INTERNATIONAL GENETIC

EPIDEMIOLOGY SOCIETY

IGES 27th Annual Meeting

October 13-16, 2018 San Diego, CA, USA Wyndham San Diego Bayside

ANNUAL MEETING AGENDA

Special Programming including Educational Symposium on Translating Genetic Discoveries into Clinical and Public Health Practice

> and the Joint Symposium with ASHG/ISCB Working with Big Data in the Cloud: Research and Privacy October 16, 2018



IGES 2018 ANNUAL Meeting Agenda

Saturday, O	ctober 13	
2:00pm-7:00pm	Registration	
5:00pm-7:00pm	IGES Board of Directors Meeting	
7:00pm-8:00pm	Welcome Reception	

Sunday, October 14

7:00am-6:00pm 7:30am-8:30am	Registration Continental Breakfast		Pacific Foyer Pacific Foyer
8:30am-9:00am	Welcome and Presidential Address • Chair: Andrew Morris		Pacific D
	enic Trait Genetics I • Chairs: Andrew Morris and Guillaume Paré		Pacific D
	Presentation	Abstract	Author
9:00am-9:15am	The Role of Human Mitochondrial DNA Variants in Common Complex Phenotypes	145	Ekaterina Yonova-Doing*
9:15am-9:30am	Tissue-wise Sub-typing of Complex Trait Based on Genetics	73	Arunabha Majumdar*
9:30am-9:45am	A New Statistical Method to Detect Novel Disease Associated Genes using Publicly Available GWAS Summary Data	148	Jianjun Zhang*
9:45am-10:00am	JM-SNP: Joint Modeling of Multiple Longitudinal and Multiple Surviva Traits to Characterize the Genetic Architecture of Complex Traits	16	Myriam Brossard*
10:00am-10:30am	Refreshment Break		Pacific Foyer
Session 2: Risk A	ssessment and Personalized Medicine • Chairs: Jaya Satagopan and	Robin Be	aumont Pacific D
	Presentation	Abstract	Author
10:30am-11:00am	Keynote Speaker: What African Genomes Tell Us About the Origins of Breast Cancer	87	Olufunmilayo I. Olopade
11:00am-11:15am	Detecting and Correcting for Bias in Mendelian Randomization Analyses using Gene-by-environment Interactions	113	Wes Spiller
11:15am-11:30am	Machine Learning in Multi-omics Data to Assess Longitudinal Predictors of Glycaemic Trait Levels	57	Marika Kaakinen
11:30am-11:45am	Combined Association of a Polygenic Risk Score with 313 Genetic Variants and Established Environmental Risk Factors in Relation to Breast Cancer Risk	76	Pooja Middha
11:45am-12:00pm	Absolute Risk of Pancreatic Cancer in the U.S. General Population	61	Jihye Kim

Sunday, October 14

12:00pm-1:30pm	Lunch on your own Young Investigators' Luncheon (Ticketed Event) Committee Lunches		Coast Ballroom
	Publications Committee Working Lunch Communications Committee Working Lunch		Captains III Captains IV
Session 3: Causal	Variants, Genes and Mechanisms I • Chairs: Ching-Ti Liu and James (Cook	Pacific D
	Presentation	Abstract	Author
1:30pm-2:00pm	Keynote Presentation: Multiethnic GWAS and Fine-mapping	21	David Conti
2:00pm-2:15pm	Efficient Fine-mapping to Identify Causal Genetic Variants and Quan- tify their Contribution to Complex Phenotypes	11	Christian Benner
2:15pm-2:30pm	Imputed Gene Associations Identify Replicable <i>Trans</i> -acting and Tar- get Gene Pairs Enriched in Transcription Factor Pathways	134	Heather E. Wheeler
2:30pm-2:45pm	Sparse Canonical Correlation Analysis (sCCA) Significantly Improves Power of Cross-tissue Transcriptome-wide Association Studies (TWAS)	33	Helian Feng [^]
2:45pm-3:00pm	Meta-analysis of ~1.3M Individuals Identifies Rare Variants Associated with Blood Pressure and Implicates Causal Genes	51	Joanna M. M. Howson
3:00pm-3:15pm	Session A: Poster Highlights • Chair: Andrew Morris		Pacific D
	Poster	Abstract	Author
	Wavelet Screaming: a Novel Approach to Analyzing GWAS Data	29	William Denault
	ELECTRONIC MEDICAL RECORDS AND GENOMICS (eMERGE) - Using EMR-Linked Biorepositories to Expand Genomic Medicine Research	56	Sheethal Jose
	Hierarchical Regularized Regression for Incorporating External Infor- mation in High-Dimensional Prediction Models	133	Garrett M. Weaver
	Genetic Association of Arterial Stiffness Index with Incident Coronary Artery Disease and Congestive Heart Failure	146	Seyedeh Maryam Zekavat
	A Rare Variant Non-parametric Linkage Method for Nuclear and Extended Pedigrees with Application to Exome and Whole Genome Sequence Data	152	Linhai Zhao
3:15pm-4:45pm	Poster Session A and Refreshment Break		Pacific ABC
Session 4: Rare Va	ariants and Sequence Analysis • Chairs: Inke König and Ekaterina Yo	nova-Doin	g Pacific D
	Presentation	Abstract	Author
4:45pm-5:00pm	A Unified Method for Rare Variant Analysis of GxE Interactions	71	Elise Lim [^]
5:00pm-5:15pm	TRIO_RVEMVS: a Fast Bayesian Variable Selection Method for Trios that Identifies Individual Rare Variants	117	Michael D. Swartz
5:15pm-5:30pm	A Fast Family-based Quantitative Trait Association Test for Nuclear and Extended Pedigrees for the Analysis of Whole Genome Sequence Data	150	Zhihui Zhang
5:30pm-5:45pm	Fast Calling of Copy Number Variations from Cohort-wide High-depth Whole Genome Sequencing	39	Arthur Gilly [^]
5:45pm-6:00pm	Efficient Gene-Environment Interaction Tests for Large-scale Sequencing Studies	131	Xinyu Wang
7:00pm-7:30pm	Cocktail Reception		Loma Vista Terrace

Sunday, October 14

7:30pm-10:00pm

Banquet in honor of Newton Morton

Harborside

The IGES Banquet this year will be held in honor of Newton Morton, who sadly passed away in February, to celebrate his life and incredible contributions to genetic epidemiology. Join us as we remember him through stories, images and the impact he had in the field of genetic epidemiology. Dress is business casual.

7:00am-5:30pm	Registration		Pacific Foyer
7:30am-8:30am	Continental Breakfast		Pacific ABC
Session 5: Causal	Variants, Genes and Mechanisms II • Chairs: Sanjay Shete and Aruna	abha Maju	ımdar Pacific D
	Presentation	Abstract	Author
8:30am-8:45am	Best Paper Presentation		
8:45am-9:00am	Significance Testing for Allelic Heterogeneity	88	Wei Pan
9:00am-9:15am	Prediction of CpG Methylation Status from SNP Genotype Data	36	James J. Fryett
9:15am-9:30am	Epigenome-wide Association Study of Change in Body Mass Index from Young- to Middle Adulthood in 626 Northern Finland Birth Cohort 1966 Participants	95	Harmen Draisma
9:30am-9:45am	Integration of mQTL Data and Enhancer-promoter Interactions with GWAS Summary Results Identifies Novel Genes	141	Chong Wu
2:45am-10:00am	Session B: Poster Highlights • Chair: Andrew Morris		Pacific D
	Poster	Abstract	Author
	Genetic Architecture of Gene Expression Traits across Diverse Populations	6	Angela Andaleon
	Generalized Linear Discriminant Analysis for High-Dimensional Genomic Data	69	Sisi Li
	Pharmacogenetic Effects in Population-based Metabolic Profiles	84	Martina Müller-Nurasyic
	Next-generation Sequencing Aligned to High-resolution Nuclear Magnetic Resonance (NMR) Measurements Reveal Role of Rare Variation in Circulating Metabolic Biomarkers	101	Fernando Riveros-Mckay Aguilera
	LASSO with Custom Penalization Based on External Information	147	Chubing Zeng
0:00am-11:30am	Poster Session B and Refreshment Break		Pacific ABC
Session 6: Polyger	nic Trait Genetics II • Chairs: Eleanor Wheeler and Iris Heid		Pacific D
	Presentation	Abstract	Author
11:30am-12:00pm	Keynote Presentation: Leveraging Big GWAS Data to Address Question about Selection, Pleiotropy, Assortative Mating and Epidemiology	126	Peter M. Visscher
12:00pm-12:15pm	Population Stratification in the Estonian Biobank and its Confounding with Complex Traits	92	Natalia Pervjakova
12:15pm-12:30pm	Maternal and Fetal Genetic Effects on Birth Weight and their Relevance to Cardio-metabolic Disease	10	Robin N. Beaumont*
12:30pm-12:45pm	Exploring Genotype-environment (GxE) Interactions Phenome-wide	79	Rachel Moore [^]

Monday, October 15

1:00pm-2:30pm	Lunch on your own		
1:00pm-2:30pm	Committee Lunches		
	Program Committee Working Lunch		Captains III
	Education Committee Working Lunch		Captains IV
	Young Investigators' Committee Working Lunch		Captains V
	Genetic Epidemiology Editorial Board Working Lunch		Embarcadero
Session 7: Genom	ics in Diverse Populations • Chairs: Andrew Morris and Jianjun 2	Zhang	Pacific D
	Dresentation		A uth a r

	Presentation	Abstract	Author
2:30pm-2:45pm	Keynote Presentation: Genetic Contribution to Obesity in African Populations: the H3Africa AWI-Gen Study	97	Michèle Ramsay
3:00pm-3:15pm	Identifying Hidden Ancestries in Publicly Available Summary Data	49	Audrey E. Hendricks
3:15pm-3:30pm	Investigation of Post-colonial Demographic Structure within the United States and Implications for Association Analyses	112	Elena Sorokin
3:30pm-3:45pm	Deriving Significance Thresholds for Genome-wide Admixture Mapping Studies	43	Kelsey E. Grinde
3:45pm-4:00pm	Large-scale Trans-ethnic Genome-wide Association Study Reveals Novel Loci, Causal Molecular Mechanisms and Effector Genes for Kidney Function	35	Nora Franceschini
4:00pm-4:30pm	Refreshment Break		Pacific Foyer
4:30pm-5:30pm	Business Meeting and Awards		Pacific D

Tuesday, October 16

Special Programming and Partner Events (separate registration required)

7:00am-1:00pm	Registration	Harbor Foyer
8:30am-11:45am	Education Symposium: Translating Genetic Discoveries into Clinical and Public Health Practice (Continental breakfast provided in Harbor Foyer from 7:30am - 8:30am)	Harborside
10:00am-10:15am	Refreshment Break	Harbor Foyer

1:00pm-4:00pm IGES/ASHG/ICSB Event: Big Data in the Cloud: a Joint Symposium San Diego Convention Center, Room 6C, Upper Level



Sunday, October 14

3:15p	m-4:45pm Poste	er Sessic	on A Pacit	ic ABC			
Poster	Presenter	Poster	Presenter	Poster	Presenter	Poster	Presenter
1	Aghababazadeh, F.A.	27	de Andrade, M.	56	Jose, S.	90	Paul, S.
3	Almli, L.M.	29	Denault, W.	59	Keaton, J.M.	93	Petty, L.E.
5	Anasanti, M.D.	31	Eslami, A.	62	Konigorski, S.	96	Raisky, J.
7	Arbeev, K.G.	34	Fouodo, C.J.K.	64	Lee, Y.	99	Reeve, N.
9	Baneye, A.D.	38	Ghasemi-Dehkordi, P.	66	Lewinger, J.P.	103	Roshandel, D.
13	Bhattacharjee, S.	41	Gola, D.	68	Li, Q.	105	Sabourin, J.A.
15	Bouzigon, E.	44	Grover, S.	70	Li, Y.	107	Schaid, D.
18	Chen, HH.	46	Guyatt, A.	74	Manning, A.	109	Shriner, D.
20	Chu, S.	48	Hass, C.B.	77	Middlebrooks, C.D.	111	Smith, J.A.
23	Coombes, B.	52	Hudson, J.	80	Morrison, J.	116	Stiemke, A.B.
25	Darst, B.F.	54	Jiang, L.	82	Moss, L.C.	119	Tekola-Ayele, F.
				86	Nguyen, J.	121	Timpson, N.

Monday, October 15

Monday, October 15 10:00am-11:30am Poster Session B Pacific ABC									
Poster	Presenter	Poster	Presenter	Poster	Presenter	Poster	Presenter		
2	Allen, R.	30	Dennis, J.	63	Laabs, BH.	98	Rausch, T.K.		
4	Almorza, D.	32	Espin-Garcia, O.	67	Lewis, D.	100	Richardson, M.		
6	Andaleon, A.	37	Génin, E.	69	Li, S.	101	Riveros-Mckay Aguilera, F.		
8	Bailey-Wilson, J.E.	40	Gkatzionis, A.	72	Lipunova, N.	102	Romanescu, R.		
12	Berrandou, T.	42	Graham, B.E.	75	Mechanic, L.	104	Saad, M.		
14	Blue, E.	45	Günther, F.	78	Miller-Fleming, T.W.	106	Sarnowski, C.		
17	Chatzigeorgiou, C.	47	Han, S.	81	Moss, L.C.	108	Shrine, N.		
19	Cherlin, S.	50	Howey, R.	83	Mukhopadhyay, I.	110	Singh, A.		
22	Cook, J.	53	Jiang, G.	84	Müller-Nurasyid, M.	114	Staley, J.		
24	Daley, D.	55	Jiang, R.	85	Musolf, A.M.	118	Takeuchi, F.		
26	Das, S.	58	Karhunen, V.	89	Pathak, G.A.	120	Thiele, D.		
28	Dejenie, T.	60	Khan, S.A.	91	Pereira, M.	122	Tounkara, F.		
				94	Pileckyte, E.	124	Ullah, E.		

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25	Viel, R.	_	Poster r	emoval hou	Ire			Pacific .	ARC
28	Wan, J.Y.	-				30am - 1:30p			
0	Wang, Q.								
33	Weaver, G.			17 18	53 54	89 93	122 1	133	m
6	Willems, E.L.			15 19	52 55	90 91	-	124	F&B
8	Wilson, C.			14 20 13 22	50 56 48 58	85 96 86 94		136 127	
0	Workalemahu, T.			12 23	48 58	84 99	118 1	138	
3	Yadav, A.			9_24	46 60	82 98	125 1	129	
6	Zekavat, S.M.	VX	- B B B B B B B B B B B B B B B B B B B						
.9	Zhang, M.			8 25 7 26	45 62 44 63	83 103 80 100	114 1 121 1		
52	Zhao, L.			7 26 6 27	44 63 42 64	80 100 81 105	_	132 143 147	7 151
				5 28	41 67	77 101	-+-	135 144	1
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oster	Presenter								
7	Ullah, E.								
7	Wang, J.			ENTRAN	CE	ENT	RANCE -		
2	Wang, X.			Ma	rk Yo	ur Cal	end	ar fi	0r
5	Wijsman, E.M.								
7	Williams, A.T.			Zoui	IUE9	Annua		GGU	ШУ
9	Woodbury-Smith, M.		57						
12	Xu, H.			1)ctobe	e r 13 -1	5.2	019	
4	Yasmeen, S.								
17	Zeng, C.				0050	on, Tex	ia5, I	121	
51	Zhang, Z.				G				
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INTERNATIONAL GENETIC EPIDEMIOLOGY SOCIETY

IGES 27th Annual Meeting INVITED SPEAKERS



Dr. David Conti

Dr. Conti is Professor in the Division of Biostatistics in the Department of Preventive Medicine and the Norris Cancer Center at USC. He was trained in Genetic Epidemiology and Biostatistics at Case Western Reserve University and was awarded the 2001 Roger Williams award from the International Genetic Epidemiology Society

(IGES). His methodological research focuses on the application of hierarchical models and Bayes model averaging to genetic association studies with a specific interest in multilocus analysis and the integration of biological knowledge in statistical modeling.



Olufunmilayo I. Olopade, MD, FACP, OON

Dr. Olufunmilayo I. OlopadeDr. Olopade is the Walter L. Palmer Distinguished Service Professor of Medicine and Human Genetics, Dean for Global Health, and Director of the Center for Innovation in Global Health at the University of Chicago. Professor Olopade develops innovative strategies for comprehensive cancer risk assess-

ment and prevention based on evolving understanding of genetic and nongenetic factors in individual patients, with a focus on women of African ancestry across the Diaspora. She is an elected member of the Association of American Physicians, National Academy of Medicine, American Academy of Arts and Sciences, and the American Philosophical Society.



Michèle Ramsay, PhD

Dr. Michele RamsayDr. Ramsay is the Director of the Sydney Benner Institute for Molecular Bioscience at the University of the Witwatersrand, Johannesburg, Professor in Human Genetics and South African Research Chair holder in Genomics and Bioinformatics of African Populations. She leads the AWI-Gen study on the genetic and envi-

ronmental contributions to obesity and cardiometabolic disease risk in six centres across four African countries. Her research aims to shed light on the role of African population genomic variation in susceptibility to diseases, given the ethnolinguistic and environmental diversity across the continent.



Peter Visscher, PhD

Dr. Peter Visscher's postdoctoral in Melbourne (Australia) was followed to Edinburgh, where he developed methods to map genetic loci underlying complex traits. He moved to a faculty position at the Institute of Evolutionary Biology of the University of Edinburgh, developing gene mapping methods and software tools, with practical appli-

cations in livestock and human populations. In 2011, he moved to the University of Queensland where he is Professor of Quantitative Genetics and Director of the Program in Complex Trait Genomics. Visscher is a Senior Principal Research Fellow of the Australian National Health and Medical Research Council and was elected a Fellow of the Australian Academy of Science in 2010.

Educational Symposium Speakers



Marylyn Ritchie, PhD

Director of the Center for Translational Bioinformatics, Associate Director for Bioinformatics in the Institute for Biomedical Informatics at the University of Pennsylvania-School of Medicine, Associate Director for the Penn Center for Precision Medicine **Exploring the World of Phenomics**



Josh Peterson, MD, PhD

Director of the Center for Translational Bioinformatics, Associate Professor of Biomedical Informatics, Associate Professor of Medicine chool of Medicine at Vanderbilt University Generating evidence for pharmacogenomics through implementation and pragmatic studies



Caroline Fox, MD

Vice President and Head, Genetics and Pharmacogenomics at Merck Research Labs Using Human Genetics to Drive Drug Discovery: the Industry Perspective



Gillian Hooker, PhD, ScM, LCGC

Vice President of Clinical Development Concert Genetics *Clinical Genetic Testing in the Genome Era: A View Across the U.S. Landscape*



Jessica Ezzell Hunter, PhD

Investigator, Center for Health Research Kaiser Permanente Northwest The impact and implementation of genetics and genomics: looking towards the future