Consortia in the author list

Cell Press journals offer several options for listing the membership of a consortium that is part of the author list of a paper. If one of the authors of your paper is a consortium, please let us know which option(s) you would like. None is required, and you can choose one option, more than one, all, or none. And of course you are welcome to reach out to Cell Press with any questions you may have.

Option 1: Secondary author list

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Am J Hum Genet. 2022 Apr 7;109(4):647-668. doi: 10.1016/j.ajhg.2022.02.010. Epub 2022 Mar 2.

The individual and global impact of copy-number variants on complex human traits

Chiara Auwerx [1], Maarja Lepamets [2], Marie C Sadler [3], Marion Patxot [4], Miloš Stojanov [5], David Baud [5], Reedik Mägi [6], Estonian Biobank Research Team [6]; Eleonora Porcu [7], Alexandre Reymond [8], Zoltán Kutalik [9]

Collaborators, Affiliations – collapse

Collaborators

Estonian Biobank Research Team: Tõnu Esko, Andres Metspalu, Lili Milani, Reedik Mägi, Mari Nelis

Affiliations

1 Center for Integrative Genomics, University of Lausanne, Lausanne 1015, Switzerland; Department of Computational Biology, University of Lausanne, Lausanne 1015, Switzerland; Swiss Institute of Bioinformatics, Lausanne 1015, Switzerland; University Center for Primary Care and Public Health, Lausanne 1010, Switzerland.

2 Institute of Molecular and Cell Biology, University of Tartu, Tartu 51010, Estonia; Estonian Genome Centre, Institute of Genomics, University of Tartu, Tartu 51010, Estonia.

3 Swiss Institute of Bioinformatics, Lausanne 1015, Switzerland; University Center for Primary Care and Public Health, Lausanne 1010, Switzerland.

Note: the **secondary author list** will not appear in the print or PDF version of the paper.

Option 2: Consortia paragraph

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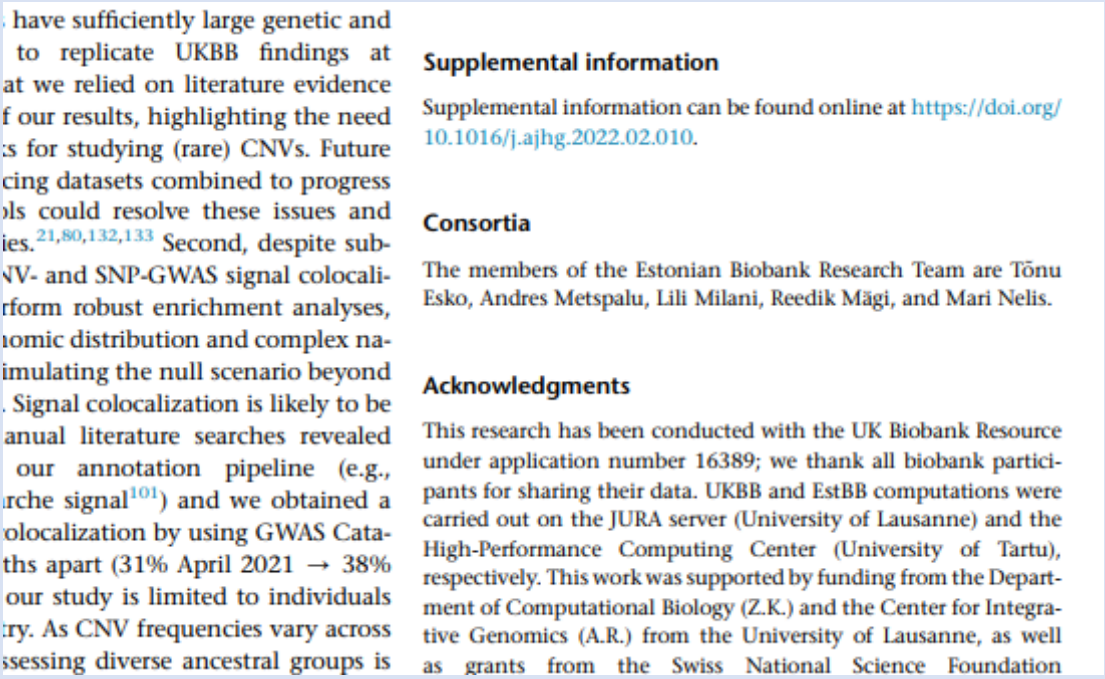

have sufficiently large genetic and
to replicate UKBB findings at
at we relied on literature evidence
f our results, highlighting the need
:s for studying (rare) CNVs. Future
cing datasets combined to progress
ols could resolve these issues and
ies.[21,80,132,133] Second, despite sub-
NV- and SNP-GWAS signal colocali-
rform robust enrichment analyses,
nomic distribution and complex na-
imulating the null scenario beyond
. Signal colocalization is likely to be
anual literature searches revealed
our annotation pipeline (e.g.,
rche signal[101]) and we obtained a
olocalization by using GWAS Cata-
ths apart (31% April 2021 → 38%
our study is limited to individuals
ry. As CNV frequencies vary across
ssessing diverse ancestral groups is

Supplemental information

Supplemental information can be found online at https://doi.org/10.1016/j.ajhg.2022.02.010.

Consortia

The members of the Estonian Biobank Research Team are Tõnu Esko, Andres Metspalu, Lili Milani, Reedik Mägi, and Mari Nelis.

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Option 3: Supplemental material

Options 1 and 2 are limited to names of consortium members. If you wish to include additional information, such as affiliations or email addresses of individuals, these details can be provided in supplemental files. We accept Excel tables and PDFs, as shown in these two examples.

	A	B	C	D	E	F
1	**Table S3. Undiagnosed disease network member list.**					
2	**First**	**Middle**	**Last**	**Affiliation**	**Email**	**Role in the UDN (ex. PI, site coordinator)**
3	Mercedes	E.	Alejandro	BCM Clinical	[redacted]@bcm.edu	Site coordinator
4	Mahshid	S.	Azamian	BCM Clinical	[redacted]@bcm.edu	Site coordinator, clinical assistant
5	Carlos	A.	Bacino	BCM Clinical	[redacted]@bcm.edu	Co-PI
6	Ashok		Balasubramanyam	BCM Clinical	[redacted]@bcm.edu	Lead clinician, adult
7	Lindsay	C.	Burrage	BCM Clinical	[redacted]@bcm.edu	Sequence analysis team, pediatric genetics
8	Hsiao-Tuan		Chao	BCM Clinical	[redacted]@bcm.edu	Pediatric neurology
9	Gary	D.	Clark	BCM Clinical	[redacted]@texaschildrens.org	Pediatric neurology
10	William	J.	Craigen	BCM Clinical	[redacted]@bcm.edu	Leadership team, pediatric genetics, biochemical genetics
11	Hongzheng		Dai	BCM Clinical	[redacted]@bcm.edu	Sequence analysis team
12	Shweta	U.	Dhar	BCM Clinical	[redacted]@bcm.edu	Adult genetics
13	Lisa	T.	Emrick	BCM Clinical	[redacted]@bcm.edu	Leadership team, pediatric neurology
14	Alica	M.	Goldman	BCM Clinical	[redacted]@bcm.edu	Adult neurology
15	Neil	A.	Hanchard	BCM Clinical	[redacted]@bcm.edu	Pediatric Genetics & Genomics Core

Table S3. UDN Member List

Myocardial Infarction Genetics Consortium

Diego Ardissino[1-3], Usman Baber[4], Matthew J. Bown[5,6], Mark D. Chaffin[7-9], Rajiv Chowdhury[10,11], John Danesh[10,12], Roberto Elosua Llanos[13-15], Valentin Fuster[4,16], Namrata Gupta[7], Sekar Kathiresan[7,8,17,18], Amit V. Khera[7-9], Ruth McPherson[19], Olle Melander[20,21], Marju Orho-Melander[22], Danish Saleheen[23,24], Nilesh Samani[5,6], Heribert Schunkert[25], Minxian X. Wang[7,8,26], Hugh Watkins[27]

1 - ASTC: Associazione per lo Studio Della Trombosi in Cardiologia, Pavia, Italy.; 2 - Azienda Ospedaliero-Universitaria di Parma,ÉParma,Italy.; 3 - Universit˜ degli Studi di Parma, Parma, Italy.; 4 - Department of Medicine, Icahn School of Medicine at Mount Sinai, New York, NY 10029.; 5 - Department of Cardiovascular Sciences, University of Leicester, Leicester, UK.; 6 - NIHR Leicester Biomedical Research Centre, Glenfield Hospital, Leicester UK.; 7 - Program in Medical and Population Genetics, Broad Institute of Harvard and MIT, Cambridge, MA, USA, 02142.; 8 - Center for Genomic Medicine, Department of Medicine, Massachusetts General Hospital, Boston, Massachusetts, USA, 02114.; 9 - Cardiovascular Research Center, Massachusetts General Hospital, Boston, MA, USA, 02114.; 10 - MRC/BHF Cardiovascular Epidemiology Unit, Department of Public Health and Primary Care, University of Cambridge, Cambridge, UK.; 11 - Centre for Non-Communicable disease Research (CNCR), Bangladesh.; 12 - The National Institute for Health Research Blood and Transplant Research Unit (NIHR BTRU) in Donor Health and Genomics at the University of Cambridge, Cambridge, UK.; 13 - Cardiovascular Epidemiology and Genetics, Hospital del Mar Research Institute, Barcelona, Spain.; 14 - CIBER Enfermedades Cardiovasculares (CIBERCV), Barcelona, Spain.; 15 - Facultat de Medicina, Universitat de Vic-Central de Catalu–a, Vic, Spain.; 16 - Centro Nacional de Investigaciones Cardiovasculares Carlos III (CNIC), Madrid, Spain.; 17 - Department of Medicine, Harvard Medical School, Boston, MA, USA, 02115.; 18 - Verve Therapeutics, Cambridge, MA, USA 02139.; 19 - Ruddy Canadian Cardiovascuar Genetics Centre, University of Ottawa Heart Institute, Ottawa, Canada.; 20 - Department of Clinical Sciences, Diabetes and Endocrinology, Lund University Diabetes Centre, Malmö, Sweden.; 21 - Department of Emergency and Internal Medicine, SkŒne University Hospital, Malmö, Sweden.; 22 - Department of Clinical Sciences, Lund University, Malmö, Sweden.; 23 - Department of Biostatistics and Epidemiology, University of Pennsylvania, Philadelphia, Pennsylvania, USA.; 24 - Center for Non-Communicable Diseases, Karachi, Pakistan.; 25 - Deutsches Herzzentrum MŸnchen, Technische UniversitŠt MŸnchen, Deutsches Zentrum fŸr Herz-Kreislauf-Forschung, MŸnchen, Germany.; 26 - Cardiovascular Disease Initaitive, Broad Institute of Harvard and MIT, Cambridge, MA, USA, 02142.; 27 - Cardiovascular Medicine, Radcliffe Department of Medicine and the Wellcome Trust Centre for Human Genetics, University of Oxford, Oxford, UK.;