BIOINFORMATICS PHD STUDE

Paris FR
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## **Experience**

Institut Pasteur Paris France

PHD STUDENT Jan 2020 - Now

- · Using machine learning for biological sequence analysis.
- Funded by the Paris Al Research Institute (PRAIRIE)
- Goal: using deep learning for more accurate contest-aware protein sequence alignment
- More: research.pasteur.fr/en/project/applying-machine-learning-to-sequence-analysis-phd-luc-blassel-prairie/

Institut Pasteur Paris France

RESEARCH ENGINEER Apr 2018 - Dec 2019

- · Using machine learning and big data to explore the drug resistance landscape in HIV.
- Using ML methods to find new Drug resistance mutations
- · Presentation of the project at MCEB 2019 conference
- Usage of Snakemake pipelines in a SLURM HPC environment
- More: research.pasteur.fr/en/project/drm-hiv/

 CapitalData
 Paris France

Data scientist Intern

Mar 2017 - Aug 2017

- · Development on automated data validation and anomaly detection pipelines using R, Golang and Python.
- Usage of R caret and Python scikit-learn libraries
- Internal client-oriented reporting with MySQL, BIME, Periscope and Mode
- More: capitaldata.fr/

## **Education**

Paris Dauphine University

Paris France

MASTER IN SCIENCE: MACHINE LEARNING Sep 2017 - Aug 2018

- · Reinforcement learning
- · Data Mining
- Deep learning
- Double diploma with AgroParisTech
- score: 15.58/20 More: dauphine.psl.eu/formations/masters/informatique/m2-intelligence-artificielle-systemes-donnees

AgroParisTech Paris France

DIPLÔME D'INGÉNIEUR: BIOLOGY - MACHINE LEARNING

Sep 2014 - Aug 2018

- · Microbiology
- Immunology
- Projects in Java, C and Matlab
- Advanced statistics and linear modeling
- · Machine learning
- score: 15.58/20 More: agroparistech.fr/ufr-info/iodaa/

École Saint Hilaire Paris France

NATIONAL COMPETITIVE EXAM: BCPST PREPARATORY CLASS

Sep 2011 - May 2014

- ranked 201/1967 (Veto)
- ranked 481/2991 (Agro)
- More: en.wikipedia.org/wiki/Classe\_preparatoire\_aux\_grandes\_ecoles

## **Publications**

- Frédéric Lemoine, Luc Blassel, Jakub Voznica and Gascuel Olivier, "COVID-Align: Accurate online alignment of hCoV-19 genomes using a profile HMM", *Bioinformatics* (2020), doi.org/10.1093/bioinformatics/btaa871
- Anna Zhukova, Luc Blassel, Frédéric Lemoine, Marie Morel, Jakub Voznica and Olivier Gascuel, "Origin, evolution and global spread of SARS-CoV-2", Comptes Rendus. Biologies (Académie des sciences) (2020), doi.org/10.5802/crbiol.29

## **Languages**

English **Native**French **Native** 

German **B1** 1 mo. program at Goëthe Institute in Germany

Chinese  $\$  Intermediate 5 and 1 mo. programs in Shanghai SISU and Beijing UIR