LINO ANDRÉ FONSECA FERREIRA www.lfe.pt linoafferreira@gmail.com

EDUCATION	University of Oxford DPhil (PhD) in Genomic Medicine and Statistics • 2019–2023
	Thesis: Analysis of Epistasis in Human Compex Traits
	• Supervisors: Profs. Simon R. Myers (primary) and Pier Francesco Palamara
	· Awarded <i>Wellcome Trust Studentship</i> (incl. £40,000 research grant) · 2019–2023
	· Awarded Jesus College Graduate Scholarship · 2022–2023
	University of Edinburgh MSc in Statistics with Data Science (Distinction) \cdot 2017–2018
	\cdot Awarded <i>School of Mathematics Prize</i> given to the top two students \cdot 2018
	University of Oxford MPhil in Economics · 2012–2014
	Universidade Nova de Lisboa Bachelor's in Economics · 2009–2012
	\cdot Awarded <i>Merit Scholarship</i> for the top combined 1st- and 2nd-year grade \cdot 2011
	· Exchange semester at Bocconi University · 2011
EXPERIENCE	University of Oxford, Department of Statistics Postdoctoral Research Associate · Feb 2024–present
	University of Oxford, Department of Statistics Teaching Assistant · Oct 2021–Jan 2022
	University of Oxford , Department of Economics Research Assistant · Sep 2015–Jul 2017
	\cdot Analysis and visualisation of survey, GIS and satellite data from East African cities
	European Commission, Directorate-General for Competition \cdot Brussels, Belgium Trainee \cdot Mar–Jul 2015
	\cdot Analysis of the economic impact of mergers in the energy and automotive sectors
	NOVAFRICA · Maputo, Mozambique Research Assistant · Jul–Nov 2014
	 Coordinated fieldwork for a research project investigating the use of mobile phone-based money transfer technology
	McKinsey & Company \cdot Lisbon, Portugal and Luanda, Angola Business Analyst Intern \cdot Jul 2013 and Jul 2012
	\cdot Worked in the Portuguese telecom and Angolan aviation industries
PREPRINT	Hu, S., Ferreira , L.A.F. , Shi, S., Hellenthal, G.,* Marchini, J.,* Lawson, D.J,* Myers, S.R.* Leveraging fine-scale population structure reveals conservation in genetic effect sizes between human populations across a range of human phenotypes. <i>bioRxiv</i> (2023)
LANGUAGES	Portuguese (native) · English (fluent) · French (intermediate)
SKILLS	Statistics: RScripting: Bash, PythonGenomics: PLINK, SnakemakeOther: Git, LATEX