## Agenda (only eligible sessions listed)

Please note as you are planning your agenda for the meeting, there is no partial credit for sessions. You may only claim credit for one full session per time block. Credits will <u>not</u> be available for on-demand content post-meeting. Credit hours are subject to change. Thank you for your understanding.

Time	Session Type	Session Number	Session Title	P.A.C.E.® Program Number	CME Credit Hours	NSGC Contact Hours	P.A.C.E.® Contact Hours
Tuesday, Nov	ember 5						
		01	A Guide to Card Sort Methods for Engaging Participants and Patients	603-001-24			Contact
10:00 am –	Workshop	02	Getting Started with Biomedical and Genomic Data in the All of Us Researcher Workbench	603-002-24	2.00	2.00	
12:00 pm		03	Using the New Ensembl Genome Browser and Variant Effect Predictor (VEP) to Analyse and Interpret Genomic Variation Data	603-003-24			
2:00 pm –	Manhah an	04	Building Understanding and Practical Skills in Community Engagement for Genetic Research	603-004-24	2.00	2.00	2.00
4:00 pm	Workshop	05	Elevating Education: Engaging with Real Data and Tools in the Cloud	603-005-24	2.00		2.00
-		06	Hidden Features of the UCSC Genome Browser	603-006-24			
4:30 pm – 5:00 pm	Plenary	07	Presidential Welcome & Address	603-007-24	0.50	0.50	0.50
5:00 pm – 6:40 pm	Plenary	08	Featured Plenary Abstract Session I	603-008-24	1.75	1.75	1.50
Wednesday,	November 6						
		09	A Heart to Heart on Cardiovascular Genetics in Health and Disease	603-009-24			
	1	10	Advancements in Molecular and Cytogenetic Diagnostics	603-010-24	1.50	1.50	1.50
		11	All the Single Cells	603-011-24			
8:00 am –	Dietferm	12	Beyond Genetic Discoveries: Novel Mechanisms of Neurodevelopmental Disorders	603-012-24			
9:30 am	Platform	13	Biobank Scale Genetic Data Resources for Studying Complex and Rare Human Diseases	603-013-24			
		14	Cancer Risk: Novel Genes and Mechanisms	603-014-24			
		15	Decoding Structural Variation at Scale	603-015-24			
		16	I See Ghosts: Archaic DNA in Our Genomes	603-016-24			
10:45 am -		17	Creative Community Engagement: Gathering Data for Better Participatory Research	603-017-24			
	18	Machine Learning and AI Applications in Human Genetics	603-018-24	1.50	4.50	1.50	
12:15 pm	Plattorm	19	Mapping the Brain in Health and Disease	603-019-24	1.50	1.50	1.50
	20	Moving Polygenic Risk Scores Closer to Clinical Implementation	603-020-24				

Wednesday, I	November 6						
		21	Multimodal Approaches to Interpreting the Non-Coding Genome:	603-021-24			
10:45 am –	Platform		Evolution, Functional Genomics, and Machine Learning				
		22	New Frontiers in Multi-ancestry Methods for Complex Traits	603-022-24	1.50	1.50	1.50
12:15 pm		23	Not Only Genetics: Integrating Other Omics Approaches	603-023-24			
		24	The Sex-Specific Landscape: Variation, Regulation, and Expression	603-024-24			
		25	Decoding Gene Expression Cis and Trans	603-025-24			
		26	Genetic Approaches Informing Drug Targets and Mechanism	603-026-24			1.00
		27	Interrogating Variant Function at Scale	603-027-24			
1:15 pm –		28	Liver, Laugh, Love: New Insights into Liver Disease	603-028-24			
2:15 pm	Platform	29	Modeling Rare Neurodevelopmental Disorders in Human iPSCs and Mice	603-029-24	1.00	1.00	
		30	Novel Aspects of Modeling Genetic Architectures of Complex Traits	603-030-24			
		31	Therapies for Genetic Disorders	603-031-24			
		32	Unifying Multimodalities: Insights from Single Cell Analyses	603-032-24			
5:00 pm –	Dlanami	34	Presidential Symposium: Mendelian Traits: Thinking about	603-034-24	1 50	1 50	1 50
6:30 pm	Plenary	54	Complexity in the World of "Simple" Genetics		1.50	1.50	1.50
Thursday, No	vember 7						
	Featured Symposium	35	Advances in Artificial Intelligence Tools to Improve Clinical Diagnoses	603-035-24			
			and Medical Genetics Education with an Emphasis on Diverse		_		1.50
			Datasets				
		36	AJHG at 75: Looking to the Future of Human Genetics Research	603-036-24		1.50	
		37	Balancing Open Science and Patient Privacy in the Era of Precision Medicine	603-037-24	1.50		
8:00 am –		38	Contributions of Tandem Repeats to Human Variation, Traits, and Disease	603-038-24			
9:30 am		39	Human Genetic Mosaicism: Diversity within Individuals	603-039-24			
		40	The Alzheimer's Disease Sequencing Project (ADSP): A Paradigm for	603-040-24			
			Identifying Genetically Driven Therapeutics for a Global Complex Disease				
		41	TOPMed 10-Year Anniversary: Ongoing Success and Future Directions	603-041-24			
		42	Unveiling Genetic Mysteries: RNA Editing's Breakthrough in Disease and Gene Therapy	603-042-24			
10:15 am – 11:45 am	Platform	43	All about Implementation	603-043-24	1.50		
		44	Alzheimer's Disease from Gene Discovery to Multi-omics Integration	603-044-24			
		45	Disease Insights from Omic-Wide Approaches	603-045-24		4.50	4.56
		46	Diverse Epigenetic Marks in Health, Diagnosis, and Disease	603-046-24		1.50	1.50
		47	From Variant to Function: Prediction and Understanding Variants	603-047-24			
		7,	Trom variant to runction. Frediction and Onderstanding variants	003-047-24			

Thursday, Nov	vember 7						
		48	Novel Genetic, Genomic, and Epigenetic Resources in the Era of Big Data	603-048-24			
10:15 am – 11:45 am	Platform	49	Polygenic Risk Scores: Novel Methods for Modeling Risk	603-049-24	1.50	1.50	1.50
		50	The Context of All in Which We Live: Gene by Environment	603-050-24			
			Interactions				
	Platform	51	3D Chromatin and Epigenomics	603-051-24		1.00	
		52	Computational Methods for Causal Variant Prioritization	603-052-24			1.00
		53	Dysfunction at the Powerhouse: Molecules, Models, and Organisms	603-053-24			
1:15 pm –		54	Expanding the Table: Considerations for Inclusion in Genetics and Genomics	603-054-24	1.00		
2:15 pm		55	Insights into Somatic Mosaicism and Human Diseases	603-055-24			
		56	Neurogenomic Approaches Translating Risk Variants to Disease	603-056-24			
		57	Population Genetics Methods Matter	603-057-24	1		
		58	Scaling Structural Birth Defects	603-058-24			
5:00 pm – 6:40 pm	Plenary	60	Featured Plenary Abstract Session II	603-060-24	1.75	1.75	1.50
Friday, Noven	nber 8						
	Featured Symposium	61	Aging, Clonal Hematopoiesis, and Our Health	603-061-24	1.50	1.50	1.50
		62	Cross-Examining the Rare and Common Variant Architecture of	603-062-24			
			Psychiatric Conditions, Brain Structure, and Function				
		63	Face the Facts: The Impact of Advances in Data Science on	603-063-24			
			Translational Research				
		64	For the Children: Genomics to Improve the Health of Pediatric	603-064-24			
			Patients and Their Families				
8:00 am – 9:30 am		65	How Do We Describe and Ascribe Clinical Significance to the Non-coding Genome?	603-065-24			
		66	Improving Health Equity in Genomics: Interventions and	603-066-24			
			Implementation Efforts to Address Disparities in Genetic Services Research				
		67	Model Organisms to the Rescue: Next Generation Animal Models for	603-067-24	1		
			Precise Phenotyping of Complex Diseases	300 00, 24			
		68	Not Only Transcription Intermediates: The Roles of R-loops in	603-068-24			
			Genome Stability and Brain Disease				
		69	Complex Traits and Other Omics	603-069-24			
		70	Exploring the Genetic Spectrum of Obesity	603-070-24			
	Platform	70 71	Exploring the Genetic Spectrum of Obesity  Long-Read Transcriptomes in Health and Disease	603-070-24	1.50	1.50	1.50
10:15 am – 11:45 am	Platform			+	1.50	1.50	1.50

Friday, Nove	mber 8						
10:15 am – 11:45 am	Platform	74	The Non-coding Genome: From Nucleotide to Protein	603-074-24	1.50	1.50	1.50
		75	Tick-Tock: The Aging Genome	603-075-24			
		76	Translating Genetics into Screening Programs	603-076-24			
		77	Exploring Omics: From Genomes to Microbiomes	603-077-24			
		78	Genetics of Human Brain: Regulation, Disease Risk, and Assortative Mating	603-078-24		1.00	1.00
		79	Lessons from Height	603-079-24			
1:15 pm –		80	Linking Non-coding Variation to Function via Diverse Epigenetic Mechanisms	603-080-24			
2:15 pm	Platform	81	Rare Variants and Admixture Modeling in Diverse Population	603-081-24	1.00		
·		82	Read All about It: Transcriptomic Insights from New Sequencing Technologies	603-082-24	-		
		83	Splice Splice Baby: Isoform Expression in Health and Disease	603-083-24			
		84	Strategies to Interpret Germline Variants in Cancer Predisposition Genes	603-084-24			
5:00 pm – 6:40 pm	Plenary	86	Featured Plenary Abstract Session III	603-086-24	1.75	1.75	1.50
Saturday, No	ovember 9				•		
	Platform	87	Framing Heritability for Complex Traits	603-087-24	1.00	1.00	
		88	Keeping It REnAL! Genetic Studies of Kidney Disease	603-088-24			1.00
8:00 am – 9:00 am		89	Long-Read Sequencing Offering New Insights into Neurological Disease	603-089-24			
		90	Tumor Genome Landscape Studies	603-090-24			
		91	Unraveling the Complexity of Polygenic Inheritance	603-091-24			
9:30 am – 10:30 am	Platform	92	Genetic Information in Breast Cancer Risk Assessment and Screening	603-092-24	1.00	1.00	
		93	Modeling Ataxia and Neuropathy	603-093-24			1.00
		94	More than One Way to Break a Gene - Variant Effects on RNA	603-094-24			
		95	Phenomenal PheWAS	603-095-24			
		96	Technology for Translation	603-096-24			
TOTAL CREDIT COUNT					25.25 CME	25.25 NSGC	24.50 P.A.C.E.