

# Hua Cailong

☎ 651 6756613 @ hua00023@umn.edu

## Education

### UNIVERSITY OF MINNESOTA - TWIN CITIES

PHD STUDENT IN ELECTRICAL AND COMPUTER ENGINEERING

📅 2019 - Present 📍 Minnesota, US  
GPA 3.947/4.0

### IMPERIAL COLLEGE LONDON

MSc. IN CONTROL SYSTEMS

📅 2016-2017 📍 London, UK  
Graduated with Distinction  
GPA: 75.4/100

### POLITECNICO DI MILANO

BSc. IN AUTOMATION

ENGINEERING

📅 2015-2016 📍 Milano, Italy  
GPA: 110/110

### TONGJI UNIVERSITY

BSc. IN ELECTRONIC INFORMATION AND ENGINEERING

📅 2012-2016 📍 Shanghai, China  
GPA: 89.52/100 (4.45/5.0)

## Links

🔗 huacailong.github.io

🐙 github.com/huacailong

in www.linkedin.com/in/cailong-hua-a9aa5719a

## Main Courses

Machine Learning • Artificial Intelligence • Optimal Control and Reinforcement Learning • Intelligent Agents • Deep Learning

## Skills

Python • Matlab • Simulink • C++ •  $\text{\LaTeX}$  • AFM • Optical Tweezer

## Languages

English • Chinese • Italian

## Honors

2019 ECE Fellowship  
2016 Outstanding Graduate  
2014 & 2015 & 2016 Third-Class Scholarship  
2015 China Scholarship Council Scholarship  
2013 China National Scholarship  
2013 First-Class Scholarship

## Research Experience

### NON-EQUILIBRIUM EXPERIMENTS STUDY

UNIVERSITY OF MINNESOTA

📅 2020 - Present

📍 Minnesota, US

- Developed algorithm for quantifying errors in non-equilibrium experiments
- Implemented the algorithm with a Python-based toolbox
- Validated the algorithm through a simulated spring-mass system under non-equilibrium conditions
- Established a proof by conducting experiments with Optical Tweezers and aligning the outcomes with a corresponding simulation

### MODELING OF MUSCLE PROTEINS

UNIVERSITY OF MINNESOTA

📅 2021 - Present

📍 Minnesota, US

- Carried out cell transfection, protein purification, and gel assays for muscle proteins utrophin and dystrophin
- Designed force spectroscopy experiments characterizing muscle proteins using Atomic Force Microscopy
- Investigated the impact of different expression systems on protein behavior through experiments
- Automated analysis of experimental data with Matlab
- Developed Monte Carlo simulation to replicate experimental processes
- Developed muscle protein modeling by conducting a statistical analysis of experimental data.

## Projects

### RECENT PROJECTS

📅 2021 - Present

📍 Minnesota, US

- Uncovered causal relations from protein pulling data and quantified the change of effects among protein properties
- Employed Langevin-based sampling techniques in conjunction with energy-based models to address the challenging partition function issue
- Employed a model based on Generative Adversarial Networks (GANs) to synthesize a realistic image of the person with the target pose based on a source image showing the person with a given pose
- Explored the effect of different fine-tuning layers and pooling strategies on the performance of the BERT model, a transformer based machine learning technique for NLP
- Designed an unsupervised agent to play the Flappy Bird game based on vector data instead of images with Reinforcement Learning technique

## Publications

Ramirez, M. P., Rajaganapathy, S., Hagerty, A. R., **Hua, C.**, Baxter, G. C., Vavra, J., ... & Ervasti, J. M. (2023). "Phosphorylation alters the mechanical stiffness of a model fragment of the dystrophin homologue utrophin." *Journal of Biological Chemistry*, 299(2).

Rajaganapathy, S., **Hua, C.** and Salapaka, M., "Confidence bounds for the Jarzynski estimator." In *APS March Meeting Abstracts (Vol. 2022, pp. S09-007)*, 2022.