

Constitution Day celebration, 17th of May (Photo by Petter Hebæk, cropped)

Target Audience

Technicians, (molecular) biologists, medical (molecular) microbiologists, infectious diseases specialists, public health specialists and bioinformaticians who would like an introduction and an update on how to use next-generation sequencing in routine diagnostics of infectious diseases.

Faculty Members

Hege Vangstein Aamot, Lørenskog, Norway Ola Brynhildsrud, Oslo, Norway Patricia Campbell, Lørenskog, Norway João André Carrico, Lisbon, Portugal Natacha Couto, Groningen, Netherlands Adrian Egli, Basel, Switzerland Nicole Fischer, Hamburg, Germany Stefan Green, Chicago, IL, United States Gilbert Greub, Lausanne, Switzerland André Ingebretsen, Oslo, Norway Mirjam Kooistra-Smid, Groningen, Netherlands Sonja Lagström, Lørenskog, Norway Aitana Lebrand, Geneva, Switzerland Jacob Moran-Gilad, Beer Sheva, Israel Paula Mölling, Örebro, Sweden Justin O'Grady, London, UK Helena Seth-Smith, Basel, Switzerland Silke Peter, Tuebingen, Germany John W.A. Rossen, Groningen, Netherlands Henrik Westh, Hvidovre, Denmark

Contact

For the Scientific Programme

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ESCMID Postgraduate Technical Workshop

Next-generation sequencing in routine clinical microbiology and infectious diseases

Lørenskog, Norway 13 – 16 May 2020



Photo by Anniken Sandvik

ESCMID Postgraduate Technical Workshop

Next-generation sequencing in routine clinical microbiology and infectious diseases

Organisers

ESCMID Study Group for Genomic and Molecular Diagnostics (ESGMD)

ESCMID Study Group for Epidemiological Markers (ESGEM)
Akershus University Hospital, Norway
University Medical Centre Groningen (UMCG), Netherlands

Course Coordinators

Hege Vangstein Aamot, Lørenskog, Norway John W.A. Rossen, Groningen, Netherlands

Course Objectives

With a multitude of benefits, Next-Generation Sequencing (NGS) technology is revolutionizing microbiological diagnostics and NGS is on the way on becoming the new gold standard. The workshop will highlight opportunities and challenges with this technology and present how it can be used in clinical diagnostic settings. As bioinformatics is a known challenge for most non-bioinformaticians, a practical e-lab will be included where we focus on data processing, analysis and interpretation of data.

Course Programme

Wednesday 13 May 2020

08:00	Registration
09:00	Welcome to Norway Merete Holth
09:10	Welcome to the course
00.10	John W.A. Rossen
Session 1: Ir	ntroduction to NGS
Chairs:	John W.A. Rossen, Hege Vangstein Aamot
09:15	NGS: the basics
	Stefan Green
10:15	Sequencing technologies and platforms Aitana Lebrand
11:15	Coffee break
11:45	Sample preparation and sequencing
12:30	Stefan Green Bioinformatics for non-bioinformaticians
	Ola Brynhildsrud
13:15	Lunch
14:15	QC of the bioinformatic pipelines
	Jacob Moran-Gilad
15:00	Analyses and visualization of metagenomic data João André Carriço
15:45	Coffee break
16:15	Added value of long read sequencing
	Justin O'Grady
17:15	Welcome reception (drinks and bites)
Thursday 4.4	Nov. 2020

Thursday 14 May 2020

Session 2: Clinical applications - Part 1

Chairs: Stefan Green and Natacha Couto

NGS from a clinical view

08:30 Challenges with current microbiological diagnostics and opportunities with NGS to provide some insight on patients prognosis by looking at the virulome Gilbert Greub

09:30 Translating NGS to the clinic: challenges and opportunities Henrik Westh 10:30 Coffee break

Amplicon sequencing

11:00 Amplicon sequencing of human papillomavirus – possible markers for cervical cancer Sonja Lagström 11:45 Added clinical value of 16S–23S amplicon-based sequencing in a clinical setting Mirjam Kooistra-Smid 12:30 Lunch

Whole genome sequencing

18:00 Social event and dinner

13:30	The use of WGS in a hospital setting André Ingebretsen
14:15	The role of whole genome sequencing in anti- microbial susceptibility testing of bacteria Silke Peter
15:00	Coffee break
15:30 16:30	NGS for outbreak detection — there is more than the core genome! Helena Seth-Smith

Friday 15 May 2020

Session 3: Clinical applications – Part 2

Chairs: Patricia Campbell, Henrik Westh

Metagenomic sequencing

09:00	Shotgun metagenomic sequencing of infected tiss
	Hege Vangstein Aamot
09:45	When dealing with difficult to culture
	microbes – identification of Neisseria gonorr-
	hoeae in throat samples
	Patricia Campbell
10:30	Coffee Break

11:00	Application of metagenomics for bacteremia
	Paula Mölling

11:45 Metagenomics for typing Natacha Couto

12:30 Lunch

13:30 The pros and cons of metagenomic sequencing for infectious disease diagnosis – is the juice worth a squeeze? Nicole Fischer

14:15 Machine learning in microbiology diagnostics Adrian Eqli

15:00 Coffee Break

15:30 Integrative and machine learning metagenomic approaches John W.A. Rossen

16:15 Farewell to participants not attending the practicals

Saturday 16 May 2020

Session 3: Bioinformatics - e-lab

Trainers: Natacha Couto, Hege Vangstein Aamot, John W. A. Rossen, Sonja Lagström, Helena Seth-Smith

08:30 Practicals on:

- 16:00 • Quality control

· Assembly and annotation

Pipelines

Visualization tools

Virulence and resistance markers

Organisation

Course Venue

Akershus University Hospital Sukehusveien 25 1478 Lørenskog, Norway

Registration Procedure

Register on the ESCMID website at www.escmid.org/education. Registration closes on 1 April 2020.

Registration Fee

EUR 400 for ESCMID members EUR 500 for all others

The fee includes the course, coffee, lunches, welcome reception and the social dinner. Travel and accommodation are not included. Attendees participating in the bioinformatics practicals will be charged an extra fee of 100 EUR (members) or 150 EUR (non-members).

Attendance Grants

ESCMID provides a number of attendance grants for ESCMID "young scientist members". The grant covers the registration fee, but not travel or accommodation costs. Please apply via the registration website by 1 March 2020. Applicants will be informed about their acceptance by 15 March 2020.

CME Accreditation

The organiser of the course will apply for European CME accreditation through EACCME.

Recommended Accommodation

Thon Hotel Arena Nesgata 1, 2004 Lillestrøm, Norway https://www.thonhotels.com/event/escmid-workshop/

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