

# 12th CNAG SYMPOSIUM ON GENOME RESEARCH:

## State-of-the-art genomic technologies

**12th  
December  
2023**



### Symposium

09:00 - 16:25 H

### Visits

16:25 - 17:35 H



### Location

Auditori Antoni Caparrós  
Parc Científic de Barcelona,  
Tower D  
Baldíri Reixac, 4  
Barcelona, SPAIN



### Free registration

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09:00 H	REGISTRATION
09:30 H	<b>Welcome to the 12th CNAG Symposium on Genome Research</b>  Ivo Gut, Centro Nacional de Análisis Genómico (CNAG)
09:45 H	<b>Human cell atlases in health and disease</b>  Holger Heyn, Centro Nacional de Análisis Genómico (CNAG)
10:10 H	<b>Decoding oncogenic processes in childhood cancer</b>  Alexandra Avgustinova, Institut de Recerca Joan de Déu (IRSJD)
10:35 H	<b>The Xenium platform - initial experiences from a core lab</b>  Susan Kloet, Leiden University Medical Center (LUMC)
10:50 H	<b>In-situ spatial transcriptomics as a tool for next-generation pathology</b>  Anna Pascual, Centro Nacional de Análisis Genómico (CNAG)
COFFEE BREAK	
11:15 H	<b>How do I choose the right spatial biology technology?</b>  Bernat Coll-Martínez, Diagnostica Longwood
11:45 H	<b>Dynamic changes of tumor and microenvironment in colorectal cancer. From bench to bedside and bench again</b>  Iosune Baraibar, Vall d'Hebron Instituto de Oncología (VHIO)
12:00 H	<b>Recent advances in Illumina technologies and their applications</b>  Pietro Gatti, Illumina
12:25 H	<b>Structure determination of genomes and genomic domains by image tracing</b>  Marc A. Martí-Renom, Centro Nacional de Análisis Genómico (CNAG)
LUNCH	
13:05 H	<b>Applying single-cell RNA sequencing and spatial transcriptomics to understand inflammatory bowel disease</b>  Azucena Salas, Inflammatory Bowel Disease group (FRCB-IDIBAPS)
14:05 H	<b>Long-Read sequencing: advances and bottlenecks</b>  Marta Gut, Centro Nacional de Análisis Genómico (CNAG)
14:30 H	<b>Diagnosing mitochondrial diseases with novel genomic technologies</b>  Rita Horvath, Department of Clinical Neurosciences, University of Cambridge
14:55 H	<b>Going beyond - why what you're missing matters and how to get there</b>  Daniel Fordham, Oxford Nanopore Technologies
15:20 H	<b>The pearly razorfish as model species for behavioural and population genomics</b>  Margarida Barceló, Institut Mediterrani d'Estudis Avançats, IMEDEA (UIB-CSIC)
15:35 H	<b>Genomics of sex chromosome evolution in cichlid fish</b>  Astrid Böhne, Museum Koenig Bonn
16:00 H	CLOSURE AND VISIT TO CNAG INSTALLATIONS
16:25 H	