

27-29 October 2025

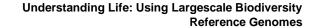
# Hybrid Conference Programme

Start	Finish	Presenter details
(BST)	(BST/)	

15:00

15:30 Refreshment break and networking

Monday 27 October 2025		
12:00	13:00	Registration, lunch and networking
12:45		Briefing for Session 1 speakers, microphone runners, chair, moderator & committee - Auditorium
13:00	13:15	Welcome and introductions
		Wellcome Connecting Science:
		Nagehan Ramazanoglu Bahadir, Wellcome Connecting Science, UK
		Scientific programme committee:
		Mark Blaxter, Welcome Sanger Institute, UK Molly Carter, Welcome Sanger Institute, UK
		Kerstin Howe, Welcome Sanger Institute, UK
		Peter Mulhair, University of Liverpool, UK
13:15	14:45	Session 1: Genome Evolution
		Chair: Peter Mulhair, University of Liverpool, UK (possibly swap roles) Moderator: Alex Twyford - TBC
13:15	13:45	Rediploidisation: unlocking the evolutionary potential of whole genome duplication <u>Aoife McLysaght, Trinity College Dublin, Ireland</u>
13:45	14:00	Biodiversity and evolution of L chromosomes using germ-line sequencing data from single wild-caught fungus gnat males Aleksandra Bliznina, Wellcome Sanger Institute, UK
14:00	14:15	Decoding the avian missing gene mystery: dot chromosomes unmask extensive gene loss and novel genetic instability  Tomas Hron, Institute of Molecular Genetics, Academy of Sciences of the Czech Republic, Czech Republic
14:15	14:30	The evolution of programmed DNA elimination in free-living rhabditine nematodes  Lewis Stevens, Wellcome Sanger Institsute, UK
14:30	14:45	The Crocus panrepeatome: dynamics, dysploidy, and diversification  Nomar Waminal, Leibniz Institute Plant Genetics And Crop Plant Research, Germany
14:45	15:00	The evolution of high-order genome architecture revealed from 1,000 species  Kai Ye, Xi'an Jiaotong University, China



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15:30 17:30 Workshop: Al and annotation

Location: Sanger Institute, Room C302

Workshop Leads:

Francoise Thibaud-Nissen, National Center for Biotechnology Information (NCBI), USA

Katharina Hoff, University of Greifswald, Germany

15:30 17:30 Workshop: Sequencing the holobiont -- how to disentangle species

Location: Sanger Institute, Room F201

Workshop Lead:

Ellen Cameron, Earlham Institute, UK

15:30 17:30 Workshop: Curation of assemblies

Location: Sanger Institute, Room C303

Workshop Lead:

Jo Wood, Wellcome Sanger Institute, UK

15:30 17:30 Workshop: Advances in Genomic Technologies

Location: Sanger Institute, Room G301 Workshop Lead:

Ana Conesa, CSIC Valencia, Spain

15:30 17:30 Workshop: Project Psyche Showcase

Location: Sanger Institute, Room G201

Workshop Leads:

Joana Meier, Wellcome Sanger Institute, UK Charlotte Wright, Wellcome Sanger Institute, UK

17:30 18:00 Poster pitch talks for poster session 1 presenters

18:00 19:30 Poster session 1

19:30 21:00 Dinner

19:30 Bar open (card payments only)



Tuesday 28 October 2025

13:45

27-29 October 2025

09:15		Briefing for Session 3 speakers, chair & moderator - Auditorium
09:30	10:30	Keynote Lecture 1
		Chair: Mark Blaxter, Welcome Sanger Institute, UK Moderator: Molly Carter, Welcome Sanger Institute, UK
		Cell type diversity and regulation in early metazoans <u>Amau Sebe Pedros, Centre for Genomic Regulation, Spain</u>
10:30	11:00	Refreshment break and networking
10:45		Briefing for session 2 speakers, chair & moderator - Auditorium
11:00	12:30	Session 2: Symbiosis Genomics
		Chair: Graeme Oatley, Wellcome Sanger Institute, UK Moderator: Elizabeth SInclair, Wellcome Sanger Institute, UK
11:00	11:30	Sample Collection, Genome Curation and How the Data is Used – A Quick Look at the Coral Hub of the Aquatic Symbiosis Genome Project
		Mike Sweet, University of Derby, UK
11:30	11:45	Mike Sweet, University of Derby, UK  What Darwin Tree of Life Genomes Have Taught Us About Microsporidia: From The Origin of Tetraploidy to Fossils in Host Genomes Amjad Khalaf, Wellcome Sanger Institute, UK
11:30 11:45	11:45 12:00	What Darwin Tree of Life Genomes Have Taught Us About Microsporidia: From The Origin of Tetraploidy to Fossils in Host Genomes
		What Darwin Tree of Life Genomes Have Taught Us About Microsporidia: From The Origin of Tetraploidy to Fossils in Host Genomes  Amjad Khalaf, Wellcome Sanger Institute, UK  The giant barrel sponge Xestospongia muta whole genome sequence has implications for reef ecology
11:45	12:00	What Darwin Tree of Life Genomes Have Taught Us About Microsporidia: From The Origin of Tetraploidy to Fossils in Host Genomes  Amjad Khalaf, Wellcome Sanger Institute, UK  The giant barrel sponge Xestospongia muta whole genome sequence has implications for reef ecology  Jose Lopez, Nova Southeastern University, USA  Disentangling Lichen Holobionts

13:45	15:15	Session 3: Genomics of Speciation
		Chair: Kerstin Howe, Welcome Sanger Institute, UK Moderator: Peter Mulhair, University of Liverpool, UK
13:45	14:15	Structural genetic diversity across the Tree of Life: how can we access evolutionary patterns and processes? Claire Mérot, University of Rennes, France
14:15	14:30	Advancing Evolutionary and Conservation Understanding of Sloth Lineages Camila Mazzoni, Leibniz Institute for Zoo and Wildlife Research, Germany
14:30	14:45	Pangenomics of the black soldier fly (Hermetia illucens) Callum Thomas, University of Cambridge, UK
14:45	15:00	Exploring genome structure evolution in the rapid radiation of large white-headed gulls  Elisa Ramos, Wellcome Sanger Institute, UK
15:00	15:15	Diving into benthic genomes: optimizing sequencing and assembly workflows for a subgroup of colonial ascidians Simona Buonanno, University of Bari, Italy
15:15	15:45	Refreshment break and networking

Briefing for session 3 speakers, chair & moderator - Auditorium

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#### 15:45 17:45 Workshop: Bat1K showcase

Location: Sanger Institute, Room E201

Workshop Leads:

Michael Hiller, LOEWE Centre for Translational Biodiversity Genomics, Germany

Emma Teeling, University College Dublin, Ireland Sonja Vernes, University of St Andrews

## 15:45 17:45 Workshop: Aquatic symbiosis genomics

Location: Sanger Institute, Room G301

Workshop Leads:

Jose Lopez, Nova Southeastern University, USA

### 15:45 17:45 Workshop: Let's be FAIR: using pipelines in genomics research

Location: Sanger Institute, Room C303

Workshop Leads:

Chris Wyatt, University College London, UK

## 15:45 17:45 Workshop: Population genomics and conservation

Location: Sanger Institute, Room C302

Workshop Leads:

Ilik Saccheri, University of Liverpool, UK Niklas Wahlberg, Lund University, Sweden

# 17:45 18:15 Poster pitch talks for poster session 2 presenters

#### Poster session 2 18:15 19:30

Performace of Eyenigma Variations: Fantasia on Codes, string quartet inspired by Ch. 6 Origin of Species by a string quartet led by Paul 19:30 19:45

Warburton.

19:45 21:00 Dinner

19:45 Bar open (card payments only)



27-29 October 2025

Wednesday 29 October 2025		
09:15		Briefing for Session 4 speakers, chair & moderator - Auditorium
09:30	11:00	Session 4: Ecological Genomics
		Chair: Mark Blaxter, Welcome Sanger Institute, UK Moderator: Molly Carter, Welcome Sanger Institute, UK
09:30	10:00	Understanding diversification in the genomic age: Noctuidae moths as an example Niklas Wahlberg, Lund University, Sweden
10:00	10:15	The genetic diversity and population genomics of sharks, rays, and other vertebrates  Amanda Gardiner, University of Cambridge, UK
10:15	10:30	Genomic adaptations to a fossorial lifestyle: Insights from the limbless Layard's snake skink (Nessia layardi) genome Ashwini Venkatanarayana Mohan, Switzerland
10:30	10:45	Evolution of the Chemical Defensome in Marine Mammals  Bram Danneels, University Of Bergen, Norway
10:45	11:00	Identifying genes under positive selection in saline tolerant tilapia species Samuel Speak, Earlham Institute, UK
11:00	11:30	Refreshment break and networking
11:15		Briefing for Keynote, chair, moderator & committee - Auditorium
11:30	12:30	Keynote 2
		Chair: Kerstin Howe, Welcome Sanger Institute, UK Moderator: Jim Downie, Wellcome Sanger Institute, UK
		Deciphering the genomic basis of sponge symbioses
		Ute Hentschel Humeida, GEOMAR Helmholtz Centre for Ocean Research, Germany
12:30	12:45	Closing Remarks
		Scientific Programme Committee:  Mark Blaxter, Welcome Sanger Institute, UK  Molly Carter, Welcome Sanger Institute, UK  Kerstin Howe, Welcome Sanger Institute, UK  Peter Mulhair, University of Liverpool, UK
12:45	13:45	Lunch and departures
13:45		Coach departures for Stansted and Heathrow airports
13:55		Coach departures for Cambridge train station and city centre