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COMMITTEE ON POPULATION

Use of Race, Ethnicity, and Ancestry as Population Descriptors in Genomics Research

**Use of Race, Ethnicity, and Ancestry as
Population Descriptors in Genomics Research**

PUBLIC WORKSHOP

April 4, 2022

All times listed in Eastern Standard Time

Register Here: <https://www.eventbrite.com/e/public-workshop-on-use-of-population-descriptors-in-genomics-research-tickets-287946785587>

11:00 am ET Welcome and Goals for the Workshop

Aravinda Chakravarti, *Committee Co-Chair*

Director, Center for Human Genetics and Genomics Muriel G & George W Singer Professor of
Neuroscience & Physiology
New York University Grossman School of Medicine

Charmaine Royal, *Committee Co-Chair*

Robert O. Keohane Professor of African & African American Studies, Biology, Global Health,
and Family Medicine & Community Health
Director, Duke Center on Genomics, Race, Identity, Difference and Duke Center for Truth,
Racial Healing & Transformation
Duke University

Session I: Historical and Current Use of Population Descriptors in Genomics Research

Moderator: Sandra Soo-Jin Lee, Columbia University

Session Objectives:

- To explore historical use of population descriptors to better understand current use
- To examine whom we study in genomic investigations
- To explore why we identify individuals and populations in genomic studies

- To examine and identify the criticisms and challenges in current use of population descriptors in genomics research

11:10 am Brief introduction to the session by the moderator

11:15 am Speakers' Talks (15 minutes each)

Pilar Ossorio

Professor of Law and Bioethics
University of Wisconsin-Madison, Law School
University of Wisconsin-Madison, Medical School

Joseph Graves

Professor of Biological Sciences
PI: IBIEM@ AT and BEACON@A&T
Associate Director, Triangle Center for Evolutionary Medicine
Department of Biology
North Carolina A&T State University

Andrew Clark

Jacob Gould Schurman Professor of Population Genetics
Nancy and Peter Meinin Family Investigator
Associate Director, Cornell Center for Comparative and Population Genomics
Interim Chair, Department of Computational Biology
Cornell University

Rina Bliss

Associate Professor of Sociology
Rutgers University

12:15 pm Q&A with speakers

1:00 pm Break

Session II: Future Use of Population Descriptors in Genomics Research

Moderator: Rick Kittles, City of Hope

Session Objectives:

- To consider the diverse types of population and individual descriptors (e.g. origins, definitions, and usage in the U.S., implications for non-U.S. participants)
- To discuss possible ideal descriptors of populations and individuals
- To consider standardized or ideal systems of population descriptors

1:30 pm Brief introduction to the session by the moderator

1:35 pm Speakers' Talks (15 minutes each)

Tesfaye Mersha

Associate Professor of Human Quantitative Genetics
Department of Pediatrics
University of Cincinnati
Cincinnati Children's Hospital Medical Center

Melinda Mills

Director, Leverhulme Centre for Demographic Sciences
Nuffield Professor of Sociology
University of Oxford

Joanna Mountain

Consultant
23andMe

Eimear Kenny

Founding Director, Institute for Genomic Health
Professor of Genetics and Medicine
Icahn School of Medicine at Mount Sinai

Stephanie Malia Fullerton

Professor of Bioethics and Humanities
University of Washington School of Medicine
Adjunct Professor
Departments of Epidemiology, Genome Sciences, and Medicine
University of Washington
Affiliate Investigator, Public Health Sciences Division
Fred Hutchinson Cancer Research Center

2:55 pm **Q&A with speakers**

3:50 pm **Break**

Session III: Community Input on Population Descriptors in Genomics Research

Moderator: Katrina Claw, University of Colorado Denver – Anschutz Medical Campus

Session Objectives:

- To hear from a variety of interested parties on the following topics:
 - What works and doesn't work about the current population descriptors used in genomics research?
 - What could be improved in current use of population descriptors in genomics research?

4:00 pm **Brief introduction to the session by the moderator**

4:05 pm **Speakers' Comments (5 minutes each)**

Catherine Potenski

Chief Editor

Nature Genetics

Donna Cryer

President & CEO

Global Liver Institute

Agustín Fuentes

Professor

Department of Anthropology

Princeton University

Charles Rotimi

President

American Society for Human Genetics

Judit Kumuthini

Bioinformatics Manager

Human Capacity Development Manager: Bioinformatics

University of Western Cape

Shishi Luo

Associate Director, Bioinformatics and Infectious Diseases
Helix Genomics

Julia Ortega

Vice President
iHope Genetic Health
Genetic Alliance

Keolu Fox

Assistant Professor
Department of Anthropology
University of California, San Diego

4:55 pm Concluding Remarks

5:00 pm Adjourn