# Hrushikesh Loya

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### **Professional Summary**

I am a Postdoctoral Researcher in the Department of Statistics at the University of Oxford, working at the intersection of statistical genetics, machine learning, and computational biology. I have extensive experience in developing methods for genome-wide association studies, polygenic score analyses, and fine-mapping to investigate the genetic basis of human phenotypes. I am proficient in Python and R, as well as statistical and machine learning methods, and I have a strong track record of publications and mentorship in genetics and statistics.

### Education

University of Oxford

Oxford, UK

D.Phil. in Genomic Medicine and Statistics

Oct 2020-Jan 2025

Supervisors: Prof. Simon Myers and Prof. Pier Palamara

Thesis: Powerful new methods for decomposing genome-wide ancestry and performing trait association

**Indian Institute of Technology Bombay** 

Mumbai, India

B. Tech. + M. Tech. in Electrical Engineering

Jul 2015-Jul 2020

Class rank: 3 out of 82, GPA: 9.54/10, Specialization in Communication & Signal Processing

## **Work Experience**

#### **Department of Statistics**

Oxford, UK

Postdoctoral Researcher

Jan 2025-Present

- Designed a genealogy-based framework that revealed hidden evolutionary events in ancient human populations
- Developed a scalable inference method that integrates deep learning-based sequence-to-function models, enabling faster and more accurate whole-genome fine-mapping of causal variants

Ossian Biosciences Inc.

Delaware, USA

Research Consultant (Part-Time)

Feb 2025-Present

- Advised on the scientific strategy and technical roadmap, contributing to the raise of \$1M in pre-seed investments
- Created an LLM-driven workflow that combined literature insights with multi-omics and GWAS data to generate therapeutic hypotheses for age-related diseases, focusing on sarcopenia

Samsung AI Center

Cambridge, UK

Research Intern

Aug 2022-Feb 2023

- Developed a novel meta-learning framework that leveraged task-level correlations across diverse search spaces to enhance neural architecture search with the aim to improve performance in under-represented domains
- Secured a first-authored UK Patent for innovative contributions to neural architecture search methodologies

### **Publications**

### **Computational Biology and Statistical Genetics**

[1] H. Loya, P.F. Palamara, L. Speidel, S.R. Myers, Genealogy-based Detection of *Target journal: Nature*Ancient Ghost Admixture Across Africa (in preparation)

[2] H. Loya, G. Kalantzis, F. Cooper, P.F. Palamara, A Scalable Variational Inference
Approach for Increased Mixed-model Association Power

Nature Genetics, 2025

[3] J. Zhu, G. Kalantzis, A. Pazokitoroudi, H. Loya, H. Chen, S. Sankararaman, P.F. Palamara, Fast Variance Component Analysis using Large-Scale Ancestral Recombination Graphs

Cell Genomics, 2025

[4]	N. Bird, H. Loya, L. Speidel, S.R. Myers, G. Hellenthal, The Reliability of Inferred Archaic Segments in Human Genomes	<i>EJHG</i> , 2025
[5]	U. Bezaljak, H. Loya, B. Kaczmarek, M. Loose, T. Saunders, Stochastic Activation and Bistability in a Rab GTPase Network	PNAS, 2020
Mac	hine Learning and Methodology	
[6]	H. Loya, Ł. Dudziak, A. Mehrotra, R. Lee, J. Fernandez-Marques, N. Lane, H. Wen, How Much Is Hidden in the NAS Benchmarks? Few-Shot Adaptation of a NAS Predictor	Preprint, 2023
[7]	C. Lance, M. Luecken, D. Burkhardt, R. Cannoodt,, Multimodal Single Cell Data Integration Challenge: Results and Lessons Learned	PMLR, 2022
[8]	E. Dupont*, H. Loya*, M. Alizadeh, A. Golinski, Y. W. Teh, A. Doucet, COIN++: Neural Compression Across Modalities	TMLR, 2022
[9]	H. Loya, P. Poduval, D. Anand, N. Kumar, A. Sethi, Uncertainty Estimation in Cancer Survival Prediction	CLR Workshop, 2020
[10]	P. Poduval, H. Loya, A. Sethi, Functional Space Variational Inference for Uncertainty Estimation in Computer Aided Diagnosis	MIDL, 2020
Awards		
[1]	Global Talent Visa: Awarded for Exceptional Promise, endorsed by the Royal Society	Sep 2025
[2]	Clarendon Scholarship: Awarded to top 200 D.Phil. applicants based on academic merit	May 2020
[3]	Medical Sciences CDT Award: D.Phil. funding, worth \$100,000	May 2020
[4]	Undergraduate Research Award: For research towards master's thesis	Apr 2020
[5]	IIT Institute Academic Prize: Awarded to the top 2 students out of $100+$	Jun 2017
Me	ntorship and Teaching	
[1]	Project Mentorship:	
	<ul> <li>D.Phil. project: Genomic Medicine and Statistics (co-supervised with Prof. Pier Palamara</li> <li>M.Sc. project: Statistical Science (co-supervised with Prof. Simon Myers)</li> </ul>	Oxford, UK Oxford, UK
[2]	<ul> <li>Teaching:</li> <li>Senior Tutor: Applied Probability; Teaching Assistant: Advanced Simulation Methods</li> <li>Teaching Assistant: Communication Systems, Differential Equations, Complex Analysis</li> </ul>	Oxford, UK Mumbai, India
Tall	ks, Academic Services, and Interests	
[1]	Oral presentation at the Ancient Genomes conference, among top 10% abstracts	Nov 2025
[2]	Oral presentation at the Probabilistic models in Genetics (ProbGen), among top 10% abstra	octs Apr 2024
[3]	Invited talks: UCL Genetics Institute (London), BDI/WHG Genomics Seminar, CHG Lunchtime Talk (Oxford)	
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	Languages: English (Fluent), Hindi (Native)	
	Interests: Traveling, Cricket, Hiking, and Sport climbing	
Skills		

**Programming:** Python, R, Julia, C++, Git, PyTorch, JAX **Bioinformatics:** Bcftools, Samtools, Picard, UKB-RAP **Statistical Genetics:** Data QC, GWAS, Fine-mapping, Polygenic scores, LD score regression, Mendelian randomization