Kilian Fatras

Senior Machine Learning Research Scientist

Research Interests: Generative Models, Protein Design, Optimal Transport, Distribution Shifts

Experiences

- 04/25-now Machine Learning Research Scientist EvolutionaryScale New York City, USA
 - Conducting foundational model research in protein sequence and structure using large-scale protein language models and generative diffusion models.
 - Developing scalable architectures for protein-related tasks such as folding and binder design.
 - Collaborating with wet-lab scientists to validate designed proteins.

06/23–04/25 Founding Machine Learning Research Scientist - Dreamfold - Montréal, Canada

- Built and trained generative models for protein design with focus on binder design, motif scaffolding, unconditional generation, and hotspot interactions.
- Co-led design of multimodal diffusion models conditioned on structure and sequence.
- Contributed to model interpretability, evaluation metrics (pLDDT, pTM), and training infrastructure.

01/22–12/23 **Postdoctoral Fellow on Distribution Shifts and Generative Modelling -** Mila & McGill University Supervisors: Professor Ioannis Mitliagkas and Professor Adam Oberman

- Published 11 papers in top ML conferences on flow matching models and distribution shifts.
- $\circ\,$ Developed and contributed to open-source software that reached thousands of stars on GitHub.

Education

- 2021 PhD in Computer Science: "Optimal Transport & Deep Learning: Learning from One Another"– INRIA –Supervisors: Professor Nicolas Courty and Professor Rémi Flamary
- 2018 Master of Science in Machine Learning and Entrepreneurship UC Berkeley & Polytechnique
- 2017 Master of Engineering in Applied Mathematics and Computer Science ENSTA Paris

Selected Publications and Open-Source Software

- 1. Generalizing Flow-Based Generative Models with Optimal Transport [URL] Alex Tong^{*}, Kilian Fatras^{*}, Nikolay Malkin^{*} et al. TMLR 2024
- SE(3)-Stochastic Flow Matching for Protein Backbone Generation [URL] -J. Bose, T. Akhound-Sadegh, G. Huguet, Kilian Fatras et al. - ICLR 2024
- 3. Sequence-Augmented SE(3)-Flow Matching for Conditional Protein Backbone Generation - [URL] - Guillaume Huguet*, James Vuckovic*, Kilian Fatras* et al. - NeurIPS 2024
- 4. Unbalanced Minibatch Optimal Transport; Applications to Domain Adaptation [URL] Kilian Fatras, Thibault Séjourné, Nicolas Courty, and Rémi Flamary ICML 2021
- 5. Learning with Minibatch Wasserstein: Asymptotic and Gradient Properties [URL] Kilian Fatras, Younes Zine, Rémi Flamary, Rémi Gribonval, and Nicolas Courty AISTATS 2020
- 6. Optimal Transport and Deep Learning: Learning from One Another [URL] Thesis
- 7. Conditional Flow Matching https://github.com/atong01/conditional-flow-matching
- 8. POT: Python Optimal Transport library https://github.com/PythonOT/POT

Selected invited talks

- 03/10/24 Xaira Therapeutics: Sequence-Augmented SE(3)-Flow Matching for Protein Backbone Generation
- 17/09/24 EvolutionaryScale: Sequence-Augmented SE(3)-Flow Matching for Protein Backbone Generation
- 11/04/23 FAIR Lab (Montréal): Designing and evaluating new domain adaptation methods
- 28/02/23 Microsoft AI Lab (Montréal) seminar: Optimal transport and deep partial domain adaptation