

### Matheus Cardoso

Computer Scientist (B. Sc.)

### **Contact**

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# Skills (0 to 5 yrs.)

# Programming Languages

Python	3+ yrs.
Shell Scripting	3+ yrs.
R	1.5+ yrs.
C/C++	1+ yrs.
SQL	6+ mos.
Java	6+ mos.
Rust	6+ mos.
Javascript	6+ mos.
RISC-V Assembly	6+ mos.
Technologies	
HTML/CSS	2.5+ yrs.
Git	2.5+ yrs.
PTEX	1.5+ yrs.

# **Biography**

Computer science student, work with data analysis in the area of genomics and proteomics, developing software that automatically generates statistics on enriched monoclonal antibodies. His work mainly involves machine learning techniques, data mining, data analysis and data visualization. As a result, in addition to working in the area of computer sciences, he also have a strong focus on biological and data sciences.

In his free time he have been contributing to open source projects as well; Spanning from data science frameworks (Pandas), to educational learning software (Anki), and productivity programs (ncspot, unclutter and hacksaw).

### **Education**

#### Computer Science | B.Sc. student

03/2021 - today

University of Brasília - UnB

I aced all my exams and projects until now.

GPA: 5/5

#### Computer Science | B.Sc. student

01/2020 - 03/2021

Federal University of Rio de Janeiro - UFRJ

Due to the pandemic I opted to move from Rio back to Brasília.

GPA: 2.9/5

### Biotechnology | B.Sc. student

01/2019 - 12/2019

University of Brasília - UnB

I had the opportunity to internship in UnB's Bioinformatics and Immunology laboratory, where I contributed to the development of a software to assess enrichment of monoclonal antibodies. There I found that more than research I really enjoy to code, so I opted to change my majors to Computer Science.

GPA: 4.8/5

# **Projects**

### ATTILA - AutomaTed Tool For Immunoglobulin

01/2019 - today

University of Brasília - UnB

ATTILA is an open source project that analyses phage display libraries in order to assess monoclonal antibody enrichment. Doing this, the software is able to predict which antibodies can more effectively bind to target molecules. As a result, ATTILA becomes a useful tool to scientists working on vaccine development.

### cdr3-parser

06/2021 - today

University of Brasília - UnB

cdr3-parser is a command line program written in Rust to convert intermediate output files generated from ATTILA into csv files containing statistics about VH-CDR3 regions presented in the input file. cdr3-parser leverages Rust's thread-safe capabilities to parse the file in the most efficiente way possible using all threads available in the UnB's Bioinformatics laboratory server.

# **Frameworks Biopython** 2.5+ yrs. **Tidyverse** 1.5+ yrs. **Tidymodels** 1.5+ yrs. Numpy 6+ mos. 6+ mos. **Pandas** 6- mos. Django Areas of expertise System administration/ 3+ yrs. Linux **Bioinformatics** 2.5+ yrs. Data science 2+ yrs. Low level programming/ 6+ mos. **Assembly** Languages **Portuguese** Native B2 **English** В1 Spanish

### Interests

- Bioinformatics/ Computational Biology
- Artificial Intelligence
- Protein Folding
- Open Source
- Embedded Systems

#### The Adventures of Lolo

University of Brasília - UnB

03/2021 - 05/2021

Offiversity of Brasilia - Offib

Remake of the game  $\it The \ Adventures \ of \ Lolo \ in \ RISC-V \ Assembly.$ 

#### Mastermind

02/2021 - 03/2021

Federal University of Rio de Janeiro - UFRJ

Remake of the game Mastermind in C.

#### Brazilian Students' Performance on ENEM

08/2020

Johns Hopkins Bloomberg School of Public Health

A data visualization project showing the performance of Brazilian students applying to a bachelors of science at Universidade de São Paulo (USP), one of the best universities in Brazil.

#### Word Predictor Model

08/2020

Johns Hopkins Bloomberg School of Public Health

A web application powered by a machine learning model able to predict the next word in a sentence.

### Awards & Certificates

### **Data Science Specialization**

01/2020 - 09/2020

Johns Hopkins Bloomberg School of Public Health

### Copenhagen Bioinformatics Hackathon

2020

BioLib

Winner of the Hackathon challenge "Variant Pathogenicity Prediction".

## Capes Prize University Talent

2019

CAPES

Awarded by the Brazilian Ministry of Education as one of the top 1000 most talented freshman students in the country.

#### Hackathon NASA Space Apps BSB

2019

NASA

Awarded as 5th out of 60+ teams in the Brazil's Midwest regional phase.

São Paulo, 17th July 2021

Matheus Cardoso de Souza