

PROFESSOR OF STATISTICAL GENETICS, UNIVERSITY OF BRISTOL

□ +44(0)7930951876 | **☑** g.hemani@bristol.ac.uk | **⋒** gibranhemani.com | **☑** explodecomputer

PRESENT APPOINTMENT

MRC Integrative Epidemiology Unit, University of Bristol

UK

Professor 2023, Associate Professor 2021, Senior Research Fellow 2018, Research

Jan-14 to present

Fellow 2014 (Advisor: George Davey Smith)

PREVIOUS APPOINTMENTS __

Queensland Brain Institute, University of Queensland *Post doctoral statistical geneticist (Advisor: Peter Visscher)*

Australia

Jan-12 to Dec-13

ACADEMIC QUALIFICATIONS _____

University of Bristol Bristol

Fellow of the Higher Education Academy, UK Sep-19 to Sep-21

University of Edinburgh

Edinburgh

PhD in Quantitative Genetics (Advisors: Chris Haley and Sara Knott)

Oct-07 to Aug-11

• Genetics Society 'Best PhD thesis in population and quantitative genetics' 2011

University of Nottingham

Nottingham

BSc (hons) Genetics (2:1)

Sep-03 to Jun-06

SPECIAL AWARDS, HONOURS AND DISTINCTIONS _____

Rising Stars of Science Award

Research.com

Ranked 2nd nationally and 137th internationally for H-index within 12 years of first paper

2022

Fellow of the Higher Education Academy.

CREATE, UoB

2022

Highly Cited Researcher Award

Clarivate / Web of Science

Top 1% of researchers internationally (assigned 'Cross-field' category)

2021-2023

Sir Kenneth Mather Memorial prize

The Genetics Society

Best PhD thesis in quantitative and population genetics

2011

Publications summary _____

• H-index: **71**

• i10-index: **152**

• Number of publications since indepdence (2018): 168

A full list of publications since independence is at the end of the document.

RESEARCH GRANTS _

Total income as PI, from 5 grant(s): 3,548,749 GBP

Total income as CI, from 7 grant(s): 4,021,252 GBP

Current income as PI, from 1 grant(s): 1,971,091 GBP

Current income as CI, from 2 grant(s): 2,540,580 GBP

Building the OpenGWAS platform towards a sustainable model for
scientific discovery and translation

Wellcome Trust	2023
• Amount: 100,000 GBP; Role: Co-PI; Dates: 2023-07-18 to 2024-07-18; Proportion: 0%	
IEU3 - Programme 1 Mendelian randomisation MRC	2022
• Amount: 1,971,091 GBP; Role: PI; Dates: 2023-04-01 to 2028-03-31; Proportion: 100%	
BRC2 - Translational Population Science Theme NIHR	2022
• Amount: 1,912,067 GBP; Role: Co-I; Dates: 2022-12-01 to 2027-11-30; Proportion: 0%	
Causes and consequences of mental disorders: The environmental and genetic influences of parents on offspring. NIH	2022
• Amount: 766,480 USD; Role: Co-I; Dates: 2022-07-15 to 2027-05-31; Proportion: 5%	
Aetiological Epidemiology Biogen	2020
• Amount: 284,525 GBP; Role: Co-I; Dates: 2020-09-01 to 2022-08-31; Proportion: 5%	
The causal map of the human phenome Wellcome Trust and Royal Society, Sir Henry Dale Fellowship • Amount: 1,356,578 GBP; Role: PI; Dates: 2018-01-04 to 2023-06-30; Proportion: 100%	2017
Classifying mechanisms of pleiotropy to improve causal modelling BBSRC and GlaxoSmithKline, CASE studentship • Amount: 100,000 GBP; Role: PI; Dates: 2017-10-01 to 2021-09-30; Proportion: 5%	2017
Pathways to self-harm: Biological mechanisms and genetic contribution Medical Research Councal and Medical Research Foundation • Amount: 372,334 GBP; Role: Co-I; Dates: 2017-10-01 to 2019-10-01; Proportion: 5%	2017
Identification of Traits and Biomarkers for Prediction of Huntington's Disease Phenotypes using Novel causal analysis Methodologies Cure Huntington's Disease Initiative • Amount: 117,059 GBP; Role: Co-I; Dates: 2017-04-01 to 2019-03-31; Proportion: 10%	2017
Translation of MR for drug target identification GlaxoSmithKline • Amount: 349,099 GBP; Role: Co-I; Dates: 2017-01-01 to 2020-01-01; Proportion: 5%	2017
Translation of MR for drug target identification Biogen Amount: 436,165 USD; Role: Co-I; Dates: 2017-01-01 to 2020-01-01; Proportion: 5%	2017

Dissecting genetic interactions in gene expression	2012
 University of Queensland, Early Career Research Grant Amount: 34,000 AUD; Role: PI; Dates: 2013-01-01 to 2013-12-31; Proportion: 2% 	2013
Dissecting genetic interactions in complex traits CASE Studentship, BBSRC and Monsanto	2007
• Amount: 100,000 GBP; Role: PI; Dates: 2007-09-01 to 2011-08-30; Proportion: 100	
INDICATIONS OF EXTERNAL RECOGNITION	
Editorships	
PLoS Computational Biology	
Invited guest editor	2017
Appointment to national or international bodies	
Appointment to national of international bodies	_
PLoS Computational Biology	
Expert evaluation panel for the Our Future Health study	2021
Invited member of the Genotype Chip Design Committee for the Our Future	2019 - 2021
Health Study, that will be used to genotype up to 5 million people in the UK	
Invitations for degree examinations	
Richard Packer	University of Leicester
PhD thesis external examiner	2022
Jisu Shin	University of South
	Australia
Dissertation of Master of precision medicine external examiner	2020
Edward Steere	Witswatersrand University,
Discortation of Master of Science in Engineering external evaminer	SA 2016
Dissertation of Master of Science in Engineering external examiner	2016
Invited lectures	
NIH/NIDA Genetics and Epigenetics meeting, Maryland USA	Invited talk
Primer in Mendelian randomisatio	2024
ECNP Network workshop, University of Muenster, Germany	Invited talk
Mendelian randomisation for causal inference of molecular phenotypes on	2024
complex traits	2024
American Society of Human Genetics	Invited talk
The importance of families in future biobanks	2023
University of Galway	Invited talk
The future of genetic biobanks	2023
University of Edinburgh	Invited talk
Recent developments in Mendelian randomisation	2023

3

	1 2 1 H
Epigenomics of Common Disease 2023, Cambridge Challenges and future directions for causal inference of molecular traits	Invited talk 2023
Centre for Health Sciences Education conference, Bristol Workshop: Decolonising the Medicul School Curriculums	Invited workshop 2023
BenevolentAl seminar series The MR-Base and OpenGWAS projects	Invited talk 2022
Cure Huntington's Disease Initiative seminar series Exploring the assumptions in Mendelian randomisation	Invited talk 2022
BILT Compassionate Conference Decolonising the Bristol medical school curriculum	Conference plenary 2022
Mendel @ 200 conference, Bristol Contemporary genome-wide association studies in the light of Mendelian inheritance	Conference plenary 2022
Broad Institute (Harvard and MIT) methods seminar Within-sibship GWAS for Mendelian randomisation	Invited talk 2022
DECOLWEEK, Bristol Decolonising your discipline	Invited panel member 2021
Mendelian Randomization conference, Bristol Results from large scale consortia	Invited session chair 2021
GSK seminar series Mendelian randomisation for molecular exposures	Invited talk 2021
Channing Network seminar series, Harvard University Using Mendelian randomisation to detect causal influences of molecular features on complex traits	Invited talk 2021
The Trøndelag Health Study, Norway Multi-omics in longitudinal cohorts	Invited workshop lead 2020
International Agency for Research on Cancer, France Collider bias in Covid-19 research	Invited talk 2020
Elizabeth Blackwell Institute data week keynote New data on Covid-19 is undermined by old statistical problems	Invited talk 2020
GRC Quantitative Genetics conference, Italy Genetic architecture of complex traits	Invited session chair 2019
Mendelian Randomization conference, Bristol Genetics of DNA methylation	Conference plenary 2019
SEGEG, University of Oxford Automating Mendelian randomization	Invited talk 2018
Edinburgh Alliance in Quantitative Genetics Machine learning in Mendelian randomization	Invited talk 2018
University College London Causal graph of the human phenome]	Invited talk 2018

Related administration Leading group on Covid-19 epidemiology MRC IEU 2020-2021 • Initiated project, and recruited group of 12 researchers • Provided analysis of ZOE symptom tracker app for external collaborators Culminated in four publications, numerous presentations including to SAGE and HDRUK Member of the UoB Covid modelling subgroup University of Bristol 2020-2021 **Initiating the OpenGWAS consortium** MRC. IEU 2020 to present • The OpenGWAS data infrastructure receives 2 million queries per week The consortium brings together researchers who develop software for GWAS summary data • Plans to expand the invitation to international collaborators **SEGEG conference organiser** South of England Genetic Epidemiology Group 2019 • Organised the long-running SEGEG conference to be held in Bristol for the first time Leading work package on MR method development (programme 1) MRC IEU 2018-2023 Line managing one post doctoral scientist Leading statistics and informatics theme in epigenetics programme 4 MRC. IEU 2018-2022 **Leading Hemani research group** MRC IEU 2018 to present • Four post-docs and four PhD students Weekly group meetings including pastoral and career support **ALSPAC Board of Directors ALSPAC** 2017 to present Leading genetics work package ALSPAC. 2017 to present Line managing one post doctoral scientist ACADEMIC LEADERSHIP AND CITIZENSHIP ______ Academic leadership in the University $_$ **MRC Integrative Epidemiology Unit 3.0** University of Bristol Programme Leader 2023-2028

• Co-leading (with George Davey Smith) Programme 1 of seven programmes within the unit

Medical Anti-Racism Taskforce University of Bristol Group member 2020 to present

Leading Special interest group on decolonising the curriculum

UoB Covid-19 Situation Report

Software developer

University of Bristol 2020 to present

• Web app developed for internal use that provides daily updates and case mapping

Used daily by the Incident Management Team in targeting Covid-19 actions

Decolonising the medical school curriculum

Co-lead

University of Bristol

2020 to present

• Co-leading (with Jo Hartland) a group of 10 volunteers to develop, pilot and deploy a curriculum review framework for the medical school (http://decolbms.org.uk)

BRMS Equality Diversity Inclusion group

University of Bristol

Group member

2018 to present

• Working in career progression subgroup

USS pension calculator

University of Bristol

Software developer

2018 to present

- Developed a web-app that allows members of the USS pension scheme to calculate changes to the pension under proposed valuation changes
- Used widely by universities across the country
- http://www.uss-pension-model.com/

Teaching peer review

University of Bristol

Reviewer 2018

• Causal inference short course, University of Bristol

Professional activities outside the University _

- 2021 Expert evaluation panel for the Our Future Health study
- 2019 Invited member of the Genotype Chip Design Committee for the Our Future Health Study, that will be used to genotype up to 5 million people in the UK
- 2017 Regular peer review for various grant bodies including the MRC, Wellcome Trust, Cancer Research UK
 - Regular contributor to open source software projects (e.g. see
- 2012 https://github.com/explodecomputer/random-metal and https://github.com/explodecomputer/ldsc/)
- 2011 Regular peer review for 20+ academic journals

Contributions to society _

- Hosting asylum seekers and refugees, working with Bristol City Council and the *Refugees at Home* charity
- 2015 Contributing member to the charity *Statisticians Without Borders*
- 2014 Provide web and software support to local vegan organisations

Entrepreneurship, enterprise and partnerships _

I have formal partnerships with GlaxoSmithKlein, Biogen, Pfizer and CHDI that arose through developing the MR-Base platform. This has led to five post-doctoral positions and one PhD position being funded by these organisations, and the development of a standardised contract system to enable future such collaborations to occur.

Good citizenship		
2023	Bristol Medical School representative of the University Research Institute task and finish group	
2022 -	Member of the University of Bristol Senate (class 4 - Academic Staff)	
2020 -	Leading a Special Interest Group on Decolonising the Curriculum, which will bring in a framework for all courses across the medical school for dealing with implicit bias within their learning materials	
2020 -	Member of the Bristol Clear mentoring scheme (currently mentoring two early career researchers)	
2020	Developed software for the University's Incident Management Team which maps new Covid-19 cases amongst students across the region in real time. This software is used in daily team management meetings	
2016 -	Participated in numerous mock interview panels for research fellowships	
2015 -	Extensive software development for the MRC IEU research community, most notably the MR-Base and OpenGWAS software platforms which I created and maintain for others to use. I am pleased that this has grown to be a platform for numerous research papers, and contributed substantially to research grants, many of which I am not named on	
2014 -	Voluntary curation and documentation of shared data resources	
2014 -	Web and software development for the ALSPAC project (e.g. see http://variables.alspac.bris.ac.uk/ and https://github.com/explodecomputer/alspac)	
2014 -	Throughout my time in Bristol I have made it a priority to provide informal training to early career researchers, particularly in genetics, software development and reproducible research.	

7