

19 September-Saturday

9.00-10.30 SESSION 11

Viral Diagnostics: Innovations and Challenges

Chairs: Adrienne Bambach, Mario Hönemann

9.00-9.30

Keynote Lecture - Sustainable laboratory Practices: Practical Strategies, Challenges, Innovations and Future Directions

Padraig Williams, Ireland

9.30-10.30

Oral presentations (OP41 - OP44)

OP41 | Improved Methods to Diagnose iciHHV-6

Alexander Lennon, United Kingdom

OP42 | Next-Generation Sequencing of Human Cytomegalovirus for the Identification of Resistance Mutations

Hope Johnston, United Kingdom

OP43 | When fever misleads atypical measles in a patient with high-risk Lassa virus exposure

Pauline Vetter, Switzerland

OP44 | Masking Recurrent Contaminants in Reference Sequences Improves Specificity of Clinical Metagenomic Sequencing

Ian Pichler, Switzerland

10.30-11.00

Coffee break

11.00-12.00

ENPEN & ENNGS Joint Session: Bridging Frontiers

CNS infections with enterovirus and parechovirus- pilot hospital based surveillance on CNS infection

Natasa Berginc, Slovenia

Pathoplexus and Nextclade : Sharing and analyzing EV genomic data

Alejandra Gonzalez Sanchez, Spain

EVOLVE : A modern, collaborative repository powering research into viral diversity, evolution and epidemiology through high-quality sequencing and standardized metadata

Cristina Andres Verges, Spain

Clinical Metagenomics in Real Life

Julianne Brown , United Kingdom

Quality control in Metagenomics: Metagenome in a bottle

F. Xavier López-Labrador, Spain

Quality control in Metagenomics: Multi-Laboratory Evaluation of NGS for HIV resistance testing

Hope Johnston, United Kingdom

12.00-12.30

Best Poster Session

12.30-13.00

Closing - Hand Over ESCV 2027