

ESHG 2022

SEQUENCE WITHOUT COMPROMISE

Register to receive copies of our content presented at ESHG

First name

Last name

Email

Organization

Country

Area of focus

Applications of interest

- ☐ All
- ☐ Epigenetics
- ☐ Gene editing
- ☐ RNA sequencing
- ☐ Metagenomics
- ☐ Pan-pathogen surveillance
- ☐ Targeted sequencing
- ☐ Viral
- ☐ WGS - De novo assembly
- ☐ Variant detection
- ☐ Other

I would like to:

- ☐ Meet a PacBio specialist
- ☐ Receive content post ESHG

Submit

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📅 11-14 June 2022

Meet the PacBio team at the ESHG conference in Vienna to learn how PacBio sequencing enables researchers to see more and do more.

At PacBio we create the world's most advanced sequencing systems to provide researchers the most complete and accurate view of genomes, transcriptomes, and epigenomes.

Highly accurate long-read sequencing, known as [HiFi sequencing](#), provides a higher resolution approach compared to previously used technologies to better understand the genetic causes of rare disease.

Below you'll discover the many ways to engage with us at the conference. We look forward to meeting you.

Highlights of the event include:

- Corporate Satellite
- In-booth presentations
- Meet our sequencing experts

Not able to attend ESHG 2022? Then make sure to fill out the form to receive copies of our posters, and other materials we'll present at the conference.

CORPORATE SATELLITE

PacBio sequencing: See what you've been missing

📅 Sunday, June 12, 2022 14:15–15:45

📍 Room Room M 1+2, Level 1

SPEAKERS



PacBio sequencing and its applications

Neil Ward, GM and Vice President EMEA, PacBio



Closing the gap: Solving complex medically relevant dark regions of the human genome

Fritz Sedlazeck, Dr. rer. nat. and Associate Professor, Baylor College of Medicine, Houston, USA



Accurate long reads for rare disease research

Alexander Hoischen, PhD and Associate Professor, Radboudumc, Nijmegen, The Netherlands



Single-cell analysis of alternative splicing in hematopoietic stem and progenitor Cells

Laura Mincarelli, PhD, Beatson Institute for Cancer Research, UK



Detecting the full spectrum of genomic variation with HiFi long reads

Michael Eberle, PhD, VP Computational Biology, PacBio

PRESENTATIONS + POSTERS

POSTER PRESENTATIONS	+
IN-BOOTH PRESENTATIONS	+

VISIT US AT BOOTH #X5-364

Stop by our booth to chat with PacBio staff and have your questions answered.

We are excited to learn about your sequencing goals and learn more about your research.

Booth hours:

- Sunday 9 June: 9:30–18:30
- Monday 10 June: 9:30–17:15
- Tuesday 11 June: 9:30–17:15

Get in touch

Speak to a sequencing specialist

Contact us

Have a collaboration you'd like to talk about

Collaborate with us

PacBio - 1305 O'Brien Dr., Menlo Park, CA 94025
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