# Inter EURLs Working Group on NGS

Valeria Michelacci

Joint Training Course on NGS June 21st, 2023





#### INTER EURLS WORKING GROUP ON NGS

#### **AIM**

to promote the use of WGS across the EURLs' networks, build WGS capacity within the EU and ensure liaison with the work of the EURLs and the work of EFSA and ECDC on the WGS mandate sent by the Commission

## from November 2017

- EURL E. coli
- EURL Listeria monocytogenes
- EURL Salmonella
- EURL Coagulase Positive Staphylococci

## Meeting twice a year (11 meetings done)

- EURL Parasites
- EURL Foodborne viruses
- EURL AMR
- EURL Campylobacter





















## Inter EURLs WG webpages

EURL *E. coli*:

https://www.iss.it/en/web/guest/-/inter-european-union-reference-laboratories-working-group-on-next-generation-sequencing-1

- EURL Salmonella
- EURL Campylobacter
- EURLAMR
- EURL Listeria
- EURL Coagulase Positive Staphylococci
- EURL Parasites
- EURL Foodborne Viruses



**Direct link!** 

Overview of conducted and planned PTs - curated by EURL Antimicrobial Resistance

Reference Whole Genome Sequencing collection – curated by EURL Salmonella

Guidance document for WGS-laboratory procedures – curated by EURL Parasites

Bioinformatics tools for basic analysis of Next Generation Sequencing data – curated by EURL VTEC

Guidance document for Whole Genome Sequencing - cluster analysis - curated by EURL Campylobacter

Guidance document for NGS-Benchmarking – curated by EURL Listeria

Inventory of training supports – curated by EURL Coagulase Positive Staphylococci

Survey on the use of NGS across the NRLs networks – curated by EURL VTEC

Supporting document for preparing high quality DNA for Whole Genome Sequencing – curated by EURL VTEC

## **Documents prepared/1**

#### **Drafting of a position paper (in review):**

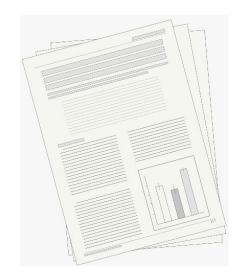
Adoption of NGS across NRLs

Harmonization of procedures

Proficiency tests

Benchmarking guidelines

Training and capacity building



#### Updated reference WGS collection curated by EURL Salmonella

Including *E. coli*, *Salmonella*, *Campylobacter*, *Listeria monocytogenes* and AMR relevant strains

e.g. E. coli: Multiple sequences of the same six **O26 STEC strains** obtained during the 1<sup>st</sup> PT on WGS (anonymzed)

Sequences available upon direct request to: crl.vtec@iss.it





## **Documents prepared/2**

#### NGS laboratory procedures curated by EURL Parasites

DNA extraction from bacteria

DNA extraction from parasites

RNA extraction from viruses

Library preparation

Standard Operating Procedures linked or appended, including protocols developed at the EURLs:

**DNA** extraction for Lm

**DNA** extraction for CPS

Library preparation (Illumina) for Lm

Library preparation (for Ion Torrent) for *E. coli* 





## **Documents produced/3**

Supporting document for DNA preparation and quality check curated by EURL VTEC

Following a request made by EFSA in the framework of the Joint EFSA-ECDC Steering Committing on Molecular Typing Data collection

AIM:

supporting the laboratories in the first steps of the NGS protocol, starting from the experiences on *E. coli, Listeria* and *Salmonella*, representing the first three pathogens for which WGS data collection has started





## **Documents produced/4**

Bioinformatic tools for basic analysis of NGS data - curated by EURL VTEC

The document aims to provide a list of **tools routinely used** by the majority of NRLs and by EURLs for NGS data analysis

Not to be interpreted as a list of validated tools, but as an **information** on those whose use is most spread among the networks of NRLs and EURLs, along with links for an easy and fast access to such tools.

Brief description of analytical steps for basic NGS analysis:

- Quality check
- Trimming

- Assembly
- Seven genes MLST

- Virulotyping
- Serotype identification
- Inference on AMR

Open source, commercial software, webservers

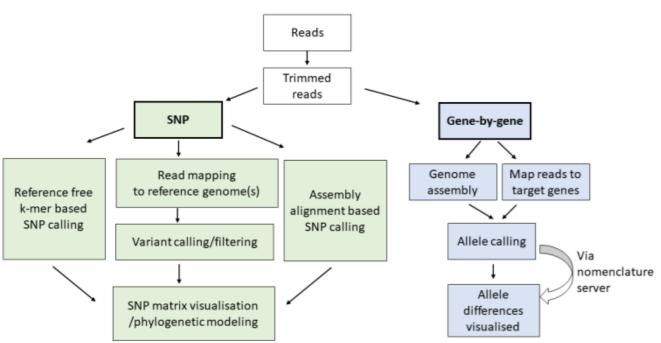




### **Documents prepared/5**

#### Guidance document for WGS - cluster analysis curated by EURL

Campylobacter





List of tools and pipelines for the different approaches for cluster analysis

Including sections dedicated to visualization tools and interpretation guidelines





## **Documents produced/6**

Guidance document for NGS-Benchmarking, curated by EURL Lm

Quality and quantity metrics

Wet and dry lab analyses

Accuracy, repeatability, reproducibility

Inventory of training supports, curated by EURL CPS

Links to resources, routinely updated

Overview of conducted and planned Proficiency Tests, curated by EURL AMR

The strategy for PT organization will be presented in a dedicated <u>webinar</u> on <u>September 29<sup>th</sup> for the benefit of the NRLs, in charge of organizing PTs at the Country level – it will be advertised soon!</u>





#### Inter EURLs initiatives/1

Joint trainings: "Introduction to Bioinformatics for genomic data mining"

• ISS (EURL *E. coli*), Rome, June 2022

24 participants from different NRLs (3 supported by EURL VTEC)

20 different countries (19 Member States + 1 Candidate Country)



RIVM (EURL Salmonella), June 2023 – here we are!





### Inter EURLs initiatives/2

#### "Science Meets Policy" conferences

- First edition: virtual conference, September 2020: over 500 participants from 49 countries worldwide
- "EU initiatives towards the large-scale use of NGS to tackle foodborne threats" organized by EFSA and the Inter EURLs WG on NGS EFSA, Parma, Italy and online, September 5<sup>th</sup>-6<sup>th</sup> 2023

Free of charge! – deadline for online participation: September 1<sup>st</sup>!









Questions?

Suggestions?