



Autumn Meeting MolecularDiagnostics.be

November 28th 2019, 9h00-12h30

BIP ('huis van het gewest')

Koningsstraat 2-4 - 1000 Brussel

Please register before November 21st on www.moleculardiagnostics.be, after logging in.

NGS variant classification and interpretation software in Hemato-oncology

Program

09h00	Welcome and announcements
09h05	OncoKDM (OncoDNA)
09h25	SOPHiA DDM (Sophia Genetics)
09h45	QCI Interpret (Qiagen)
10h05	PierianDX (PierianDX for Illumina)
10h25	Coffee
11h00	Oncomine Reporter (ThermoFischer Scientific)
11h20	ALISSA & MASTR reporter (Agilent Technologies)
11h40	NAVIFY (Roche)
12h00	To be confirmed
12h20	Lunch



Next-generation Sequencing in the routine microbiological diagnostic laboratory

Program

- 13h30** Welcome and announcements
- 13h35** Understanding and overcoming the pitfalls and biases of next-generation sequencing (NGS) methods for use in the routine clinical microbiological diagnostic laboratory – Dr. John P. Hays (Erasmus MC, the Netherlands)
- 14h05** Explore the pore: Real time, long read nanopore sequencing for clinical research – Dr. Leila Luheshi (Oxford Nanopore Technologies Ltd, United Kingdom)
- 14h25** Culture-independent targeted next generation sequencing of the 16S-23S rRNA region for the identification of bacterial species directly from clinical samples: opportunities and challenges - Dr. Mirjam Kooistra-Smid (Certe, Groningen, the Netherlands)
- 14h45** Coffee
- 15h15** The nasal microbiota: comparison of Illumina and Nanopore 16S rRNA gene sequencing – Dr. Astrid Heikema (Erasmus MC, the Netherlands)
- 15h45** Microbiome: From sample to clinical insight: Unravel all the 16S V-regions with 1 kit – PhD Hélène Bauby (Qiagen, France)
- 17h05** Reliable species identification with SmartGene 16S Centroids - Stefan Emler (Smartgene, Switzerland)
- 17h25** Closing remarks & goodbye drinks