto carlo simulation on a directed kmer graph constructed from the segmented signal.
- Developed an vector quantized variational autoencoder (VQ-VAE) based model to discovered subtle signal difference that corresponding RNA post-transcript modification, a graphical model is used as encoder for interpretability and a convolutional recurrent neural network (CRNN) is used as decoder for high classification accuracy.
- Achieved state-of-art accuracy in RNA methylation detection, and is the first kind of model established methylation-aware basecalling.

Ph.D. Research Project, combine deep learning and probabilistic graphical model to clustering the spatial transcriptomics data in single cell level. 2019-09-2021-02

Advisor: Prof. Ziv Bar-Joseph, Machine Learning & Computational Biology department, SCS, CMU

- Developed a dimensional reduction tool for gene expression profile (Program page: https://github.com/haotianteng/GECT).
- Developed a probabilistic Graphical model to cluster the cell from the spatial gene expression data (Program page: https://github.com/haotianteng/FICT).
- Produce a simulation pipeline for validating the spatial transcriptomics clustering tools and benchmark and visualization the clustering result. (Program page: https://github.com/haotianteng/FICT-SAMPLE)
- Conduct differential gene expression analysis and GO term annotation.

Masters Research Project, Using Machine learning algorithm in Nanopore Basecalling Advisor: Prof. Lachlan Coin, Institute for Molecular Bioscience, University of Queensland

2017-02 - 2018-07

- Built a deep learning-based basecaller **Chiron** using Tensorflow, for Oxford Nanopore sequencer basecalling (Program page: https://github.com/haotianteng/Chiron)
- Developed a preprocessing tool **Nanopre** to identify the polyA region in the Nanopore RNA sequencing platform. (Program page: https://github.com/haotianteng/Nanopre Tool)
- Prepared training dataset of DNA and RNA Nanopore basecalling reads, using Nanoraw and Graphmap to label the data.
- Implemented a pipeline in Google Cloud and Google Compute engine for end-to-end genome analysis.

Internship and Winter Scholar, The development of spontaneous neural activity in the zebrafish Advisor: Prof. Geoffrey Goodhill, Queensland Brain Institute, University of Queensland

- Built a pipeline for laboratory automation and data analysis in Zebrafish neuron experiment with Arduino, LabVIEW, and MATLAB.
- Constructed PHANTOM toolbox for projecting visual stimulation with conformal transformation, used for zebrafish tectum research. Program page in Github: https://github.com/haotianteng/PHANTOM-toolbox
- Developed algorithms for functional connectivity reconstruction using regularization method under scale-free assumption, correct the false positive correlation due to common input, transition connection, and latent common input.

Internship, Feedback in AIY neurons in Thermotaxis behavior of C.elegans

2015-07 - 2015-12

Advisor: Prof.Aravinthan D.T. Samuel, Center for Brain Science, Harvard University, Boston

- Studied thermotaxis in C.elegans with tracking and multi-neuron fluorescent marked.
- Cross & keep the worm, experiment using a spinning disk confocal microscope and the afterward data acquisition & processing with the combination of ImageJ (Miji) and Matlab
- Proved the derivation dependence between AFD neuron and temperature, designed and conducted the experiment to measure the parameters of the AFD-temperature relationship with temperature signal input under different shapes.

Research Assistant, Locomotion and PH sensoring mechanism in C.elegans & fast reaction tracking System development

Advisor: Dr. Louis Tao, Center for Bioinformatics, Peking University, Beijing

- Marked GCaMP6 into the C.elegans ASH, AWC, ASE neurons to testify and determine the neuron responsible for PH
- Developed a neuro-muscle model of C.elegans motor system and proved the theoretical prediction of gait adaptation in C.elegans.
- Recorded and analyzed long-term locomotion parameters of C.elegans by using a tracking and photographing system.
- Developed a visualization tool with openGL to describe and simplify the neuron network in C.elegans, and enabled the tool to search the whole neural pathway through any two given neurons.
- Built a tracking system as one of the contributors, which could achieve high-precision (accuracy below 1 micron) tracking and photographing and simultaneous data collection & processing
- Modified and developed a "snake" model based algorithm for robust and precise C.elegans center line extraction.

Professional EXPERIENCE

Algorithm Engineer Winter Intern

2019-01 - 2019-02

Alibaba, Hangzhou, China

Intelligence cache prediction using deep learning model based on biometric information.

Bioinformatics Engineer

Novogene Europe, Beijing, China

2018-09 - 2019-01

- Optimized the human resequencing and laboratory automation pipeline.
- Designed and developed the long-read sequencing platform.

Senior Research Technician

Institute for Molecular Bioscience, University of Queensland, Australia

2017-06 - 2018-07

Worked on Oxford Nanopore Technologies Long-read Nanopore direct RNA sequencing data processing, improve the sequencing accuracy and efficiency, improve the succeeded sequencing reads ratio by 15X compared to the original pipeline for long poly-A tail reads.

2014-07 - 2014-10

Biodynamic Optical Imaging Center, PKU, Beijing, China

- Micro-fluid chip preparation and fabrication.
- Developed a Computational Fluid Dynamics (CFD) module for the microfluid chips fluid field calculation in Fluent, which could draw the flow field from the CAD design sketch.

PUBLICATIONS

- Teng, H., Yuan, Y. and Bar-Joseph, Z., 2021. Clustering Spatial Transcriptomics Data. Bioinformatics.
- Pitt, M. E., Nguyen, S. H., Duarte, T. P., Teng, H., Blaskovich, M. A., Cooper, M. A., & Coin, L. J. (2020). Evaluating the genome and resistome of extensively drug-resistant Klebsiella pneumoniae using native DNA and RNA Nanopore sequencing. GigaScience, 9(2), giaa002.
- Teng, H., Cao, M. D., Hall, M. B., Duarte, T., Wang, S., & Coin, L. J. (2018). Chiron: translating nanopore raw signal directly into nucleotide sequence using deep learning. *GigaScience*, 7(5), *giy037*.
- Avitan, L., Pujic, Z., Mölter, J., Van De Poll, M., Sun, B., Teng, H., Amor, R., Scott, E.K. and Goodhill, G.J., 2017. Spontaneous activity in the zebrafish tectum reorganizes over development and is influenced by visual experience. Current Biology, 27(16), pp.2407-2419.
- Teng, H. "A neuron-muscle circuit model of C.elegans's locomotion." Bachelor of Science Thesis: Peking University, 2015

EDUCATION

Carnegie Mellon University, Pittsburgh, United States

Ph.D., Computational Biology

Advisor: Ziv Bar-Joseph, Machine Learning Department and Computational Biology department,

School of Computer Science, Carnegie Mellon University.

2020 - present

2016 - 2019

Featured courses: Probabilistic Graphical Models (10-708) A+, Deep Reinforcement Learning&Control (10-703), Convex Optimization (10-725) A+

University of Queensland, Queensland, Australia

M.S., Bioinformatics

- Advisor: Prof. Lichlan Coin, Institute of Molecular Bioscience, University of Queensland
- Advisor: Prof. Geoffrey Goodhill, Queensland Brain Institute, University of Queensland

Peking University, Beijing, China

B.S., Physics 2011 - 2015

HONORS AND AWARDS

•	The 1 st Prize at 27 th Chinese Physics Olympiad, Zhejiang Province (rank 1/1232 in theory part)	2011
•	The Silver Medal at 27 th Chinese Physics Olympiad, Finals	2011
•	The 1 st Prize at 29 th Parts of the National College Students Physics Competition	2012

SKILLS

- Programming: Python, C, C++, Matlab, R, Linux, LaTeX,
- Packages&Platforms: Tensorflow, MXNet, Caffe, CUDA, cuDNN, OpenGL, BWA, SAMtools, Velvet, DIAMOND, BLAST+, Minimap2, H5py, Psychtoolbox, LabVIEW, Arduino.
- Software: PyMOL, Fluent(ANSYS), Origin, AutoCAD, Primer Premier, DNA Man, Microsoft Office,
- Wet-lab experiment skill: Molecular cloning, Microinjection
- Language: Chinese(Mother Language), English(Fluent), Spanish(basic), German (Pizza-orderable) TOEFL: Cumulative 103 (R 29, L 29, S 23, W 22); GRE: V 150, Q 169, AW 3.0
- Proficient in Piano playing, learned since 6 years old. Skillful in saxophone.