

## Skills

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**Programming:** Python3, C++, C#, Bash, C, HTML, CSS

**Technologies:** Git, CI/CD (GitHub Actions), Docker, Unity, Django, CMake, SQL, MongoDB

**Languages:** French (native), English (proficient), Mandarin (beginner)

## Experience

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### Research Assistant & Research engineer

2019 – 2024

Laboratory of Theoretical Biochemistry, IBPC · CNRS , Paris

- Developed a Python algorithm to estimate interaction parameters between a reduced-resolution particle system and an analytical field, allowing large-scale dynamic simulations [1]
- Standardized multi-threaded workflows (C++, C#) for real-time molecular simulation control and monitoring, ensuring seamless and extensible data exchange
- Enhanced a 3D visualization interface (Unity, C#) for real-time Human-computer interaction and visualization of the complex dynamics of molecular shapes and their spatial evolution [1, 2]
- Implemented a CI/CD pipeline (GitHub Actions, Docker), automating software releases, resolving dependency issues, and improving accessibility for end-users
- Led the writing of the first review article on interactive molecular simulations (IMS) [3], synthesizing decades of community research

### Bioinformatics Drug Design Intern

Feb 2019 – June 2019

Bioinformatics and biophysics team, IMPMC · Sorbonne université , Paris

- Developed a workflow combining molecular docking and sampling simulations using a national supercomputer, with results compared to alternative methods
- Designed cyclic peptides using molecular visualization and modeling tools

### Bioinformatics Structural Analysis Intern

July 2018

Bioinformatics Research Team, ISYEB · Sorbonne université , Paris

- Optimized a protein similarity search by re-implementing Bellman's algorithm (Python) to find the longest path through shared motifs, and developing heuristic algorithms for faster matching

## Education

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**Université Paris Cité** – Ph.D. in Bioinformatics

2023

Advisor: Dr Marc Baaden

**Sorbonne Université, Paris** – B.S. in Biology & M.S. in Bioinformatics

2019

## Publications

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- [1] André Lanrezac and Marc Baaden. "UNILIPID, a Methodology for Energetically Accurate Prediction of Protein Insertion into Implicit Membranes of Arbitrary Shape". *Membranes* (Mar. 2023). DOI: 10.3390/membranes13030362.
- [2] André Lanrezac et al. "Fast and Interactive Positioning of Proteins within Membranes". *Algorithms* (Nov. 2022). DOI: 10.3390/a15110415.
- [3] André Lanrezac, Nicolas Férey, and Marc Baaden. "Wielding the power of interactive molecular simulations". *WIREs Computational Molecular Science* (July 2022). DOI: 10.1002/wcms.1594.
- [4] André Lanrezac, Nicolas Férey, and Marc Baaden. "Interactive Molecular Dynamics". *Reference Module in Chemistry, Molecular Sciences and Chemical Engineering*. Elsevier, 2023. DOI: 10.1016/B978-0-12-821978-2.00115-X.