

Virtual Conference Agenda

Start (GMT)	Finish (GMT)	Presenter details
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Tuesday 2 November 2021

12:00 12:10 Welcome

Scientific Programme Committee:

[Marta Mirazon Lahr, University of Cambridge, UK](#)

[Lluis Quintana-Murci, Institut Pasteur & Collège de France, France](#)

[Carina Schlebusch, Uppsala University, Sweden](#)

[Michael Westaway, The University of Queensland, Australia](#)

12:10 13:00 Keynote lecture 1

Introduction to the session

Chair: Lluis Quintana-Murci

12:10 12:40 The history of human functional evolution: adaptations to high altitude

[Anna Di Rienzo, University of Chicago, USA](#)

12:40 13:00 Q&A

Chair: Lluis Quintana-Murci

Moderator: Marta Mirazon Lahr

13:00 13:20 Break

13:20 14:50 Session 1: Human adaptation

Introduction to the session

Chair: Lluis Quintana-Murci

13:20 13:40 Humans adapting to their new homes: the fine-scale evolution of cold receptor *TRPM8* in Europe

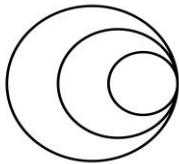
[Aida Andres, UCL, UK](#)

13:40 14:00 The Middle Pleistocene cranium from Harbin establishes a new human lineage in China

[Chris Stringer, Natural History Museum, UK](#)

14:00 14:10 Human-chimp hybrid cells reveal gene regulatory evolution underlying skeletal divergence

David Gokhman, The Weizmann Institute of Science, Israel



14:10 14:20 The genomic signatures of natural selection in recently admixed human populations
Sebastian Cuadros Espinoza, Institut Pasteur, France

14:20 14:50 Q&A
Chair: Lluis Quintana-Murci
Moderator: Marta Mirazon Lahr

14:50 16:50 Poster session I

14:50 15:05 Poster session I lightning talks

15:05 15:35 Poster session I

15:35 17:25 Session 2: Human evolution and disease

Introduction to the session

Chair: Marta Mirazon Lahr

15:35 15:55 The relevance of life history hypotheses to nutrition intervention promoting child growth and development

[Rihlat Said Mohamed, University of Cambridge, UK](#)

15:55 16:15 Changing patterns in health and disease: Insights into complex hunter-gatherers and farmers on the periphery of Southeast Asia

[Marc Oxenham, University of Aberdeen, UK](#)

16:15 16:35 Using ancient and present-day DNA to infer the changing landscape of natural selection

[Iain Mathieson, University of Pennsylvania, USA](#)

16:35 16:45 The architecture of gene regulation in Indonesia identifies QTLs associated with global and local ancestries

Irene Gallego Romero, University of Melbourne, Australia

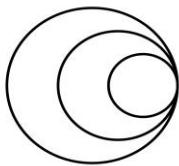
16:45 16:55 The impact of demography in functional variation: whole-exome analysis in Tunisian Imazighen and Arabs

Marcel Lucas-Sánchez, Universitat Pompeu Fabra, Spain

16:55 17:25 Q&A

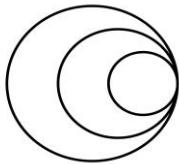
Chair: Marta Mirazon Lahr

Moderator: Lluis Quintana-Murci

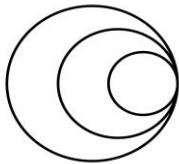


Wednesday 3 November 2021

12:00	13:30	Session 3: Ancient and modern genetic admixture
		Introduction to the session
		Chair: Carina Schlebusch
12:00	12:20	Genomic insights into the population history and biological adaptation in Pacific Islands Etienne Patin, Institut Pasteur, France
12:20	12:40	The population history of northern and southern East Asia Qiaomei Fu, Chinese Academy of Sciences, China
12:40	12:50	A Philippine ethnic group with the most Denisovan DNA Maximilian Larena, Uppsala University, Sweden
12:50	13:00	A Genetic History of Continuity and Mobility in the Iron Age Central Mediterranean Hannah Moots, University of Chicago, USA
13:00	13:30	Q&A Chair: Carina Schlebusch Moderator: Michael Westaway
13:30	13:50	Break
13:50	15:20	Session 4: Inferring human demography from genomic data
		Introduction to the session
		Chair: Lluis Quintana-Murci
13:50	14:10	Human germline mutation and the erratic evolutionary clock Priya Moorjani, UC Berkeley, USA
14:10	14:30	The phenotypic impact of Neandertal introgression in modern humans Michael Dannemann, Tartu University, Estonia
14:30	14:40	The mixed genetic origin of the first farmers of Europe Nina Marchi, Institute for Ecology and Evolution, Switzerland
14:40	14:50	Population diversity of Austroasiatic speakers: Complex substructure, multiple admixture and language replacement Debashree Tagore, National Institute of Biomedical Genomics, India
14:50	15:20	Q&A Chair: Lluis Quintana-Murci Moderator: Carina Schlebusch



15:20	16:05	Poster session II
15:20	15:35	Poster session II lightning talks
15:35	16:05	Poster session II
16:05	17:35	Session 5: Population structure
		Introduction to the session
		Chair: Carina Schlebusch
16:05	16:25	Population structure and demographic history of Bantu speakers from South Africa Dhriti Sengupta, University of the Witwatersrand, South Africa
16:25	16:45	Reconstructing human migrations from genomic data across Latin America and the Pacific Andres Moreno Estrada, Langebio, Cinvestav, Mexico
16:45	16:55	Haplotype-based recent ancestry sharing methods reveal fine-scale genetic structure in West and Central Africa Nancy Bird, University College London, UK
16:55	17:05	Population structure in Europe has been stable from the Iron Age onward, despite high mobility Margaret Antonio, Stanford University, USA
17:05	17:35	Q&A Chair: Carina Schlebusch Moderator: Marta Mirazon Lahr



Thursday 4 November 2021

12:00 – 13:40 Session 6: Palaeoanthropology/human evolution

Introduction to the session

Chair: Marta Mirazon Lahr

12:00 – 12:20 New Neanderthal remains from Shanidar Cave, Iraqi Kurdistan
[Emma Pomeroy, University of Cambridge, UK](#)

12:20 – 12:40 The Paleobiology, paleoecology, and biogeography of the earliest *Australopithecus*:
New insights from Woranso-Mille (Ethiopia)
[Yohannes Haile-Selassie, Cleveland Museum of Natural History, USA](#)

12:40 – 12:50 A revision of the Border Cave 1 cranium
Amelie Beaudet, University of Cambridge, UK

12:50 – 13:00 Deciphering the biological bases of endocranial globularity in modern humans
Barbara Molz, Max Planck Institute for Psycholinguistics, The Netherlands

13:00 – 13:10 The Developmental Impacts of Natural Selection on Human Pelvic Morphology
Terence Capellini, Harvard University, USA

13:10 – 13:40 Q&A
Chair: Marta Mirazon Lahr
Moderator: Michael Westaway

13:40 – 14:00 Break

14:00 – 15:20 Session 7: Languages, culture and history

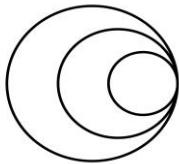
Introduction to the session

Chair: Carina Schlebusch

14:00 – 14:20 Language and human history
[Michael Dunn, Uppsala University, Sweden](#)

14:20 – 14:40 The roles of contact and inheritance in generating the linguistic diversity of pre-Bantu
southern Africa
[Anne-Maria Fehn, CIBIO, Portugal](#)

14:40 – 14:50 Assessing human genome-wide variation in the Massim region of Papua New Guinea and
implications for the Kula trading tradition
Dang Liu, Max Planck Institute for Evolutionary Anthropology, Germany



14:50 15:20 Q&A
Chair: Carina Schlebusch
Moderator: Marta Mirazon Lahr

15:20 15:40 Break

15:40 16:30 Keynote lecture 2

Introduction to the session
Chair: Marta Mirazon Lahr

15:40 16:10 The Sapiens-Neandertal Connection
[Jean-Jacques Hublin, Max Planck Institute/College de France, Germany/France](#)

16:10 16:30 Q&A
Chair: Marta Mirazon Lahr
Moderator: Lluis Quintana-Murci

16:30 16:40 Closing remarks

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