

# One Health Whole Genome Sequencing system

#### **Ernesto Liebana**

Head of Biological Hazards & Animal Health and Welfare (a.i.)



Trusted science for safe food



### Timelines



Launch of molecular typing database

October 2015

April 2017

Mandate on feasibility study for extension of the system to WGS data ECDC-EFSA
WGS
Roadmap
December
2019

December 2019

Mandate on development of One Health WGS system

### 2019 EC Mandate



Subject:

Request for the implementation of a 'One Health' system for the collection and analysis of whole-genome sequencing (WGS) data from human and food/animal isolates



#### **SCOPE**

Serving public health interests and protecting European consumers



#### **APPLICABILITY**

• WGS data of Salmonella, Listeria monocytogenes and STEC isolates



#### **OBJECTIVE**

- Promoting cross-sectorial data sharing:
  - for the purpose of molecular surveillance and continuous monitoring of foodborne pathogens
  - in case of ongoing multi-country foodborne outbreaks for the purpose of supporting the investigation of the events

### 2019 EC Mandate



### REQUIREMENT

 Implementation of two systems for the collection of WGS data of food-borne pathogens from human isolates (ECDC) and nonhuman isolates (EFSA)



- Full interoperability between the two systems
  - System interoperability
  - Semantic interoperability

The European Union needs a robust and sensitive tool for rapid detection and management of multi-country foodborne outbreaks with the ultimate purpose of serving public health interests and protecting European consumers. Emphasis will be placed on full interoperability of this joint molecular typing system between the food and public health pillars in order to ensure the protection of consumers within the EU single market in the context of a 'One Health' system approach.

### Benefit for stakeholders – EU/EEA MSs



#### Role:

✓ Data providers/Users

#### • Outcome:

- ✓ <u>Secured system</u> → data confidentiality
- ✓ <u>Customer-oriented service</u> → based on Member States' capacity
- ✓ <u>Capacity building</u> → decentralisation of the bioinformatic analysis
- ✓ <u>Database searchable</u> → data comparison and visualisation

#### • Added value:

- ✓ More efficient investigation at country level
- ✓ Trust in the EU food safety system
- ✓ Engagement





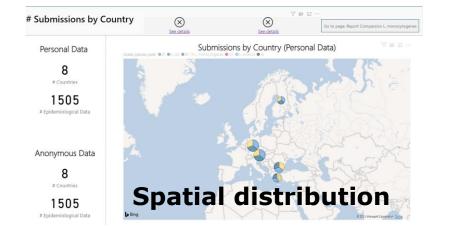
**European Food Safety Authority** 

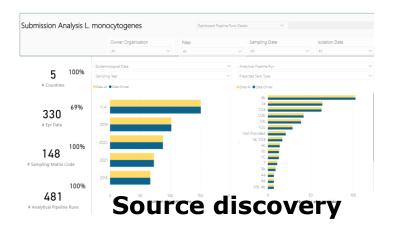
#### **EFSA One Health WGS System Portal**

The EFSA One Health WGS System has been developed by EFSA as response to the European Commission mandate for the collection of sequencing data of isolates from food, feed, animals, and the farm and food processing environment, for the purpose of supporting ECDC in rapid detection of multicountry foodborne outbreaks

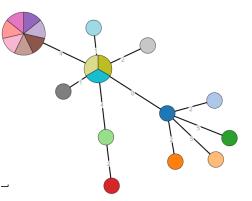


efsa■





These are prototypes based on fictional data



#### **Phylogenetic** analysis

### **Promoting EFSA core values**



#### Role:

√ Coordinator

#### • Outcome:



- ✓ Real-time data availability
- ✓ System incorporated in <u>IT EFSA strategy and architecture</u> → Azure cloud system



- ✓ Use of open-source community-validated bioinformatic tools → Agility and resilience to technological changes
- ✓ Open-source distribution of bioinformatic pipeline → <u>Transparency</u>

#### • Added value:

✓ Capacity to <u>promptly support</u> outbreak investigation at EU level

### One Health WGS system

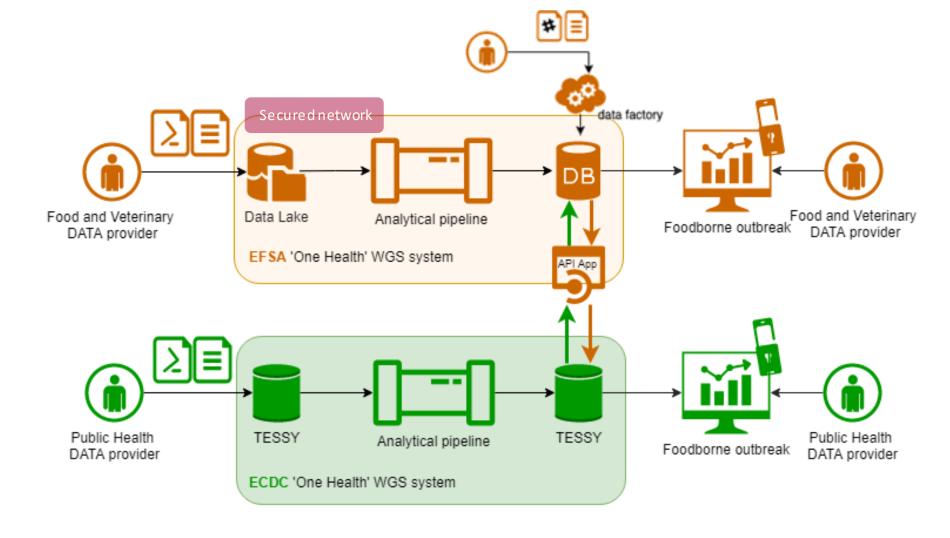






Cross-sector matches





### EFSA-ECDC interoperability



Genome profile of the Human representative isolate



Matching profiles = isolates showing similarity in their genomic profiles based on a specific threshold

EFSA checks possible matches for clusters from ECDC



ECDC calculates clusters



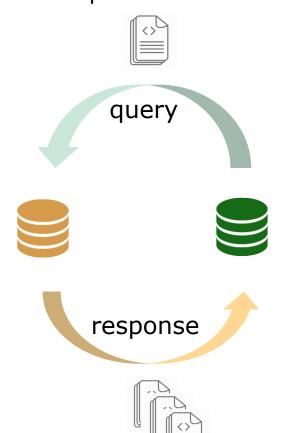
Genome profiles of Matching food/animal isolates

### An example



#### Genome profile of the Human representative isolate

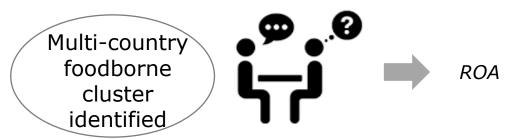
Yes!! we found 3 food isolates from fish products countries X and Y all from 2019 with 3, 4, and 6 allelic differences to any human isolates.



There is a cluster of 5 human isolates of Listeria monocytogenes. Do you have isolates with less of 8 alleles of difference in cgMLST?



Genome profiles of Matching food/animal isolates



### Total cost



- Setup (from December 2019 to December 2022)
  - 1.5 Mio
  - Staff (3.0 FTE) = 0.3 Mio
- Running
  - IT Cost 0.20-0.35 Mio/year
  - Staff (2.2 FTE) = 0.22 Mio /year

### **Timelines**



Functional system opened to final users

June 2022

## December 2022

Consolidation and troubleshooting

System fully operational

From January 2023

### Thank you for your attention





### **Acknowledgments**

EFSA: Mirko Rossi, Valentina Rizzi, Denise Pezzutto, Ernesto Liebana, Chiara Bianchi, Giancarlo Costa, Giovanni Iacono, Giulio Di Piazza

**ECDC** 

EURL E. coli

EURL *Listeria monocytogenes* 

EURL Salmonella

SANTE G4