



Clinical Genomics and NGS

**Bertinoro - Italy
April 28 – May 3, 2019**

32nd Course jointly organized by ESGM, ESHG AND CEUB

University Residential Centre
Via Frangipane, 6 – Bertinoro

Course Directors:

Han Brunner (Nijmegen and Maastricht, the Netherlands); Christian Gilissen (Nijmegen, the Netherlands); Alexander Hoischen (Nijmegen, the Netherlands); Tommaso Pippucci (Bologna, Italy); Brunhilde Wirth (Cologne, Germany)



ALMA MATER STUDIORUM
UNIVERSITÀ DI BOLOGNA



COURSE PROGRAM

SATURDAY APRIL 27TH

Arrival and dinner

SUNDAY APRIL 28TH

Morning Lectures:

Medical Genetics concepts and principles

8:30 – 9:00

Participants Registration

9:00 – 9:15

Introduction to the course – **H. Brunner** – with video of **G. Romeo**

9:15 – 10:00

Genomic Medicine – **D. Donnai**

10:00 – 10:45

Phenotype to genotype – **H. Brunner**

10:45 – 11:00

Coffee break

11:00 – 11:45

Cytogenetics and arrays – **M. Spielman**

11:45 – 12:30

Complex disorders and classical gene identification – **A. Read**

12:30 – 13:15

Discussion of the morning lectures

13:30 – 14:30

Lunch Break

Afternoon Workshops

Session I (14:30 – 16:00)

Mutation patterns – **H. Brunner**

Interpreting CNVs for beginners – **M. Spielman**

Computer room

Dysmorphology – **D. Donnai**

16:00 – 16:30

Coffee break

Session II (16:30 – 18:00)

Mutation patterns – **H. Brunner**

Interpreting CNVs for beginners – **M. Spielman**

Computer room

Dysmorphology – **D. Donnai**

MONDAY APRIL 29TH

Morning Lectures:

Basics of NGS for Mendelian disorders

09:00 – 09:45

Basics of next generation sequencing technology – **A. Hoischen**

09:45 – 10:30

Basics of NGS bioinformatics – **C. Gilissen**

10:30 – 11:00

Coffee break

11:00 – 11:45

NGS in the clinic – **H. Brunner**

11:45 – 12:30

Long-read sequencing – **E. E. Eichler**

12:30 – 13:15

Discussion of the morning lectures

13:30 – 14:30

Lunch Break

Afternoon Workshops

Session I (14:30 – 16:00) NGS Bioinformatics Basics – **C. Gilissen & T. Pippucci**
Computer room

Targeted NGS approaches – **A. Hoischen**

Clinical Considerations for NGS – **H. Brunner**

16:00 – 16:30 Coffee break

Session II (16:30 – 18:00) NGS Bioinformatics Basics - **C. Gilissen & T. Pippucci**
Computer room

Rarity in the clinic – **J. Burn**

Copy number variations – **E. E. Eichler**

TUESDAY APRIL 30TH

Morning Lectures: **Therapy and novel disease mechanisms in the NGS era**

09:00 – 09:45 Therapy and cancer – **J. Burn**

09:45 – 10:30 SMA: From gene and modifier to therapy – **B. Wirth**

10:30 – 11:00 Coffee break

11:00 – 11:45 Molecular inversion probes and mosaicism – **A. Hoischen**

11:45 – 12:30 Epigenetics, imprinting, clinical – **K. Temple**

12:30 – 13:15 Discussion of the morning lectures

13:30 – 14:30 Lunch Break

Afternoon Workshops

Session I (14:30 – 16:00) NGS Bioinformatics, variant interpretation – **C. Gilissen & T. Pippucci**
Computer room

Genetic Imprinting – **K. Temple**

New workshop tbd – **David Fitzpatrick**

16:00 – 16:30 Coffee break

Session II (16:30 – 18:00) NGS Bioinformatics, variant interpretation – **C. Gilissen & T. Pippucci**
 Computer room

Ethics of medical genetics – **A. Read**

From your newly discovered candidate gene to its function – **B. Wirth**

18:00 Poster viewing session with aperitif (session I)

WEDNESDAY MAY 1ST

Morning Lectures: Prenatal testing in the NGS era and complex mechanisms of disease

09:00 – 09:45 Non-invasive prenatal testing – **J. Weiss**
 09:45 – 10:30 Single cell sequencing and applications to PGD – **T. Voet**
 10:30 – 11:00 Coffee break
 11:00 – 11:45 Non-coding mutations/long-range effects – **M. Spielman**
 11:45 – 12:30 Massively parallel functional assays – **L. Starita**

12:30 – 13:15 Discussion of the morning lectures

13:30 – 14:30 Lunch Break

Afternoon Excursion

THURSDAY MAY 2ND

Morning Lectures: Novel NGS applications

09:00 – 09:45 Discovering structural variants in cancer using NGS data - **T. Rausch**
 09:45 – 10:30 Phenotype and NGS integration/HPO benefits – **D. Fitzpatrick**
 10:30 – 11:00 Coffee break
 11:00 – 11:45 GWAS with NGS - **M. Nothnagel**
 11:45 – 12:30 Patterns of rare variation contributing to disease – **tbd**

12:30 – 13:15 Discussion of the morning lectures

13:30 – 14:30 Lunch Break

Afternoon Workshops

Session I (14:30 – 16:00) Discovering structural variants in cancer using NGS data, par. 1 – **T. Rausch**
Computer room

Multiplexed functional assays/variant interpretations – **L. Starita**

How to do single cell genomics? – **T. Voet**

Practical considerations for NIPT – **J. Weiss**

16:00 – 16:30 Coffee break

Session II (16:30 – 18:00) Discovering structural variants in cancer using NGS data, par. 2 – **T. Rausch**
Computer room

How to do GWAS – **M. Nothnagel**

Interpretation of rare variants – **tbd**

18:00 Poster viewing session with aperitif (session II)

Social dinner and farewell party

FRIDAY MAY 3RD

Morning Lectures:

Highlights of the 2019 course

09:30 – 10:30 New McKusick-Romeo Lecture: INVITED SPEAKER NN

10:30 – 11:30 Quiz on: “What have you learned”?

11:30 – 12:00 Best poster pitches by students

12:00 – 13:00 Lunch Break

Departure after lunch