

### Jérémie KALFON

Phone +33 652663443

Email jkobject@gmail.com

Website jkobject.com

LinkedIn www.linkedin.com/in/jkobject

Github www.github.com/jkobject

Twitter @jkobject

Address 76 avenue de suffren, 75015

Paris, France

### **Profile**

Motivated graduate student in Biomed Eng. & ML. Intern in Research & R&D companies.

I wish to do a PhD. CEO at PiPle & leader of school associations, I am keen on project development and highly independent. I am interested in improving Genomics & Biomedical research thanks to data-science and machine learning.

## Technical skills

Data Science

Imaging

Neuroscience

CS

Maths & stats
genomics theory

Electronics

English

Team work

# **Experience**

### **Projects**

 Developing comparison tools and notebooks for the Calcium Imaging Analysis software (CalmAn)

RESEARCH, IMAGING, ML, VISION

201

 CEO of PiPle, a startup focused on bringing the next generation messaging application

PROJECT DEV. ENTREPRENEUR

2016

- Developing PyCUB, Master Thesis in ML for the Codon Usage Bias
   ML, BIG DATA, RESEARCH, GENOMICS 2017 -
- Developing VCF2Ancestry, Hackathon for the BroadInstitute

ML, BIG DATA, GENOMICS

201

### Exchanges and Internships

 Internship at the Simons Foundation's CCB, a research facility in NYC under the supervision of Dmitri Chklovskii

An exchange semester in Mexico at TEC de Monterrey
 CRARLEY

 Intern at HighLife Med., a Parisian valve implant R&D start-up SIMULATION, DATA ANALYTICS
 2019

#### Additional & Societies

- Leader of the NoLarsen music Festival gathering 1500 people in Paris

  Music Society
- Teacher of high school & undergraduate students weekly for 2 years
   Teaching Society

### **Education**

French Diplôme d'ingénieur, Biomed. Eng., ECE paris
PARIS, FRANCE
2013-2018

M.Sc. in Computational Intelligence, University of Kent

CANTERBURY. ENGLAND

2017-2018

### Interests













- CalmAn: An open source tool for scalable Calcium Imaging data Analysis, Andrea Giovanucci. eLife. DOI:10.7554/eLife.38173
- PyCUB: A machine exploration of the Codon Usage Bias, Master Thesis. (superv. of Tobias Von der Harr & Dominique Chu) www.jkobject.com/projects/PyCUB.pdf
- Hidden Patterns of the Codon Usage Bias, Yun Deng, Jeremie Kalfon, Tobias Von de Harr, Dominiaue Chu, in review at Nucleic Acid Research, DOI:10.1101/478016