

# Maximiliano Eaton

MaxEaton.github.io | maximiliano.k.eaton@gmail.com | linkedin.com/in/maximiliano-eaton | github.com/MaxEaton

## Education

**University of Colorado, Boulder** – BS in Computer Science and Applied Mathematics Expected May 2026

- Major GPA: 4.000; Cumulative GPA: 3.988
- Computer Engineering Minor
- Engineering Honors Program
- Relevant Coursework: Algorithms, Theory of Computation, Linear Programming

## Skills

**Technical:** C/C++, Python, Rust, Java, Git, GDB, numpy, TensorFlow, SystemVerilog

**Languages:** English, Japanese

## Experience

**Course Assistant**, University of Colorado, Boulder – Boulder, CO June 2024 – Present

- Assisted both Computer Systems (CSCI 2400) and Operating Systems (CSCI 3753) over the summer
- Conducted office hours to provide additional support and clarification on course material
- Collaborated with course teams to organize lab help and midterms reviews to support students

**Undergraduate Researcher**, University of Colorado, Boulder – Boulder, CO May 2023 – October 2023

- Optimized computer simulation for archaic genomics to trace lineage of neanderthals
- Parallelized tasks on CU's Research Computing facilities for better performance of machine learning models
- Contributed on poster for Society for Molecular Biology and Evolution conference (2023)

**Intern**, Dataquest – Boulder, CO July 2021 – October 2022

- Implemented checkers to determine correct user submissions for the online platform to learn data science skills
- Resolved bugs submitted by users and beta tested new content whenever deployed

## Projects

**Primomata** github.com/MaxEaton/Primomata

- Created tool that converts between different forms of regular languages, namely  $\epsilon$ -NFA, DFA, and regex
- Implemented different algorithms to manipulate DFAs to find properties of prime numbers
- Visualized automatas using dotLang for both NFAs and DFAs to see relationships
- Inspired me to further pursue subject by taking a formal Theory of Computation course

**Genome Tracing** github.com/MaxEaton/GenomeGraph

- Derived minimal gene mutation paths to determine possible lineage of genetic strain
- Pruned branches and utilized bit masking to increase efficiency

**Probabilistic Sudoku** github.com/MaxEaton/sudoku

- Generated random sudoku puzzles and ensured feasibility by backtracking to determine existence of unique solution
- Attempted to design a statistical solver to more efficiently solve NP-complete problem

**Grid Combinatorics** github.com/MaxEaton/Grid

- Computed the number of unique patterns of dots in a grid under rotational and translational symmetry
- Utilized bit manipulations and specialized data structures for optimization