









Organizing Committee

Sarah Harris, University of Leeds, United Kingdom Stephen Levene, University of Texas at Dallas, USA Julia Salzman, Stanford University, USA Massa Shoura, Stanford University, USA

Scientific sessions will be held in the Santa Cruz Room and poster sessions will be held in the New Brighton Room, unless otherwise noted.

Sunday, August 19, 2018

15:00 – 18:00	Registration/Information	Santa Cruz Room
18:00 – 19:00	Welcome Reception	. Courtyard Terrace
19:00 – 20:00	Dinner	Sunset Restaurant

Monday, August 20, 2018

	8:30 – 17:00	Registration/Information	Santa Cruz Room
Session	n I	Junk or Not Junk: Structure and Sequence of Coding and Nonco Sarah Harris, University of Leeds, United Kingdom, Chair	ding DNA
	9:00 – 9:45	Sergei Mirkin, Tufts University, USA RNA-DNA Hybrids Promote the Expansion of Friedreich's Ataxia (GAA)n Repevia Break-induced Replication	ats
	9:45 – 10:30	David Levens, NIH, USA The Regulatory Roles of DNA Topology and Conformation in Mammalian Gen	ne Expression
	10:30 – 10:50	Coffee Break	Seacliff Lounge & Terrace
	10:50 – 11:10	Anton Goloborodko, MIT, USA* A Pathway for Mitotic Chromosome Formation	
	11:10 – 11:55	Charles Dorman, Trinity College, Ireland Bacterial Decision-making Operating Through Tuneable Binary Genetic Switch	hes
	12:00 – 14:00	Lunch	Sunset Restaurant
Session	ı II	Biophysical Approaches to Understanding Chromatin Structure Stephen Levene, University of Texas at Dallas, USA, Chair	
	14:00 – 14:45	Wilma Olson, Rutgers University, USA Contributions of DNA Sequence in 3D Genomic Architectures	
	14:45 – 15:30	Tamar Schlick, New York University, USA Modeling Gene Elements at Nucleosome Resolution	
	15:30 – 15:50	Coffee Break	Seacliff Lounge & Terrace
	15:50 – 16:35	Andrzej Stasiak, University of Lausanne, Switzerland *Transcription-induced Supercoiling and TADs Formation*	
	16:35 – 17:20	Karsten Rippe, Heidelberg University, Germany Establishing Chromatin Subcompartments That Are Both Stable and Plastic	
	17:20 – 18:00	Nature Hike	Chaminade Red & Blue Trails

Tuesday, August 21, 2018

18:00 - 20:00

8:30 – 17:00	Registration/Information San	nta Cruz Room

Session III **Exploring the Physical Genome I**

Wilma Olson, Rutgers University, USA, Chair

Javier Arsuaga, University of California, Davis, USA 9:00 - 9:45

Biophysical Models of DNA Organization inside Viral Capsids

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9:45 – 10:30	Shinichi Morishita, University of Tokyo, Japan Understanding Tandem Repeats and Methylation with Long-read Sequencing
10:30 – 10:50	Coffee Break Seacliff Lounge & Terrace
10:50 – 11:10	Thomas Bishop, Louisiana Tech University, USA* G-Dash: A Genomics Dashboard that Unites Physics and Informatics Studies of Chromatin
11:10 – 11:55	Alexandra Zidovska, New York University, USA The "Self-stirred" Genome: Bulk and Surface Dynamics of the Chromatin Globule
12:00 – 14:00	Lunch Sunset Restaurant
Session IV	Exploring the Physical Genome II Stephen Levene, University of Texas at Dallas, USA, Chair
14:00 – 14:45	Xaiver Darzacq, University of California, Berkeley Nuclear Organization and Transcription Regulation Mechanisms Studied by Live Cell Imaging
14:45 – 15:30	Martin Depken, Delft University of Technology, The Netherlands **Bottom-up Physical Modelling for CRISPR//Cas Target Prediction**
15:30 – 15:50	Coffee Break Seacliff Lounge & Terrace
15:50 – 16:35	Laura Landweber, Columbia University, USA RNA-programmed Genome Rearrangement in the Ciliate Oxytricha
16:35 – 16:55	Katerina Kraft, Stanford University, USA* Genomic Rearrangement Induced Gene Activation by Architectural Stripes
17:00 – 18:00	David Schwartz, University of Wisconsin, USA From Big DNA Molecules to Big Data Keynote Speaker
18:00 – 20:00	Dinner
Wednesday, Augu	st 22, 2018
8:30 – 17:00	Registration/Information Santa Cruz Room
Session V	Fundamental Limits of Sequencing Accuracy, Sensitivity, and Uniqueness: What Do, and What Don't, We Know? Marc Salit, NIST, USA, Chair
9:00 – 9:45	Marc Salit, NIST, USA Metrology of Genome-scale Measurements: Standards and Systematics to Get Comparability and Confidence

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9:00 – 9:45	Marc Salit, NIST, USA Metrology of Genome-scale Measurements: Standards and Systematics to Get Comparability and Confidence	
9:45 – 10:05	Miten Jain, University of California, Santa Cruz, USA* A Reference Human Transcriptome Based on Native RNA Sequencing	
10:05 – 10:25	Idan Gabdank, Stanford University, USA* Portable and Reproducible Computational Analyses	
10:25 – 10:50	Coffee Break Seacliff Loui	nge & Terrace
10:50 – 11:10	Stephen Lincoln, Invitae, USA* Complex Genetic Variants: Implications for Clinical Sequencing Methods and Validation Appr	oaches
11:10 – 12:00	Formal Discussions	
12:00 – 14:00	Lunch	et Restaurant
Session VI	Single-cell Genomics and Single-Molecule Sequencing Tim J. Stevens, MRC Laboratory of Molecular Biology, United Kingdom, Chair	
14:00 – 14:45	Tim J. Stevens, MRC Laboratory of Molecular Biology, United Kingdom Capturing the 3D Folds of Whole Mammalian Genomes in Single Cells	
14:45 – 15:30	Bo Wang, Stanford University, USA Self-assembling Manifolds in Single-Cell RNA Sequencing Data	
15:30 – 16:15	Cristian Micheletti, SISSA Trieste, Italy Nanopore Translocation of Knotted DNA	

16:15 – 18:00

Poster Session

20:00	Pool Night
Thursday, August	: 23, 2018
8:30 – 17:00	Registration/Information Santa Cruz Roor
Session VII	The RNA World: Structure, Dynamics, and Interaction Andrew Fire, Stanford University, USA, Chair
9:00 – 9:45	Andrew Fire, Stanford University, USA Long-term RNA-based Transmission of Biological States
9:45 – 10:30	Alan Lambowitz, University of Texas at Austin, USA Thermostable Group II Intron Reverse Transcriptases (TGIRTs) and Their Use in RNA-seq
10:30 – 10:50	Coffee Break Seacliff Lounge & Terrac
10:50 – 11:10	Priscilla L. Boon, National University of Singapore* How Dengue Capsid Protein Assists in Organizing Dengue Virus Genomic RNA
11:10 – 11:55	Sarah Woodson, Johns Hopkins University, USA Sequential Folding of RNA
12:00 – 14:00	LunchSunset Restaurar
Session VIII	Genomics of Gene Regulation Massa Shoura, Stanford University, USA, Chair
14:00 – 14:45	Nadav Ahituv, University of California, San Francisco, USA Functional Characterization of Gene Regulatory Elements
14:45 – 15:30	Polly Fordyce, Stanford University, USA Quantitative Mapping of Transcription Factor Binding Energy Landscapes
15:30 – 15:50	Coffee Break Seacliff Lounge & Terrac
15:50 – 16:35	Zeba Wunderlich, University of California, Irvine, USA <i>Noise in the Shadows</i>
16:35 – 16:55	Ariel Afek, Duke University, USA* Mismatched Base-pairs Locally Distort DNA Structure and Can Induce Increased DNA-binding by Transcription Factor Proteins
16:55 – 17:15	Alexander Wood, Newcastle University, United Kingdom* What Gene Expression Noise Tells about the Spatiotemporal Organization of Gene Regulatory Networks
17:15 – 18:00	Illumina Workshop
18:00 – 20:00	Dinner Sunset Restaurar
Friday, August 24	, 2018
8:30 – 12:00	Registration/Information Santa Cruz Roor
Session IX	Extreme Genomes Massa Shoura, Stanford University, USA, Chair
9:00 – 9:45	Ami Bhatt, Stanford University, USA Culture-free Microbial Genome Assembly and Tracking in Hospitalized Patients
9:45 – 10:05	Jason DeRouchey, University of Kentucky, USA* DNA in Tight Spaces: Linking Structure, Stability, and Protection in Sperm Chromatin
10:05 – 10:50	Joanna Kelley, Washington State University, USA Eukaryotic Genome Evolution in Extreme Environments
10:50 – 11:20	Closing Remarks Co-Organizers: Sarah Harris, Stephen Levene, Massa Shoura
12.00 14.00	Lunch Suncat Postauran

Dinner Sunset Restaurant

12:00 - 14:00

18:00 - 20:00

LunchSunset Restaurant



We never stop seeking

We are driven to know more—to relentlessly search for the answers that will advance the understanding of genomics to improve human health. And we realize we can't do it alone. We're counting on the next generation of scientific minds to help us keep up the momentum. As the leading developer of genomic solutions and services, we help accelerate genetic research and its use in the fields of cancer, hereditary disease, reproductive health, infectious disease, and agriculture. Together, we'll realize the promise of personalized medicine.

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