

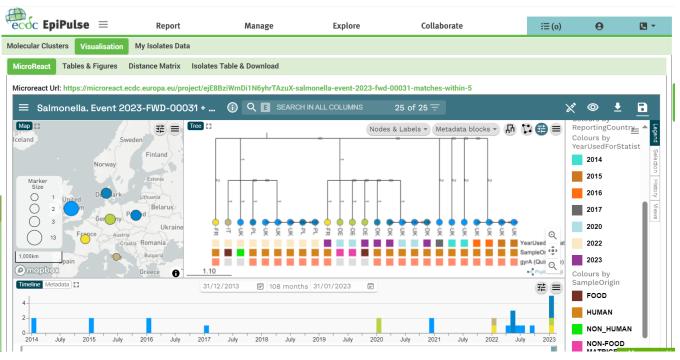
FWD AMR-RefLabCap, 2nd Multidisciplinary training workshop

EU level surveillance using EpiPulse molecular typing tool

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Microbiology and Molecular Surveillance group, ECDC







Legislation

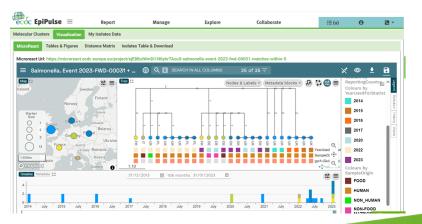


 New Regulation On Serious Cross Borders Threats to Health (2022/2371)

Article 13

Epidemiological surveillance

- 3. The national competent authorities referred to in paragraph 1 shall communicate the following information, based on agreed indicators and standards, to the participating authorities of the network for epidemiological surveillance:
 - (d) molecular pathogen data, if required for detecting or investigating serious cross-border threats to health



Core components of ECDC WGS-enhanced surveillance



- TESSy The system for submitting data to ECDC
- Epipulse The ECDC platform for nominated users
 - Molecular Typing Tool A tool for visualising WGS-based results integrated with epidemiological data
 - Events, Forum & News Notification and discussion platform, can also connect to the Molecular Typing Tool
- ECDC Bioinformatics infrastructure
 - Bionumerics
 - Open-source tools to complement Bionumerics

Use of Public data



- ECDC downloads public data when appropriate for ongoing investigations
- The public data are analysed using the same ECDC bioinformatics pipelines as data submitted through TESSy
- Several methods are used to find relevant public data, mainly:
 - NCBI Pathogen Detection
 - Enterobase

Matching with EFSA



- Every week, every cluster for Listeria (where we perform continuous surveillance) is queried to the EFSA system with a 7 AD cut-off
- Any matches are saved in the ECDC system temporarily (updated every week if they are found again)

 ECDC staff can also perform ad-hoc queries whenever it is deemed necessary for an ongoing investigation, this can currently be performed for Listeria, and Salmonella, and with up to 20 AD cut-off

Data visibility restrictions



- National identifiers can only be seen by ECDC and the submitting country
- For EFSA data, country of origin can only be seen by ECDC and the same country, and further restrictions on data visibility can be applied by the EFSA users
- These restrictions on EFSA data do not apply to data that are also in the public domain

Access rights to the typing tool for the FWD domain



 Access rights correspond to TESSy Download rights for subjects LIST/LISTISO (*Listeria*), SALM/SALMISO (*Salmonella*), CAMP/CAMPISO (*Campylobacter*), and ECOLI/ECOLIISO (*E. coli*)



Welcome to EpiPulse - the European surveillance portal for infectious diseases!

Through this portal you can report and share information on cases or events of infectious diseases/pathogens, explore European data or exchange information with other nominated users.

Note: EPIS ELDSNet continues to work as usual, migration will occur on a later stage. The function and use of EWRS remain unaffected.

TESSy can also be accessed through the EpiPulse menu. To report cases or molecular typing data go to 'Report' \rightarrow 'Cases' or 'Sequence data'. The existing TESSy URL and functionalities have not changed.

Should you have any further questions or suggestions for improvement, please send them to epipulse@ecdc.europa.eu.

Click on the menu above to get started. Through the help button on the top right-hand side of the screen, you can access built-in tutorials to guide you around. You can also watch the videos to learn how to use EpiPulse:



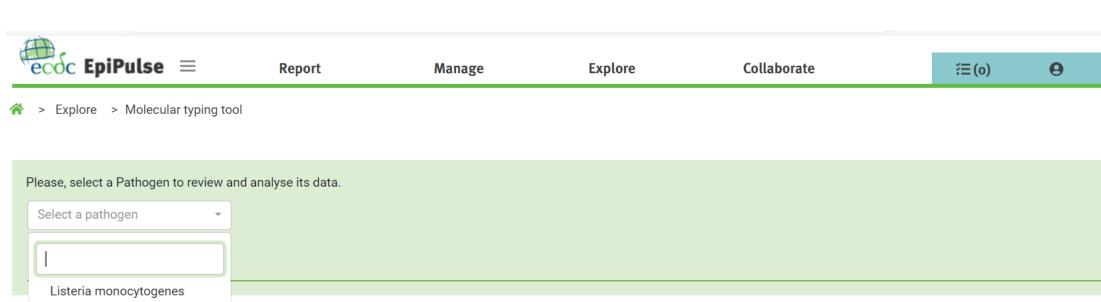












Salmonella

Escherichia coli

Campylobacter

My Isolates Data

Please, select a Pathogen to review and analyse its data.

Pseudomonas aeruginosa Neisseria meningitidis

Legionella pneumophila

Klebsiella pneumoniae

C. difficile

Mycobacterium tuberculosis

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ECDC Tools

Manage cookies





Report

Manage

Explore

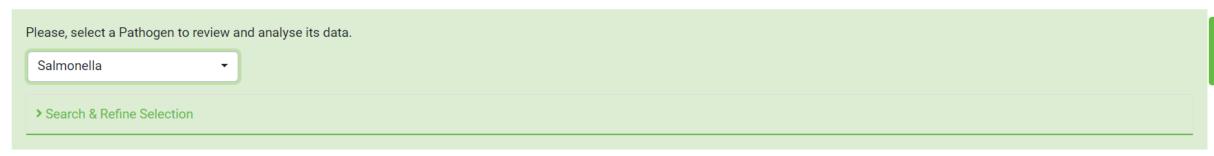
Collaborate

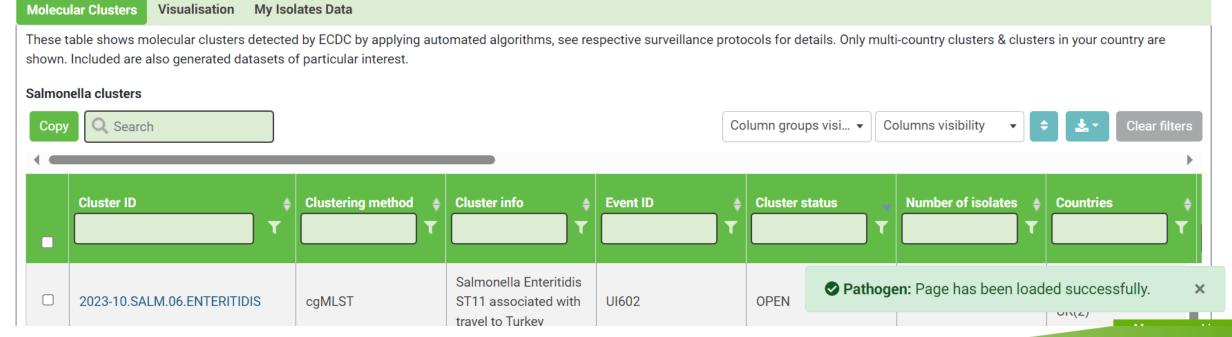
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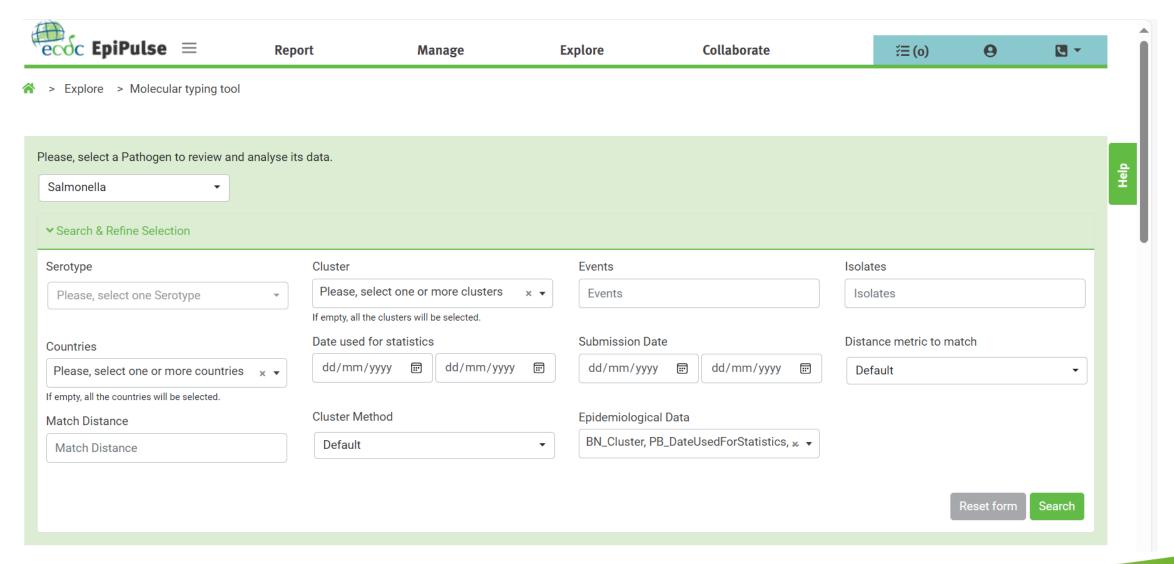
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> Explore > Molecular typing tool

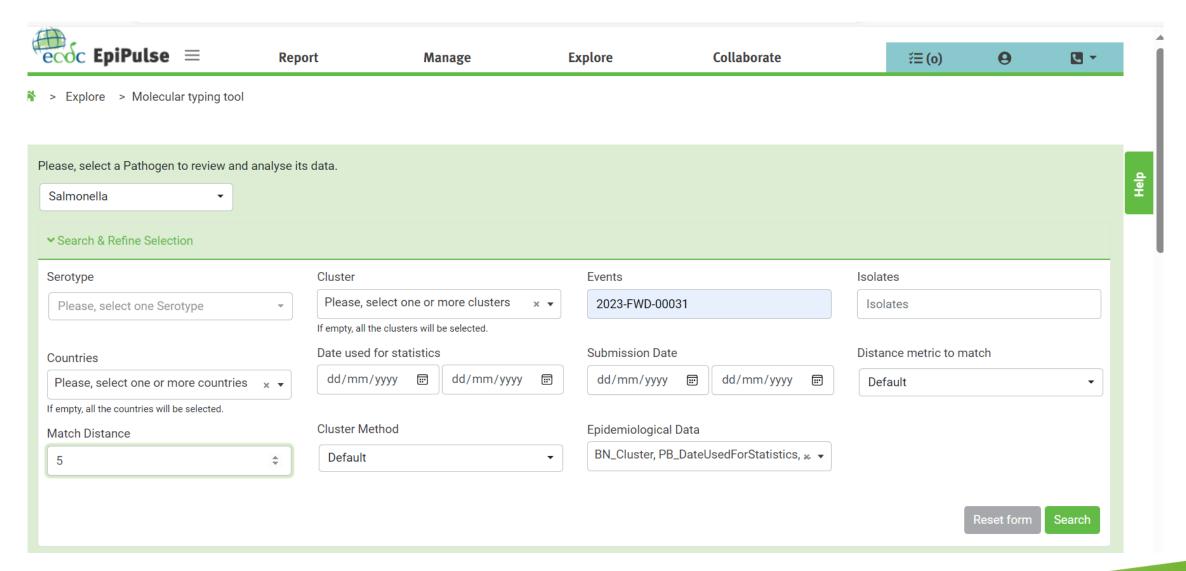


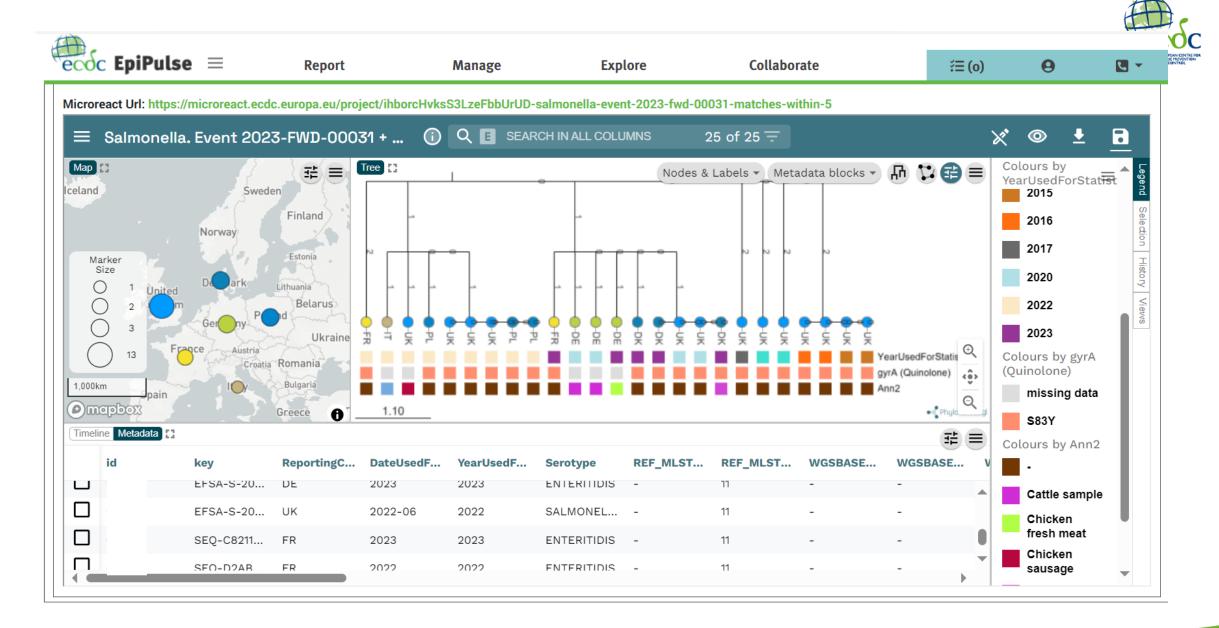












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Report

Manage

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Explore Collaborate

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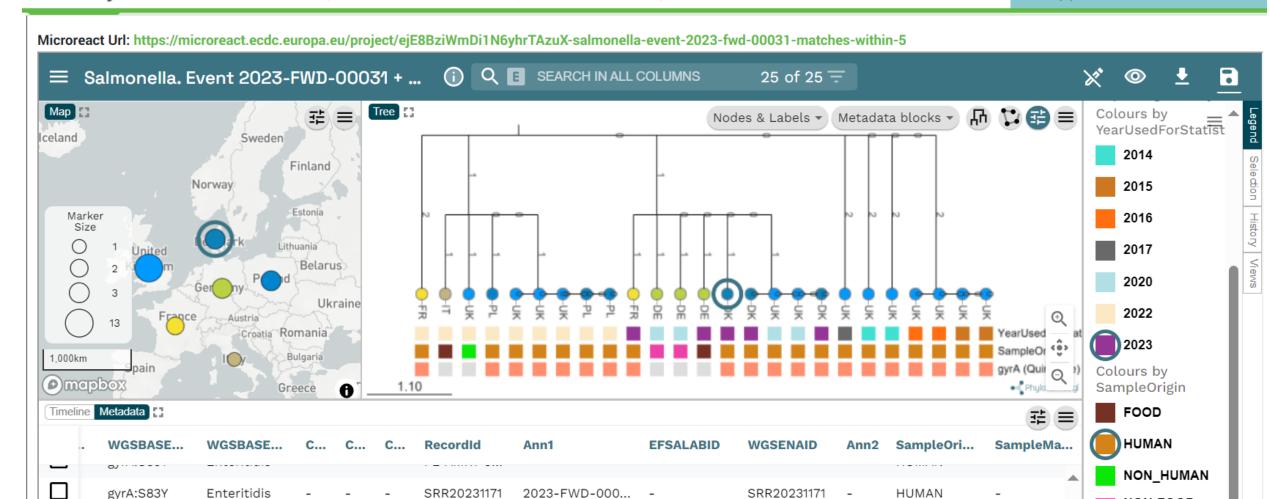
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NON-FOOD MATRICES

miceina data

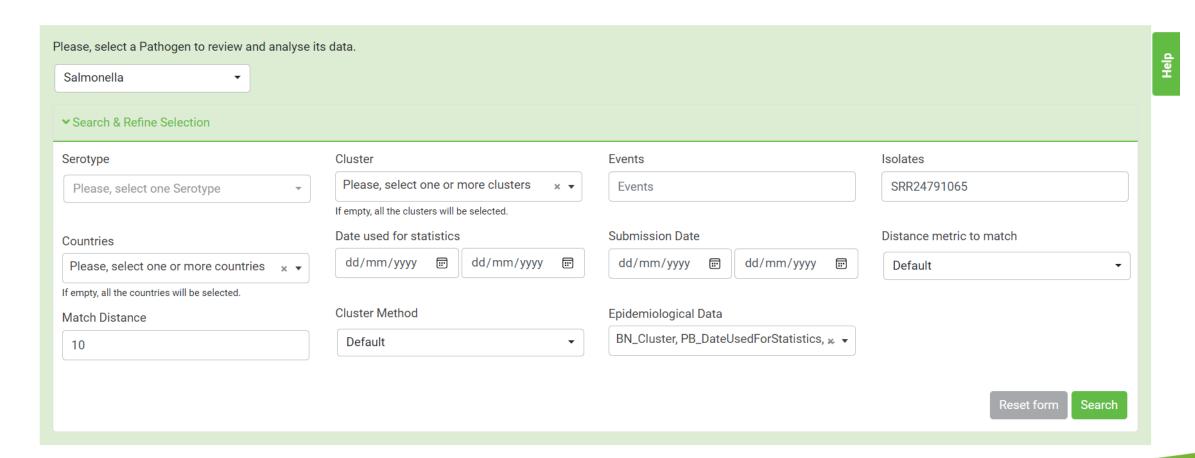
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(Quinolone)

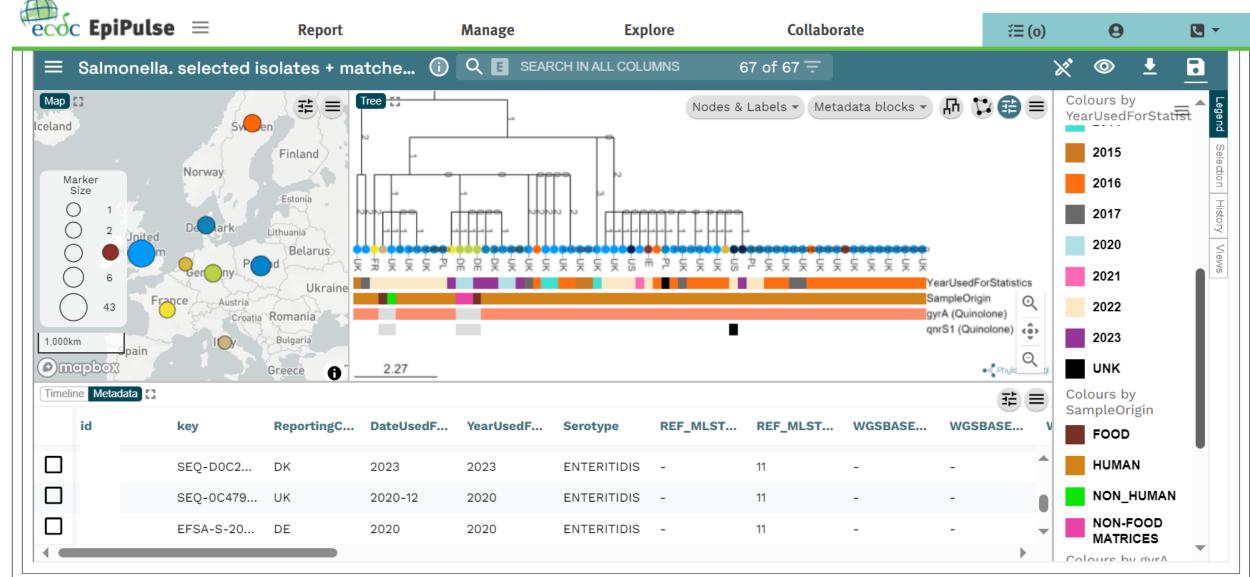




> Explore > Molecular typing tool

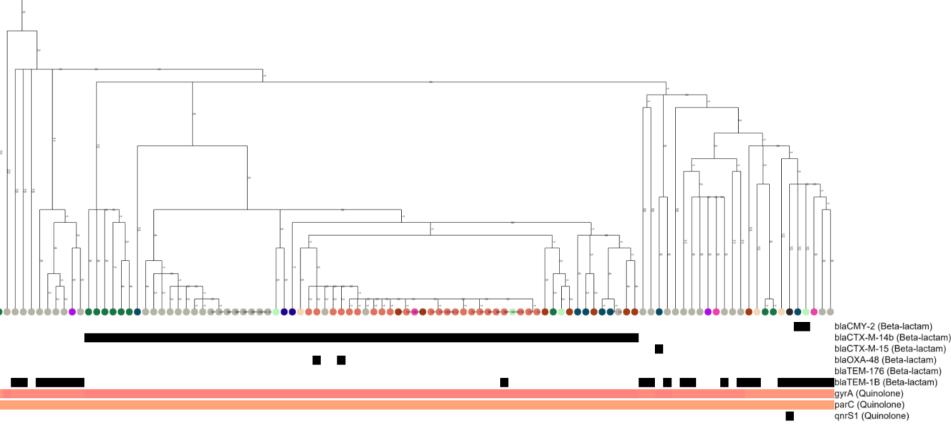






AMR visualisation

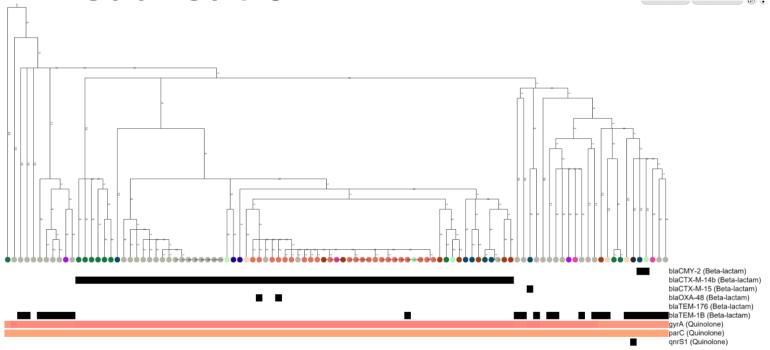




- Currently absence/presence of resistance genes are shown, as well as resistance point mutations
- There is also a detailed table with length/identity values for each detected gene

AMR visualisation





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To be able to use EpiPulse Molecular Typing tool to visualise your own data in the context of the data in ECDC db and those that would cluster with isolates from EFSA db you need:

- Have access rights for TESSy Download rights for subjects SALMISO (and SALM), CAMPISO (and CAMP).
- If you do not have this your country's National Coordinator need to nominate you. You will then be granted access to the system.
- Contact us if you need support with this process
- Contact us if you need support with sequence data uploads or with learning more about the Molecular typing tool.



Thank you!