EFSA WGS PLATFORM FOR SUPPORTING AMR DATA COLLECTION

Mirko Rossi, 23/05/2023



OBJECTIVE OF THE AMR GENE DETECTION SERVICE

To offer to MS the opportunity to use EFSA computing resources for extracting AMR genes from WGS data

To give access to User-Friendly Interface for accessing and managing the data in a secured and private environment, no sharing of data

Automate mapping against the PARAM catalogue and different downloads options for the results



DEFINE THE AMR GENE DETECTION SERVICE

• The AMR gene detection service is...

The AMR gene detection service is not...

a web application for extracting AMR genes and mutations conferring resistance to beta-lactams in a standardized format, already mapped against EFSA PARAM catalogue

designed so that all users of the same organization access the same environment and users from other organizations cannot see the data a data submission tool, submission remains trough Data Collection Framework (DCF) of EFSA. EFSA cannot use the data temporarily stored in the system

a tool for sharing information among MSs and between MS and EFSA

a system allowing permanent storage of raw data



Adaptation of WGS portal implemented for One Health WGS system

Use **same pipeline** implemented for the EFSA One Health WGS system - AMR genes are extracted with ResFinder 4.1

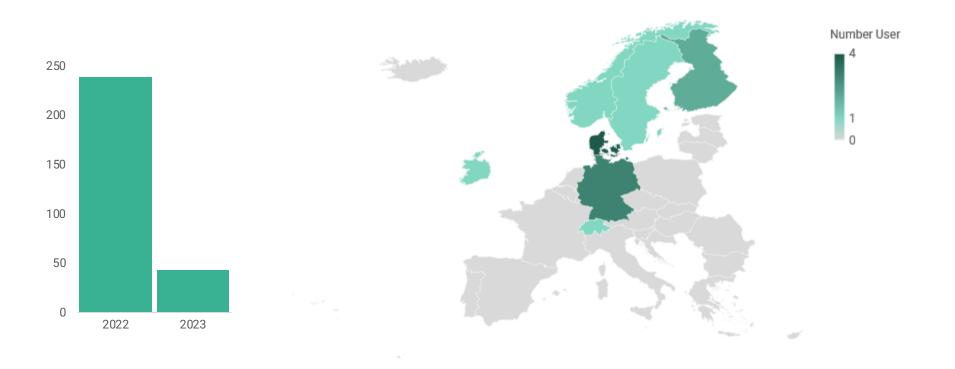
Catalogue enrichment of the analytical result \rightarrow genes found by ResFinder mapped against PARAM catalogue

Option to **download AMR results** in several formats, including precompiled for EFSA submission tool

Access is granted to data providers identified by the Reporting Officer



AMR USERS AND USAGE AS OF MAY 2023

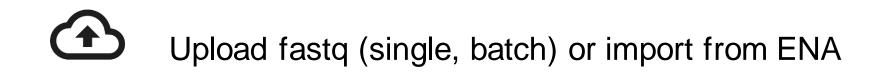


1.1.1

Powered by Bing © GeoNames, Microsoft, OpenStreetMap, TomTom



USER ACTIONS





Execute analytical pipeline



Monitor execution and results



Download results (also in DCF compatible format)



Store or delete



INFORMATION REQUESTED FOR UPLOAD

Isolate ID

data provider's isolate identification code

Instrument Model

the model of instrument used for sequencing

Species

"Salmonella enterica" or "Escherichia coli"

Library Layout

SINGLE or PAIRED

Raw Reads

Names of the FASTQ-files

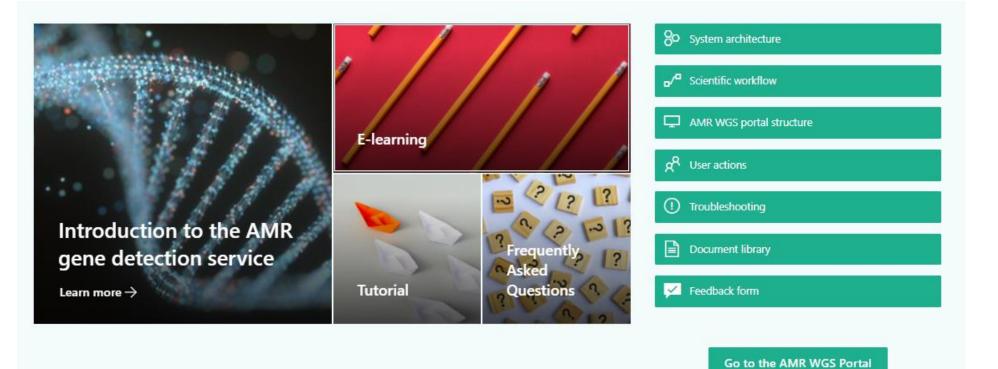




LEARNING MATERIALS

• Sharepoint website:

https://efsa815.sharepoint.com/sites/AMRWGSportaltraining



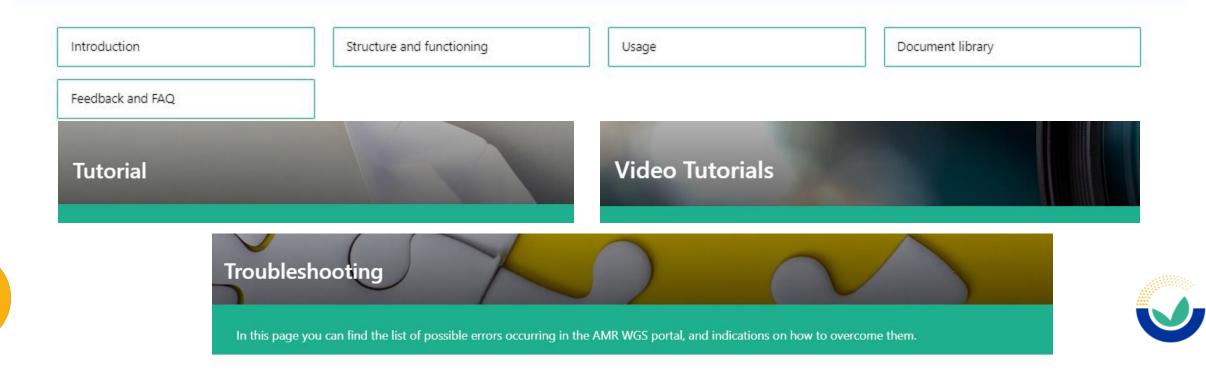
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LEARNING MATERIALS

E-learning for AMR gene detection service

Follow the links to the site pages to learn more about different aspects of the AMR gene detection service.

For a comprehensive e-learning experience, follow the modules in the suggested order.



TAKE HOME MESSAGES

EFSA doesn't consider WGS generated for the scope of AMR monitoring and uploaded to the AMR Gene Extraction Service as submission for outbreak investigation

AMR Gene Extraction Service is not a data collection tool, EFSA cannot use the data upload to the system

Only Data Providers formally appointed from MS for the scope of submitting WGS data for outbreak investigation submit data to the EFSA One Health WGS System

It is on discretion of the MS through the appointed Data providers of the EFSA One Health WGS System to decide if the WGS data generated for AMR monitoring activity can be submitted also with the scope of supporting outbreak investigation



Microsoft Teams

AMR demo

2023-05-23 06:44 UTC

Recorded by ROSSI Mirko Organized by ROSSI Mirko

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