# Jaldert François

PhD Candidate in Bioscience Engineering (Bioinformatics)

I am a **Bioinformatician (M.Sc.)** working on **Protein Engineering, Applied Machine Learning** and Statistical Analysis at STADIUS (Engineering) and CSB (Bioscience Engineering).



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# **SKILLS**

Machine Learning

Bioinformatics

Statistical Analysis

Pyhon, R, Bash, SQL

Databases, Data Mining

# **LANGUAGES**

#### English

Native or Bilingual Proficiency

#### Dutch

Native or Bilingual Proficiency

#### **AWARDS**

Honours College Program - University of Antwerp (09/2017 -06/2019)

FWO- SB Research Grant (11/2022 - 11/2026)

## **EDUCATION**

# **Bioinformatics (Master)**

### **KU Leuven**

09/2019 - 06/2021

Cum Laude

Courses

Electives: (1) Neural Computing (2)
 Univariate Data Analysis and Modelling (3)
 Gene Technology

 Thesis: "Exploring mutational patterns of evolution experiments: Meta-analyses of microbial data from the CAMEL database"

# **Biochemistry and Biotechnology (Bachelor)** University of Antwerp

09/2016 - 06/2019

Summa Cum Laude

Additional Course Work

- Honours College Program

# **WORK EXPERIENCE**

## **PhD Candidate**

FWO - KU Leuven

10/2021 - 11/2026 Leuven

Skills & Research Domains

- Protein/Enzyme Engineering, Proteomics, Computational Protein annotation
- Applied Machine Learning for predictive modelling
- Applied Deep Learning (ANN, CNN, GNN), neural network optimisation & LLM fine-tuning
- Data mining and database development (SQL, NoSQL and Graph databases)
- Statistical Analysis & Software Engineering
- Teaching Bioinformatics (3rd Bachelor Bioscience Engineering) & Introduction to Object
   Oriented Programming (Master of Bioinformatics & Master of Artificial Intelligence)

# Research Internship University of Antwerp

08/2018 - 09/2018

Antwerp

Achievements/Tasks

- "Examining novel mutations in Charcot-Marie-Tooth"
- DNA/RNA extraction & analysis

## **PROJECTS**

Enzymares - https://catalisti.be/project/enzymares/ (10/2021 - 10/2024)

- Development of a toolbox for enzyme selection, including a custom database linking UniProt, Brenda, Rhea. etc.
- Implementation and development of machine learning models for enzyme (parameter) prediction, and efficient enzyme screening

FWO - SB (10/2022 - 10/2026)

- □ Title: "Optimising the enzyme discovery pathway through omics integration and AI"
- Exploring current approaches for enzyme classification, reaction parameter prediction and substrate affinity
- Developing novel machine learning models for enzyme classification and reaction parameter prediction
- Development of an enzyme selection toolbox/software