

NANOPORE COMMUNITY MEETING 2023: HOUSTON

AGENDA

The following pages will guide you through all we have in store for you at this year's event.

TECHNICAL CLINICS

One-to-one support session with our experts

FLOW CELL LOADING

Perfect your flow cell loading technique in this hands-on, expert-led demonstration



ONSITE

These events are viewable on site



ONLINE

These events are viewable online



NCM
Houston





Wednesday 6th December 2023



08:00–09:00 Pre-event activities



08:00–08:45 CST  **REGISTRATION AND BREAKFAST**

08:00–08:30 CST  **POSTER REGISTRATION**

09:00–10:35 Session 1

09:00–09:15 CST   **WELCOME TO THE NANOPORE COMMUNITY MEETING 2023: HOUSTON** **AUDITORIUM**
Gordon Sanghera | CEO, Oxford Nanopore Technologies

09:15–09:40 CST   **PLENARY 1** **AUDITORIUM**
Nanopore sequencing of cell-free DNA for methylation-based breast cancer detection
Billy Lau | Stanford University School of Medicine, USA
■ **CANCER RESEARCH**
■ **EPIGENETICS**

09:40–10:05 CST   **PLENARY 2** **AUDITORIUM**
Future diagnostic potential for long-read sequencing as a single assay for imprinting disorders
Cate Paschal | Seattle Children's Hospital & University of Washington, USA
■ **HUMAN & CLINICAL RESEARCH**
■ **EPIGENETICS**

10:05–10:35 CST



LIGHTNING TALKS

AUDITORIUM

Liquid biopsy therapy response monitoring of pediatric high-grade gliomas

Jack Wadden | University of Michigan, USA

■ CANCER RESEARCH

De novo genome assembly of *Candida parapsilosis* from a case of neonatal sepsis

Per Aspera Adastra | Baylor College of Medicine & Texas Children's Hospital, USA

■ MICROBIOLOGY & INFECTIOUS DISEASE

Enrichment strategies for recovery of Avian influenza virus from samples using MinION

Maria Chaves | Iowa State University, USA

■ MICROBIOLOGY & INFECTIOUS DISEASE

On-site nanopore sequencing reveals microbial diversity of Colombian Pacific Coast mangrove soils

Felipe Báez Aguirre | University of the Andes, Colombia

10:35–11:35

Poster networking break

10:35–11:35 CST



FLOW CELL LOADING, CUSTOMER SERVICES, TECHNICAL CLINICS, PRODUCT TABLES

**PRODUCT
DISPLAY
AREA**

10:35–11:25 CST



POSTER NETWORKING

POSTERS

11:00–11:15 CST



MINI THEATRE

Setting new standards in clinical testing with nanopore sequencing and short-read WGS

Lucy Kaplun | Variantyx, Inc, USA

■ HUMAN & CLINICAL RESEARCH

**BREAKOUT
ROOM 1**

11:35–12:35

Session 2: Breakouts

11:35–12:35 CST



BIOINFORMATICS

MethPhaser: methylation-based haplotype phasing of human genomes

Yilei Fu | Rice University, USA

■ BIOINFORMATICS

Haplotypes, isoforms, and fusions: towards a richer cancer transcriptome

Colette Felton | University of California, Santa Cruz, USA

■ BIOINFORMATICS

STRspy 2.0: unlocking the potential of long reads for forensic DNA profiles

Rupesh Kesharwani | Baylor College of Medicine HGSC & University of North Texas HSC, USA

■ BIOINFORMATICS

BREAKOUT ROOM 1

11:35–12:35 CST



INFECTIOUS DISEASE

Nanopore sequencing for enhanced antimicrobial resistance gene surveillance

Erin L. Young | Department of Health and Human Services, State of Utah, USA

■ MICROBIOLOGY & INFECTIOUS DISEASE

Metagenomic sequencing of air samples to identify human viral pathogens

Christina Newman | University of Wisconsin-Madison, USA

■ MICROBIOLOGY & INFECTIOUS DISEASE

Building and applying global pathogen surveillance in a post/peri-pandemic world

Noah C. Hull | Association of Public Health Laboratories, USA

■ MICROBIOLOGY & INFECTIOUS DISEASE

BREAKOUT ROOM 2

11:35–12:35 CST



RARE AND COMPLEX DISORDERS

Understanding normal patterns of human structural variation with nanopore sequencing

J. (Gus) Gustafson | University of Washington, USA

■ HUMAN & CLINICAL RESEARCH

Unraveling complex Mendelian diseases with nanopore sequencing

Medhat Mahmoud | Baylor College of Medicine, Human Genome Sequencing Center, USA

■ HUMAN & CLINICAL RESEARCH

Potential personalized diagnosis and treatment of autism spectrum disorder in the era of long-read sequencing

Sarah Dada | Canada's Michael Smith Genome Sciences Centre & The University of British Columbia, Canada

■ HUMAN & CLINICAL RESEARCH

**BREAKOUT
ROOM 3**

11:35–12:35 CST



CANCER RESEARCH

Analysis of melanoma evolution using nanopore long-read sequencing data

Yuelin Liu | University of Maryland & National Cancer Institute, USA

■ CANCER RESEARCH

GoT-Splice: unraveling cell-type-specific impact of splicing factor mutations

Mariela Cortés López | Weill Cornell Medicine & New York Genome Center, USA

■ CANCER RESEARCH

■ SINGLE-CELL

Nanopore sequencing reveal structural heterogeneity in canine osteosarcoma

Mikhail Kolmogorov | National Cancer Institute, NIH, USA

■ CANCER RESEARCH

**BREAKOUT
ROOM 4**

12:35–13:45

Lunch

12:35–13:45 CST



FLOW CELL LOADING, CUSTOMER SERVICES, TECHNICAL CLINICS, PRODUCT TABLES












**PRODUCT
DISPLAY
AREA**

12:35–13:15 CST



NETWORKING LUNCH

TBC

13:10–13:25 CST	 MINI THEATRE AmplideX PCR and nanopore sequencing for accessible carrier screening for any lab Bradley Hall Asuragen, USA ■ HUMAN & CLINICAL RESEARCH	BREAKOUT ROOM 1
13:45–15:35 Session 3		
13:45–14:10 CST	  PLENARY 3 Update from the Applications team Sissel Juul VP, Emerging and Commercial Applications, Oxford Nanopore Technologies	AUDITORIUM
14:10–14:35 CST	  PLENARY 4 Telomere dynamics in aging and cancer by nanopore long-read sequencing Tobias T. Schmidt Salk Institute for Biological Studies, USA ■ CANCER RESEARCH	AUDITORIUM
14:35–15:00 CST	  PLENARY 5 2043: a MinION space odyssey Sarah Castro-Wallace NASA, USA ■ MICROBIOLOGY & INFECTIOUS DISEASE	AUDITORIUM
15:00–15:25 CST	  PLENARY 6 Improving bacterial disease public health testing with nanopore sequencing Kimberlee Musser Wadsworth Center, New York State Department of Health, USA ■ MICROBIOLOGY & INFECTIOUS DISEASE	AUDITORIUM
15:35–16:20 Coffee break		
15:35–16:20 CST	 FLOW CELL LOADING, CUSTOMER SERVICES, TECHNICAL CLINICS, PRODUCT TABLES	PRODUCT DISPLAY AREA
15:50–16:10 CST	 PRODUCT DEMO	PRODUCT DISPLAY AREA

16:20–17:50

Session 4

16:20–16:45 CST



PLENARY 7

Complex phased variants in inherited retinal diseases with long-read sequencing

Debarshi Mustafi | University of Washington, USA

■ **HUMAN & CLINICAL RESEARCH**

AUDITORIUM

16:45–17:45 CST



UPDATE FROM OXFORD NANOPORE TECHNOLOGIES

Update from Oxford Nanopore Technologies

Rosemary Sinclair Dokos | SVP, Product & Programme Management, Oxford Nanopore Technologies

AUDITORIUM

17:45–17:50 CST



CLOSING REMARKS

AUDITORIUM

17:50–19:30

Evening networking event