

YI-YUAN LEE

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EDUCATION

M.S., Carnegie Mellon University, Pittsburgh, PA

May 2020

Master of Science in Computational Biology, School of Computer Science
Awarded Academic Achievement Fellowship

B.S., National Taiwan University, Taipei, Taiwan

June 2014

Bachelor of Science in Biochemical Science and Technology

Relevant Course List:

Introduction to Deep Learning, Introduction to Machine Learning, Biological Modeling and Simulation, Computational Methods for Proteogenomics and Metabolomics, Algorithm and Advanced Data Structure, Introduction to Computer Systems, Bioinformatics Data Integration Practicum, Immunology, Physiology, Statistics.

PUBLICATIONS

- **Yi-Yuan Lee**, Haodong Liu, Neel Mittal, Stephanie Eristoff, Liu Cao, Hosein Mohimani, "hypoNPAtlas: an atlas of hypothetical natural product for mass spectrometry database search". [06/15/2021 submitted to Nature Microbiology, <https://bit.ly/37lTlJj>]
- Liu Cao, Mustafa Guler, Azat Tagirdzhanov, **Yi-Yuan Lee**, Alexey Gurevich, Hosein Mohimani, "MolDiscovery: Learning Mass Spectrometry Fragmentation of Small Molecules". [06/04/2021 accepted, Nature Communications]
- Michelle et. al, including **Yi-Yuan Lee**, "A community resource for paired genomic and metabolomic data mining", Nature Chemical Biology, 2021.
- W.C. Su, S.-F. Hsu, **Y.-Y. Lee**, et al., "A Nucleolar Protein, Ribosomal RNA Processing 1 Homolog B (RRP1B), Enhances the Recruitment of Cellular mRNA in Influenza Virus Transcription", Journal of Virology, 2015.
- Tzu-Hui Hsu, Yu-Chan Chang, **Yi-Yuan Lee**, Chi-Long Chen, Michael Hsiao, Fan-Ru Lin, Li-Han Chen, Chun-Hung Lin, Takashi Angata, Fu-Tong Liu and Kuo-I Lin, "B4GALT1-dependent galectin-8 binding with TGF β receptor suppresses colorectal cancer metastasis". [in preparation]

RESEARCH EXPERIENCE

Computational Biology Department, Carnegie Mellon University

Sep 2018 - Now

Graduate Research Associate

Prof. Hosein Mohimani Ph.D.

- Developed deep neural networks for discovering novel ribosomally synthesized and post-translationally modified peptides (RiPPs), a class of natural products from microbial genomes. Both models are written in PyTorch and outperform the-state-of-the-art models in similar tasks.
- Implemented a subgraph-isomorphism-based chemical structure predictor, which generates hypothetical structures given a core peptide and a list of tailoring enzymes. Written in C++ and Rust.
- Mentoring three undergraduate students in research of *in silico* natural product discovery.

Genomics Research Center, Academia Sinica, Taiwan

Aug 2015 - May 2018

Research Assistant

Prof. Kuo-I Lin Ph.D.

- Project 1: "The role of B cell receptor (BCR) signaling in chronic lymphocytic leukemia (CLL)."
 - Established four drug-resistant clones for drug-resistance study.
 - Identified a kinase responses to BCR activity and stimulates the downstream transcriptional factor activation.
- Project 2: "The role of diSia motif in B cell immunity."
 - Identified the role of alpha-2,8-sialyltransferase 6 (ST8-6) in acute inflammatory response.
- Project 3: "The function of galectin-8 in colon cancer."
 - Performed extra splenic injection to confirm the anti-cancerous activity of galectin-8 with IVIS system.
 - Conducted animal experiments for lab members, and trained new lab members experimental techniques.

National Taiwan University, Taiwan

Jan 2013 - June 2014

Undergraduate Volunteer

Prof. Bi-Fong Lin, Ph.D., Bor-Luen Chiang, M.D./Ph.D.

- Confirmed the effect of additional nutrients on mesenchymal stem cells' (MSCs) immunoregulatory ability.

National Institute of Genetics (NIG), Japan

July 2012 - Sep 2012

Summer Intern

Prof. Inoue Ituro, M.D.

- Identified the mutation of *KDM5B* is one of the causalities of ovarian cancer.

Institute of Molecular Biology, Academia Sinica, Taiwan*Undergraduate Volunteer*

Jan 2012 - June 2012

Prof. Michael M.C. Lai, M.D./Ph.D.

- Supported the postdoctoral researcher to identify the role of human gene RRP1B in the infection and replication of *influenza A virus*. The result was published on Journal of Virology in 2015.

Institute of Molecular Biology, Academia Sinica, Taiwan*Summer Intern*

July 2011 - Sep 2011

Prof. Meng-Chao Yao, Ph.D.

- Verified the chromosome breakage executing time is later than dsRNA guided DNA deletion in *Tetrahymena thermophila* duplication.

PROJECTS

ChiGAN: A deep generative neural network to discover new chemical flavor compounds using a deep generative neural network with chemical structure data from FooDB.

SimCLL: A signaling and transcriptional factor network to simulate responses of healthy B cells, and Ibrutinib treated CLL cell with BioNetGen.

VireTap: A viral gene detection pipeline in human disease transcriptomes, by integrating bioinformatics tools such as TopHat, Trinity, and Blastn.

ProLang: A high level programming language parser and simulator to convert a code into cell-like structures. Written in Golang.

LEADERSHIP AND ACTIVITIES**Department of Life Science, National Taiwan University***Teaching Assistant in Biotechnological Core Techniques Course*

Feb 2014 - June 2014

- Coached students to complete molecular biology experiments.

Department of Information Management, Executive Government, Taiwan*Substitutive Military Service*

Aug 2014 - July 2015

Supervisor: Ming-Chung Lee

- Launched the guiding system and official website of History Exhibit in Executive Yuan.
- Solved problems of exigent program and system, and managed SVN subversion system.

iGEM NTU-Taiwan Team, Hong Kong*Team-leader*

Dec 2012 - Oct 2013

Advisor: Chern, Edward Yen-Rong

- Established a new iGEM team, recruited interdisciplinary members in NTU and raised funding.
- Awarded silver medal in the competition. Designed a biological heating device by transformed *SrUCP* into *Saccharomyces cerevisiae* and simulated the regulation circuit with different types of promoters by computational model. Enhanced about 150% heat production in our device. (Website: 2013.igem.org/team:NTU-Taiwan/index.html)

Novartis Biocamp, Taiwan*Participant*

May 2014 - May 2014

Organizer: Novartis Taiwan

- Only 30 students was selected out each year in Taiwan from various backgrounds.
- Acquired knowledge about how pharmaceutical industry works and worked with group members to create a business plan of hypertension drug.

SKILLS

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| Programming Language | Python, C, C++, Rust, Go, Bash, R |
| Bioinformatics | HMMER, BLAST, RDKit, Dereplicator+, MolDiscovery, BWA, SAMtools, NGS |
| Deep Learning | PyTorch, TensorFlow, Numpy, Scipy, Sklearn, modAL |
| Development Tools | Linux, AWS, Git, Slurm, L ^A T _E X, VIM, tmux |
| Wet lab skills | mouse breeding and handling, simple survival surgery in mice and blood collection, primary B cell isolation, cell line and primary cell culture, flow cytometry, plasmid construction, point mutation, protein purification, genotyping, PCR, qRT-PCR, transfection, western blotting, virus preparation |
| Language | English (fluent), Mandarin Chinese (native speaker), Japanese (conversational) |

TEST SCORE**TOEFL:** 111/120 (Reading: 29, Listening: 30, Speaking: 27, Writing: 25)

Sep 2019

GRE: Verbal 152/170, Quantitative 167/170, Analytic Writing Analysis: 3.5/6.0

Nov 2017