



4th Training Course
***“Quality assurance, variant
interpretation and data
management in the NGS
diagnostics era”***

October 19-21, 2022
ONLINE



This Training has received funding from the European Union's Horizon 2020 Research and Innovation Programme under Grant Agreement No 825575 - European Joint Programme on Rare Diseases.

Program of the Training Course

19-21 October 2022 | online

Wednesday, 19 October 2022

Berlin time (CEST)

10:00 Welcome & Introduction

Holm Graessner, University of Tübingen, Tübingen, Germany & Gert Matthijs, KU Leuven, Leuven, Belgium

10:10 Basics of NGS diagnostics

NGS-based diagnostic approaches for rare diseases

Tobias Haack, University of Tübingen, Tübingen, Germany

Variant classification and re-classification

Jorge Pinto-Basto, Centogene GmbH, Rostock, Germany

11:40 Break

12:10 Basics of NGS diagnostics – continued –

Value of re-analysis

Katja Lohmann, University of Lübeck, Lübeck, Germany

12:55 Hands-on: variant filtering and prioritisation

Introduction

Sergi Beltran, CNAG-CRG, Barcelona, Spain

Case demo introducing key concepts

Steve Laurie, CNAG-CRG, Barcelona, Spain

- Enter structured phenotypic information (PhenoStore)
- Explanation and use of phenotypic data to filter and prioritise genomic data
- Explanation and use of in silico panels (PanelApp Genomics England)

Explain cases for homework

Questions

14:00 End of day 1



Thursday, 20 October 2022

Berlin time (CEST)

10:00 Hands-on: variant filtering and prioritisation

Review homework & questions from day 1
Steve Laurie, CNAG-CRG, Barcelona, Spain

10:20 Quality in NGS diagnostics

[Recommendations for whole genome sequencing in diagnostics for rare diseases](#)

Introduction and diagnostic strategies
Janneke (Marjan) Weiss, Radboud UMC, Nijmegen, the Netherlands
Bioinformatics and quality assessment
Erika Souche, KU Leuven, Leuven, Belgium

12:00 Break

12:30 Quality in NGS diagnostics – continued –

Ethical considerations and reporting
Helger Yntema, Radboud UMC, Nijmegen, the Netherlands

13:10 Communicating genomic results

[Co-designing models for the communication of genomic results for rare diseases: a comparative study in the Czech Republic and the United Kingdom](#)

Alessia Costa, Kings College London, London, UK

13:40 Hands-on: variant classification

Explanation and use of variant classification following ACMG guidelines
Leslie Matalonga, CNAG-CRG, Barcelona, Spain

- a) Use of webtools such as VarSome and Franklin – including CNV classification
- b) How to improve computational classification (add criterias manually: e.g. *de novo* parentally confirmed, etc.)
- c) TAG a variant accordingly in the system

Explain cases for homework

14:00 End of day 2



Friday, 21 October 2022

Berlin time (CEST)

10:00 Hands-on: variant classification

Review homework & questions from day 2
Leslie Matalonga, CNAG-CRG, Barcelona, Spain

10:20 Innovation & research

Genome data: How could (should) we use it in the clinic?
Olaf Riess, University of Tübingen, Tübingen, Germany

Economic evaluation of diagnostic innovations in medical genetics
Lisenka Vissers, Radboud UMC, Nijmegen, the Netherlands

11:50 Break**12:20 Innovation & research – continued –**

Long read sequencing for rare diseases
Alexander Hoischen, Radboud UMC, Nijmegen, the Netherlands

Treatabome
Antonio Atalaia, Sorbonne University, Paris, France

13:50 Wrap-up

Gert Matthijs, KU Leuven, Leuven, Belgium

14:00 End of workshop

Agenda may be subject to change.



SPEAKERS

Antonio Atalaia, Sorbonne University, Paris, France

Sergi Beltran, CNAG-CRG, Barcelona, Spain

Alessia Costa, Kings College London, London, UK

Tobias Haack, University of Tübingen, Tübingen, Germany

Alexander Hoischen, Radboud UMC, Nijmegen, the Netherlands

Steve Laurie, CNAG-CRG, Barcelona, Spain

Katja Lohmann, University of Lübeck, Lübeck, Germany

Leslie Matalonga, CNAG-CRG, Barcelona, Spain

Jorge Pinto-Basto, Centogene GmbH, Rostock, Germany

Olaf Riess, University of Tübingen, Tübingen, Germany

Erika Souche, KU Leuven, Leuven, Belgium

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Janneke (Marjan) Weiss, Radboud UMC, Nijmegen, the Netherlands

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