hj2368@columbia.edu • 646-643-1348 • helenjin.me

Helen Jin

EDUCATION

Columbia College, Columbia University, New York, NY

B.A. Candidate in Mathematics, Computer Science

- Relevant Coursework: Data Structures in Java, Programming Lang (Python), Discrete Math, CS Theory, Linear Algebra, Modern Algebra I, Calculus-Based Intro to Statistics, Probability Theory
- Honors/Awards: Dean's List (all semesters)
- **Teaching Assistant** for Calculus I (Fall 2017), Calculus IV (Spring 2018) | **Grader** for Calculus-Based Intro to Statistics (Fall 2017)

Stuyvesant High School, New York, NY

Sep 2012 – Jun 2016

Expected: May 2020

GPA: 3.8 / 4.0

GPA: 4.0 / 4.0

Advanced Regents Diploma

- Activities: Girls' Varsity Swimming, Lifeguard, Stuyvesant Spectator (News and Art), Math Team, CS Dojo and Writing Center Tutor, Red Cross Volunteer
- Honors/Awards: AP National Scholar, National Merit Finalist, ARISTA Honor Society, National Latin Society

SELECTED EXPERIENCE

Grantham, Mayo, van Otterloo & Co. LLC (GMO), Boston, MA

Expected: Jul - Aug 2018

Incoming Quantitative Equity Analyst Intern

· Global Equities team, will work with MATLAB and SQL

Computational and Systems Biology, MSKCC, New York, NY

May 2017 – Present

Undergraduate Researcher

• Worked with Dr. Dana Pe'er to develop computational methods/packages to analyze single cell data using Python and R

SELECTED LEADERSHIP

Girls Who Code (GWC) at Columbia University, New York, NY

Feb 2017 - Present

Vice President of External Affairs, Executive Board

- Oversaw high school outreach and recruitment for program that holds weekly CS classes for 50 high school girls each semester; worked with other Executive board members to better improve program
- Managed ~30 people on the Managing Board (Finance, Programming, Publicity, HS Recruitment)
- Planned on-site visits, created fundraisers, and reached out to companies and local businesses for sponsorship

SELECTED PROJECTS

SCAnalysis

Built a Python package and Jupyter notebook to analyze single cell RNA-seq data that incorporates MAGIC and Wishbone
technologies, with additional features (including Palantir). Wishbone is an algorithm to align single cells from
differentiation systems with bifurcating branches. MAGIC (Markov-Affinity Based Graph Imputation of Cells) is an
interactive tool to impute missing values in single-cell data and restore the structure of the data.

Tetris Remix

• A modified re-make of the classic game Tetris in Processing; has Levels mode (twenty successive levels) and Infinite mode

SKILLS & INTERESTS

Technologies: Java, Python, R, Git, Jupyter Notebook, LaTeX, HTML/CSS/JavaScript, Processing, Racket, NetLogo, Microsoft Office (Excel, Word, PowerPoint), G Suite (Gmail, Drive, etc.)

Languages: English (native), Mandarin Chinese (intermediate), Korean (intermediate), Spanish (basic), Japanese (elementary)

Personal Interests: Philosophy, Music, Visual Arts, Swimming, Yoga