Time	Monday, April 8, 2024			
09:00 – 10:45	[Main Hall] Educational workshop 1 "Knowledge and Skills for Multi-disciplinary Genomic Healthcare"	[Room 2] Educational workshop 2 "Variant Interpretation / Variant Curation"	[Room 3] Educational workshop 3 "All about Tools (OMIM, ClinGen, GA4GH, Biobank tools, Phenopackets)"	[Room 4] Educational workshop 4 "Pangenomes Workshop"
	- Andreas Laner	- Ada Hamosh, Erin Riggs	- Karen Miga	
10:45 – 11:15	Coffee Break & Exhibition			
11:15 – 12:30	Workshop 1 Continued	Workshop 2 Continued	Workshop 3 Continued	Workshop 4 Continued
12:30 – 13:30	Break			
13:30 – 14:00	[Main Hall] Corporate symposium 1		[Room 2] Corporate symposium 2	
14:00 – 14:10	Opening by Ada Hamosh, Giuseppe Novelli, Juergen Reichardt			
14:10 – 14:45	[Main Hall] Plenary Mike Snyder, Stanford University, US			
15:00 – 16:15	[Main Hall] Symposium 1 - Precision Health		[Room 2] Symposium 2 - Reproductive Genetics	
12:00 – 10:12	Francesca Forzano, NHS, UK		Diana Bianchi, NIH, US "The NIH IDENTIFY study: A prospective evaluation of pregnant	

	Marc Sultan, Roche, Switzerland "Applications of genomics technologies in early drug development at Roche"	women with prenatal cfDNA sequencing results that suggest maternal malignancy" Liborio Stuppia, University of Chieti-Pescara, Italy	
16:15 – 17:15	Poster session & Exhibition & Coffee Break		
17:30 – 18:30	[Main Hall] Introduction to HUGO & HUGO Committee Activities & Representatives from other organizations		
	- Walter Bodmer: HUGO Past, Present, and Future - HUGO Committees Introduction		
18:30 -	[Main Hall] Welcome Reception		

Time	Tuesday, April 9, 2024		
9:00 – 10:15	[Main Hall]  Symposium 3 - Gene Therapy  Alessandro Aiuti, San Raffaele Telethon Institute for Gene Therapy, Italy  Luigi Naldini, Vita-Salute San Raffaele University, Italy	[Room 2] Symposium 4 - Non-coding RNA genes  Roderic Guigo, CRG, Spain "GENCODE: Uncovering and cataloging the transcriptional complexity of the human and mouse genomes" Tim Coorens, Broad Institute, US	
10:15 – 10:45	Coffee Break		
10:45 – 11:00	[Main Hall] Special Session GA4GH Introduction by Peter Goodhand		

	[Main Hall]		
11:00 – 11:20	Special Session		
	HUGO Education Committee Introduction by Dhavendra Kumar		
	[Main Hall] Symposium 5 - Genetics of Complex Diseases	[Room 2] Symposium 6 - Rare Diseases	
11:30 – 12:45	Nicole Soranzo, Human Technopole, Italy	Steven Laurie, CRG, Spain "Solve-RD, Pan-European co-operation elevates the diagnostic rate in the most challenging rare disease cases"	
	Naomi Wray, University of Queensland, Australia "Genetics of psychiatric vs gastrointestinal disorders"	William Gahl, NIH, US  "Rare and undiagnosed diseases: Discoveries and Insights from the NIH undiagnosed diseases program"	
12:45 – 14:00	Lunch		
14:00 – 14:30	[Main Hall] After-lunch corporate symposium 3	[Room 2] After-lunch corporate symposium 4	
14:45 – 16:00	[Main Hall]  Symposium 7 - Computational Biology & AI  Alistair Forrest, University of Western Australia, Australia  "Spatial transcriptomics reveals discrete tumour microenvironments and autocrine loops within ovarian cancer subclones"  Susanna Zucca, enGenome, Italy  "Advancing genomic variant interpretation with artificial intelligence"	[Room 2]  Symposium 8 - Genetics of Infectious Diseases  Jean Laurent Casanova, The Rockefeller University, US  "Human genetic and immunological determinants of infectious diseases"  Christian Happi, Redeemer's University, Nigeria	
16:00 – 17:00	Poster session & Exhibition & Coffee Break		
17:00 – 18:15	[Main Hall] Symposium 9 - Cancer Genomics	[Room 2] Symposium 10 - Single-cell Genomics / Spatial Genomics	
	Serena Nik-Zainal, University of Cambridge, UK	Kerstin Meyer, Wellcome Sanger Institute, UK	

	"Whole cancer genomes: Insights and updates" John Burn, NHS, UK "Lynch syndrome: A common rare disease, hiding in plain sight, the UK experience"	Efrat Shema, Weizmann Institute of Science, Israel "Decoding the epigenome for cancer research and diagnostics"
20:15 –	[Terrazza Caffarelli] Gala Dinner	

Time	Wednesday, April 10, 2024	
9:00 – 09:50	[Main Hall] Opening by Director of Sapienza University of Rome/Ministry of Health	
	[Main Hall]	
9:50 – 10:20	HUGO African Prize	
	Leon Mutesa, University of Rwanda, Rwanda "Intergenerational and epigenetic effects of trauma and PTSD"	
10:20 – 10:50	Coffee Break	
	[Main Hall] HUGO Chen Awards	
10:50 – 12:00	Chen Award for Distinguished Academic Achievement in Human Genetic and Genomic Research : Peter Visscher, University of Queensland, Australia "Height as a model trait in human complex trait genetics"	
	Chen Award of Excellence : Zornitza Stark, University of Melbourne, Australia "Accelerating rare disease diagnosis"	
12:00 – 13:00	Lunch	

13:00 – 13:30	[Main Hall]	[Room 2]
	After-lunch corporate symposium 5	After-lunch corporate symposium 6
13:45 – 15:00	[Main Hall]  Symposium 11 - Transgenerational Inheritance / Epigenetics  Jill Fahrner, Johns Hopkins University, US  "Mendelian disorders of the epigenetic machinery and the use of DNA methylation profiling in diagnosis and discovery"  Simona Giunta, Sapienza University of Rome, Italy  "Complete diploid isogenomic reference human genomes for epigenetic precision"	[Room 2]  Symposium 12 - Challenges in Assessing Pathogenicity of Variants  Alan Rubin, University of Melbourne, Australia "Enabling clinical translation of high-throughput mutagenesis Data with MaveDB"  Doug Speed, Aarhus University, Denmark "Computational tools to improve our understanding of the biological basis of complex traits"  Melissa Cline, UCSC, US "Engaging researchers with patients"
15:00 – 16:00	Poster session & Exhibition & Coffee Break	
16:00 – 17:15	[Main Hall]  Symposium 13 - Pangenomes and Genomic Diversity  Karen Miga, UCSC, US  Charles Lee, The Jackson Laboratory, US	[Room 2] Symposium 14 - Microbiome and Metagenomics  Mireia Valles-Colomer, University of Trento, Italy "The microbiome and health: Focusing on interpersonal transmission" Serena Sanna, Institute for Genetic and Biomedical Research, Italy
17:15 –	[Main Hall] Poster Prize Introduction to <i>HGM2025</i> in <i>Durban, South Africa</i> Closing	·