

EU Reference Laboratory for *E. coli* Department of Food Safety, Nutrition and Veterinary Public Health Unit of Food Microbiology and Foodborne Diseases Istituto Superiore di Sanità



# Training Course on WGS data use: bioinformatics tools for aiding STEC outbreak investigation

19-20 October, 2020

**Online course** 

Istituto Superiore di Sanità



Organized by:

The EU Reference Laboratory for *E. coli* 

Funded by the European Commission – DG SANTE

#### DIRECTOR OF THE COURSE

Stefano MORABITO EU Reference Laboratory for *E. coli* Department of Food Safety, Nutrition and Veterinary Public Health Istituto Superiore di Sanità

#### SPEAKERS/TUTORS

EU Reference Laboratory for E. coli, ISS, Rome, Italy

Federica GIGLIUCCI Arnold KNIJN Antonella MAUGLIANI Valeria MICHELACCI Stefano MORABITO Rosangela TOZZOLI

#### **TECHNICAL SECRETARIAT**

Guendalina FORNARI LUSWERGH

EU Reference Laboratory for *E. coli* Department of Food Safety, Nutrition and Veterinary Public Health Istituto Superiore di Sanità Rome, Italy

#### **GENERAL INFORMATION**

This event is part of the scientific and tutorial activities of the EURL-VTEC, funded by the European Commission – DG SANTE

For any information regarding the event, please send an email to crl.vtec@iss.it

## **DRAFT PROGRAM**

October 19th, Monday				
09.00	Start of the training stage			
09.10	Welcome, housekeeping, and general overview on the training course	Stefano Morabito		
	Session 1			
09.20	Introduction to Next Generation Sequencing data formats, quality check and basic analytical tools	Valeria Michelacci		
09.50	Bioinformatic analysis of NGS data: approaches and opportunities	Rosangela Tozzoli		
10.10	Introduction to the ARIES webserver user-interface	Antonella Maugliani		
10.30	Basic characterization: 7-genes Multi Locus Sequence Typing (MLST), virulotyping and serotyping	Federica Gigliucci		
11.00	Explanation of the tasks assigned to the trainees to be carried out during the hands-on session (Quality check and Trimming of WGS, 7-genes MLST, virulotyping and serotyping)			

11.20 End of session 1

### Session 2

14.00	Results discussion	Valeria Michelacci Federica Gigliucci
14.30	Assembly and assembly statistics	Valeria Michelacci
14.40	Introduction to core genome MLST (cgMLST)	Federica Gigliucci
15.00	Understanding the cgMLST and the concept of strain relatedness. Explanation of the tasks assigned to the trainees to be carried out during the hands-on session (Assembly statistics and cgMLST)	Valeria Michelacci

October 20th, Tuesday				
	Session 3			
09.00	Results discussion	Valeria Michelacci Federica Gigliucci Stefano Morabito Valeria Michelacci		
09.30	Introduction to the Outbreak investigation exercise			
10.00	Hands on: Outbreak investigation exercise			
	End of session 3			
12.00	Discussion on the outcome of outbreak investigation exercise	Stefano Morabito		
13.00	Closure			