



Jonathan Harrison

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<https://github.com/shug3502>

Scientific expertise

- Machine Learning
- Molecular Property Prediction
- Drug Discovery
- Bayesian Optimization/Active Learning
- Bayesian Statistics

Education

2019. DPhil in Systems Biology
University of Oxford
Developing and applying modelling and Bayesian inference tools for cell biology
Supervised by Prof. R.E. Baker

2014. MMath in Mathematics
University of Oxford
1st class

Languages

Python ●●● R ●●●
Julia ●●○ Stan ●●○
C++ ●○○ Git ●●○

Current position

November 2024 – Present **Senior AI Research Scientist, Recursion**

July 2022 – October 2024 **AI Research Scientist, Exscientia**

The focus of my work is on molecular property prediction, including uncertainty quantification for machine learning models and with applications in Bayesian Optimization/Active Learning. I have experience leading a product team in this area.

I have developed strong software skills, taking projects from ideation to deployment and practical real world use. I have trained and fine-tuned foundation models for molecular property prediction. I am a core contributor to **Molflux** - a high profile open source ecosystem for machine learning for chemistry. Machine learning models and active learning software I have developed are used day-to-day by dozens of drug designers in their standard workflow.

Previous experience

Jan. 2019 – July 2022 **Postdoctoral Research Fellow, University of Warwick** (With Prof. N.J. Burroughs; Prof. A.D. McAinsh)

Using computational tools from Bayesian statistics, differential equations, stochastic modelling and image analysis, I provided insight into cell biology. I modelled chromosome dynamics allowing understanding of how and why cell division goes wrong. Conceptualized and supervised MSc rotation projects for 2 PhD students.

Jan. - Apr. 2018 **Data Science Intern at Hudl**

Worked within research and development team on a project investigating event detection in football matches using deep learning.

Selected Achievements, Presentations and Publications

Achieved 2nd place in **public molecular property prediction competition run by Polaris (2025)**

Recursion **annual 10-K report** included my analysis showing "a >2.5x increased efficiency in detecting new bioactive scaffolds with a >40% reduction in the flagging of likely cytotoxic compounds." (2025)

Invited talk at **Active Learning in Pharma Symposium (2024)**

O. Sen*, J.U. Harrison*, N.J. Burroughs, and A.D. McAinsh (2021) *Kinetochores life histories reveal the origins of chromosome mis-segregation and correction mechanisms* Developmental Cell <https://doi.org/10.1016/j.devcel.2021.10.007> (All computational work - JUH, experiments - OS)

J.U. Harrison and R.E. Baker (2020) *An automatic adaptive method to combine summary statistics in approximate Bayesian computation* PLOS One <https://doi.org/10.1371/journal.pone.0236954>

For full list of publications see [my Google Scholar profile](#)